

Contents lists available at ScienceDirect

Computer Methods and Programs in Biomedicine

journal homepage: http://www.elsevier.com/locate/cmpb



Non-Invasive Estimation of Coronary Velocity: A Computational Framework for Computed Tomography Perfusion Imaging

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ARTICLE INFO

Article history:
Received 1 May 2013
Received in final form 10 May 2013
Accepted 13 May 2013
Available online 15 May 2013

Keywords: Coronary Artery Imaging, CT Myocardial Perfusion Imaging, Computational Fluid Dynamics, Blood Velocity Estimation, Non-Invasive Coronary Flow Assessment

ABSTRACT

Background and Objectives: Ischemic heart disease (IHD) results from insufficient blood flow to the myocardium, primarily caused by coronary artery stenosis. Existing clinical methods rely on invasive procedures for assessing coronary hemodynamics. Computational methods, particularly those based on coronary CT angiography (cCTA) and CT myocardial perfusion imaging (CT-MPI), offer non-invasive alternatives.

Methods: This study introduces a computational framework leveraging CT-MPI for estimating blood velocity in stenosed coronary arteries under steady flow conditions. The framework integrates diverse simulation techniques, including computational fluid dynamics (CFD) and advection-diffusion simulations.

Results: Results demonstrate promising agreement between estimated and assigned velocities, suggesting the potential of this method for non-invasive coronary flow assessment. The observed dependence on Reynolds number highlights the need for further refinement and investigation into estimation methods.

Conclusions: Future work may involve exploring advanced models to mitigate biases and enhance accuracy. Ultimately, this research contributes to the potential of non-invasive assessment of blood flow dynamics from routine scans, with implications for improved diagnosis and targeted therapeutic interventions.

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1. Introduction

Ischemic heart disease (IHD) arises from inadequate blood flow to the myocardium, primarily due to coronary artery stenosis. While coronary artery anatomy plays a crucial role in understanding blood perfusion, it often fails to directly correlate with the pathophysiol-

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ogy of stenosis causing flow restriction. Consequently, the current clinical standard for assessing the physiological effects of coronary disease on hemodynamics relies on invasive pressure catheterization during angiography. However, the lack of a routine non-invasive flow assessment remains a significant challenge, hindering our understanding of the underlying mechanisms of coronary flow limitations [? ?].

Computational methods based on coronary CT angiography (cCTA) offer a promising avenue for capturing hemodynamic features of the blood flow in coronary arteries. By leveraging cCTA-derived coronary geometry, computational fluid dynamics (CFD) simulations enable comprehensive assessments of hemodynamics, including pressure, velocity, and wall shear stress [?????]. However, the accuracy of these assessments hinges on accurate boundary conditions and flow split assumptions [??].

Another potentially valuable CT-based technique for visualizing and quantifying coronary blood flow is CT myocardial perfusion imaging (CT-MPI). Although unexplored for this purpose, the dynamic transit of contrast through the coronary arteries holds promise for calculating mean blood velocity. In 2015, Eslami [1] pioneered the transluminal attenuation flow encoding (TAFE) method, utilizing experimental phantom CT images to estimate blood velocity from contrast transition along a tapered vessel. Assuming unidirectional flow and dominant convective forces, TAFE calculates mean flow rate. Eslami subsequently introduced a CFD-based correction factor to further refine accuracy [2]. While their studies focused on experimental data under laminar flow conditions, we have demonstrated the potential of this approach under a wider range of flow conditions, suggesting its applicability to CT-MPI-based applications.

In this study, we present a computational framework for estimating blood velocity in stenosed coronary arteries with steady flow based on contrast agent transition. Our pipeline encompasses diverse simulation techniques to model blood flow, contrast agent transit, and velocity calculation. Following a detailed description of the theoretical background and simulation setup, we validate our method by comparing calculated velocities with preassigned values under steady flow conditions. Finally, we conclude by discussing the implications and potential future directions of this work.

