



**Application for a Postgraduate Scholarship  
or Postdoctoral Fellowship  
(FORM 201)**

AID
CTTEE
Date  2017/09/29

Type of Award <b>PGS D</b>	Reference No. 401226308
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Family name of applicant  Cassidy	Given name  Tyler	Initial(s) of all given names  TT	Personal identification no. (PIN)  <b>Valid</b> 439027
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**ADDRESSES. Changes to any of the information below must be sent to [schol@nserc-crsng.gc.ca](mailto:schol@nserc-crsng.gc.ca).**

Current mailing address 6- 9230 Rue Lajeunesse Montreal, QC CANADA  H2M 1S2	Permanent address (if different from current mailing address) 11 Reighley Close Red Deer, AB CANADA  T4P 3V7
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If current mailing address is temporary, indicate leaving date	Telephone number at permanent address
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Telephone number 11 (403) 8726089	Facsimile number	E-mail address NSERC will use this information as the initial point of contact. tyler.cassidy@mail.mcgill.ca
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**CITIZENSHIP**

<input checked="" type="checkbox"/> Canadian citizen	<input type="checkbox"/> Permanent resident of Canada	<input type="checkbox"/> Other
Indicate date of landing as stated on official immigration document		Indicate country of citizenship

**LANGUAGE OF CORRESPONDENCE**

I wish to receive my correspondence in:

<input checked="" type="checkbox"/> English	<input type="checkbox"/> French
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**University responsible for the internal selection process (Not applicable for PGS applications submitted directly and PDF applications.)**

McGill  
Mathematics and Statistics



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**ACADEMIC BACKGROUND (include only current and past degree programs)**

Degree	Name of discipline	Department, institution and country	Month and year started	Month and year awarded/expected
Bachelor's	Honors Applied Mathematics	Science, Faculty of Alberta, CANADA	9 / 2011	4 / 2015
Master's	Applied Mathematics	Mathematics and Statistics McGill, CANADA	9 / 2015	Transferred to Ph.D.
Doctorate	Applied Mathematics	Mathematics and Statistics McGill, CANADA	1 / 2017	9 / 2020



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<b>ACADEMIC, RESEARCH AND OTHER RELEVANT WORK EXPERIENCE</b>			
Position held and nature of work (begin with current) Full Time - Part Time	Organization and department	Supervisor	Period (mm/yyyy-mm/yyyy)
Dynamical Systems Modeller - Full Time Quantitative systems biology scientist in the internal medicine research unit.	Pfizer Inc Internal Medicine Research Unit	Dr. Cynthia J. Musante	6/2017 - 9/2017
Teaching Assistant - Part Time Teaching assistant for Calculus for Management, Calculus II	McGill University Mathematics and Statistics	Dr. Sidney Trudeau	9/2016 - 12/2017
Research Assistant - Full Time NSERC USRA research assistant	University of Alberta Mathematics and Statistics	Dr. Hassan Safouhi	5/2015 - 9/2015
Research Assistant - Full Time NSERC USRA research assistant	University of Alberta Mathematics and Statistics	Dr. Hassan Safouhi	5/2014 - 9/2014
Teaching Assistant - Part Time Teaching assistant for Statistics I, Calculus I,II and linear algebra I courses.	University of Alberta Campus Saint-Jean	Dr. Sarah Pelletier	1/2013 - 4/2015

Personal identification no. (PIN)

**Valid** 439027

Family name, given name and initial(s) of applicant

Cassidy, Tyler TT

**AWARD APPLIED FOR**

Type of award

Postgraduate Scholarships - PGS D

Proposed starting date of award

2018/09

Proposed degree program  
(e.g. Masters, Doctorate)

Doctorate

Proposed