2. Methods

2.1. Contrast Transport and Blood Flow

To rigorously validate the accuracy and robustness of our velocity estimation pipeline, we established a multistep computational framework simulating both blood flow and contrast agent dispersion within a stenosed arterial segment. This framework encompassed two key components:

- Stenosed Vessel Geometry: This study employed a three-dimensional, idealized model of a non-axisymmetric stenosis to represent the complex geometric variations observed in diverse vasculatures. The modeled artery extended along the x-axis from -9 cm to 72 cm, with the stenosis located at x = 0 causing a 75% reduction in the cross-sectional area. Notably, the stenosis throat incorporated an eccentricity to promote flow instability without requiring additional velocity field perturbations. The specific geometry is depicted in Figure ??. This configuration was inspired by a prior study investigating turbulent flow transition in stenosed arteries [reference].
- 2. Modeling blood flow in stenosed arteries: In order to implement our pipeline of flow and contrast transition in the stenosed artery, two partial differential equations are the key components: Navier-Stokes equations, simulating the blood flow assuming a newtonian and incompressible fluid [], and Advection-Diffusion equation, modeling the contrast agent transport in the blood []. The governing equations of the Navier-Stokes equation is presented below:

$$\frac{\partial U}{\partial t} + (U.\nabla)U + \frac{\nabla P}{\rho} = \nu \nabla^2 U, \qquad \nabla \cdot U = 0$$
(1)

Where U is the time-dependent velocity vector field, P is the pressure field inside the geometry domain, ρ is the blood flow density, and ν is the dynamic viscosity. A Neumann type boundary condition was assigned to the inlet of the vessel and a non-slip boundary condition (u = 0) was assigned to the walls. To encompass the range of flow regimes encountered in diverse arterial beds, we varied the Reynolds number (Re) from 100 to 1000 in increments of 100. This parameter selection captures the spectrum of physiological flow conditions observed in coronaries and cerebral arteries (Re 100-300), carotid arteries (Re 600), and the aorta (Re 700-1000). By simulating across this extensive Re range, our investigation encompasses the complex, multi-regime flow dynamics found in human vasculature.

3. Modeling contrast agent transition in coronaries: The next crucial step in our framework involves simulating the spatiotemporal evolution of the contrast agent concentration within the vasculature. This necessitates incorporating the com-

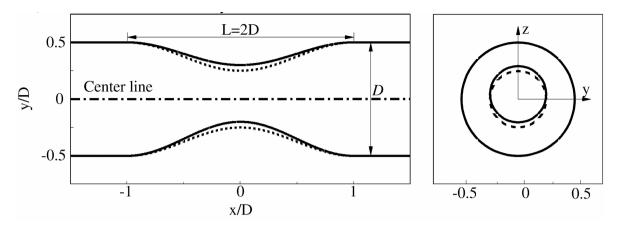


Fig. 1. Eccentric Stenosis Geometry

plex interplay between advection, driven by the bulk blood flow, and diffusion, arising from microscopic Brownian motion. To adequately capture this interplay, we employ the advection-diffusion equation, a fundamental partial differential equation widely used in fluid dynamics and mass transport phenomena. In our context, the advection-diffusion equation takes the following form, where $C(x,y,z;t)(\frac{ml}{mg})$ represents the contrast agent concentration at spatial location (x,y,z) and time t:

$$\frac{\partial C}{\partial t} + (U.\nabla)C = D\nabla^2 C \tag{2}$$

Here, U(x,y,z) denotes the blood velocity at the spatial location (x,y,z), which derives the transport of contrast agent in the blood and is determined by the preceding CFD simulation. D is the diffusion coefficient, which quantifies the rate of contrast spread due to random molecular motion, with a value of 0.04 cm²/s as reported by []. To assign the inlet boundary conditions of the contrast agent we used the following equation []:

$$C_{inlet} = C_{min} + 0.5(C_{max} - C_{min})[1 - cos(\pi \frac{t - T_s}{T_d})], \label{eq:cos}$$

where C_{min} and C_{max} are the minimum and maximum values of the contrast in the inlet of the artery, chosen to be 0 and 1 respectively. And T_s is the time at which the bolus arrives and T_d is time for which the advection-diffusion is considered.

4. Contrast Dispersion: Following the simulations, we extract mean velocity information from the observed contrast transition. Based on the simulated contrast bolus, we perform an inverse estimation to compute the centerline velocity along the arterial segment. While solving the full contrast disper-

sion equation would be ideal, the inherent limitations of CT-MPI in terms of spatial resolution and noise hinder the accurate reconstruction of the 3D velocity field, particularly in the context of coronary arteries. Therefore, to account for these limitations, we adopted a simplified approach assuming unidirectional flow with negligible radial dispersion of contrast. This allowed us to estimate the average velocity across the arterial lumen, albeit sacrificing the full spatiotemporal resolution offered by the advection-diffusion equation:

$$v_{mean} = \frac{\frac{\partial C}{\partial t}}{-\frac{\partial C}{\partial x}} \tag{4}$$

This equation relates the centerline velocity (V) at a specific axial location (x) along the artery to the temporal and spatial variations in contrast concentration (C) observed during the simulation. The numerator, representing the rate of change in contrast over time $(\frac{\partial C}{\partial t})$, is obtained by computing the instantaneous derivative of the time-intensity curve at the desired location. Likewise, the denominator, reflecting the spatial gradient of contrast along the centerline $(\frac{\partial C}{\partial x})$, is calculated by taking the derivative of the contrast profile with respect to the axial coordinate at the peak contrast intensity.

2.2. Simulations Setup

CFD simulations To characterize the onset and nature of turbulence transition in a stenosed artery, we employed direct numerical simulations (DNS) of the steady flow within a non-axisymmetric 3D stenosis model (Khan et al., 2018). Using the Oasis flow solver, we implemented CFD simulations of the Navier-Stokes equations for incompressible Newtonian blood ($\mu = 0.004 Pa * s, \rho = 1057 \frac{kg}{m^3}$). To accurately capture



Fig. 2. The results of implementing CFD simulations in stenosed pipe. changing Re=100-1000

the evolving turbulent features, time steps were meticulously chosen as 0.0001 seconds. A pulsatile Womersley profile was applied at the inlet to mimic realistic blood flow, while the outlet boundary conditions were dynamically adjusted through a pressure coupling scheme. The Navier-Stokes equations were solved for blood pressure and velocity using a second-order polynomial interpolation for velocity and a first-order scheme for pressure. The resulting velocity fields were subsequently exported for further analysis in the Advection-Diffusion simulation stage. This DNS approach was repeated across a range of Reynolds numbers (Re = 100 - 1000). The stenosed artery geometry was meshed with approximately 6000 triangular elements. The simulations were done on Niagara supercomputers of the Canada Research Alliance []. Advection Diffusion Simulations To model the transport of a contrast agent within the fluid flow, we employed the FEniCS framework [reference] to numerically solve the advection-diffusion equation. The velocity field, computed a priori using CFD simulations, was incorporated as the convection velocity within the advection term of the governing equation. The FEniCS framework facilitated the solution of the variational form of the equation using finite element methods. For each time step, the corresponding CFD velocity field was loaded to ensure consistency with the evolving flow dynamics. Inlet boundary conditions were prescribed in accordance with equation, with $C_{min} = 0$, $C_{max} = 1$, $T_s = 0$, and T_d adjusted between 12 and 85 seconds to accommodate the range of Reynolds numbers investigated. The spatiotemporal evolution of contrast agent concentration along the vessel is visualized in Figure ??. Contrast Dispersion Simulations To quantify the mean velocity within the vessel, we employed a backcalculation procedure based on the advection-diffusion equation (Equation). This process involved two key steps:

- Extraction of Attenuation Curves We first extracted
 the contrast agent's attenuation over time at the inlet of the vessel from the advection-diffusion simulation results, forming the time-attenuation curve.
 Additionally, we extracted the attenuation profile
 along the vessel's centerline at the peak time of contrast agent concentration, creating the centerlineattenuation curve.
- 2. Linear Regression Modeling and Mean Velocity Calculation Each extracted curve was then subjected to linear regression analysis, establishing a linear relationship between attenuation and the respective independent variable (time for the time-attenuation curve and spatial position for the centerline-attenuation curve). The slopes of the fitted linear models, representing the temporal and spatial concentration gradients ($\frac{\partial C}{\partial t}$ and $\frac{\partial C}{\partial x}$, respectively), were subsequently incorporated into equation to back-calculate the mean velocity within the vessel.

This approach enabled us to leverage the simulated advection-diffusion data to quantify the flow velocity without requiring direct measurement, providing valuable insights into the fluid dynamics within the vessel.

3. Results

In our framework, multiple simulations were conducted to derive meaningful insights. The obtained results are summarized as follows:

3.1. CFD Simulations

Figure ?? illustrates the Q-criterion, providing a visual representation of turbulent transition along the artery at various Reynolds numbers (Re=100-1000). The CFD simulations aimed to assign a steady velocity to the

stenotic pipe, ranging from 1.33 cm/s to 13.3 cm/s at different Reynolds numbers.

3.2. Advection-Diffusion Simulation

The advection-diffusion setup was adjusted and validated against theoretical formulations. Figure $\ref{eq:contrast}$ depicts the contrast dynamics at Re=1000 at four time points (t=1,5,10,15 seconds), showcasing contrast perfusion along the artery. The final output illustrates the motion of contrast agents driven by the assigned velocity.

3.3. Contrast Dispersion Simulation

Results of the contrast dispersion, estimating velocity, were compared to the mean velocity assigned in the CFD simulations and summarized in Table ??. The mean velocity error ranged from 0.1% to 9%, with an average of 2.5% for different Reynolds numbers. The transit delay varied from 46% to 65% of the simulation runtime. An observed increase in the transit delay to simulation run-time ratio was noted with higher Reynolds numbers. The correlation coefficient for calculated velocity vs. Reynold's number.

Table 1. Comparison of the CFD assigned velocity and the estimated velocity using our contrast dispersion method at varying Revnolds numbers and Advection-Diffusion simulation run-times.

cynoids numbers and rection-Diffusion simulation run-time				
Re	run-time (s)	V_{CFD}	V_{est}	Err (%)
100	85	1.3	1.20	9.22
200	45	2.6	2.52	5.06
300	30	3.9	3.63	9.05
400	25	5.3	4.83	9.25
500	18	6.6	6.50	2.38
600	16	7.9	7.75	3.01
700	15	9.3	9.04	3.05
800	14	10.6	10.46	1.86
900	13	11.9	11.88	0.90
1000	12	13.3	13.31	0.12
	Re 100 200 300 400 500 600 700 800 900	Re run-time (s) 100 85 200 45 300 30 400 25 500 18 600 16 700 15 800 14 900 13	Re run-time (s) V _{CFD} 100 85 1.3 200 45 2.6 300 30 3.9 400 25 5.3 500 18 6.6 600 16 7.9 700 15 9.3 800 14 10.6 900 13 11.9	Re run-time (s) V _{CFD} V _{est} 100 85 1.3 1.20 200 45 2.6 2.52 300 30 3.9 3.63 400 25 5.3 4.83 500 18 6.6 6.50 600 16 7.9 7.75 700 15 9.3 9.04 800 14 10.6 10.46 900 13 11.9 11.88

4. Discussion

Our findings underscore the potential of estimating lumen velocity through contrast perfusion imaging, despite inherent errors influenced by the flow regime. Notably, the estimated velocities closely align with the assigned velocities in CFD simulations across the Reynolds number spectrum, with errors predominantly below 10%. Importantly, higher Reynolds numbers correspond to shorter simulation run-times and smaller percentage errors. In aggregate, the data suggests robust agreement between estimated and assigned velocities, indicating the efficacy of our method across diverse flow conditions.

These results bear significant implications for the expanding domain of non-invasive coronary flow assessment employing CT-MPI. The observed correlation between accuracy and Reynolds number necessitates indepth exploration and refinement of estimation methods. Investigating advanced models that incorporate periodic flow assumptions or account for Reynolds number-specific corrections holds promise in mitigating biases and enhancing the precision of velocity and flow rate estimations. Ultimately, our research has the potential to facilitate non-invasive evaluation of blood flow dynamics directly from routine scans, contributing to improved diagnoses and targeted therapeutic interventions.

4.1. Limitations and Future Work

While our study demonstrates promising results, it is essential to acknowledge certain limitations. The current method's dependence on Reynolds number prompts further investigation into refining estimation techniques for varying flow conditions. Future work may explore additional factors influencing accuracy and employ more sophisticated models to address these considerations. Additionally, clinical validation and comparisons with established methods will be crucial to establishing the broader applicability and reliability of our approach.

5. Conclusions

In this study, we have presented a comprehensive computational framework for estimating blood velocity in stenosed coronary arteries using contrast perfusion imaging. Our results indicate strong agreement between estimated and assigned velocities across varying Reynolds numbers, underscoring the method's potential for non-invasive coronary flow assessment. The observed dependence on Reynolds number highlights the need for further refinement and investigation into estimation methods. Future work may involve exploring advanced models to mitigate biases and enhance accuracy. Ultimately, this research contributes to the potential of non-invasive assessment of blood

Acknowledgements

MOK would like to acknowledge funding from the Natural Science and Engineering Research Council of Canada.

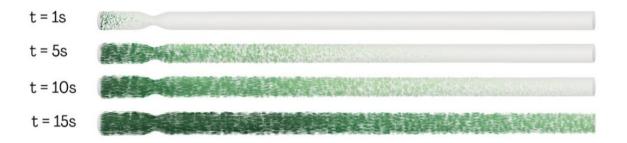


Fig. 3. Advection Diffusion Simulation Results, Contrast transition as the result of convection-diffusion simulations for Re=1000 in after 1, 5, 10 and 15 sec of running the simulations.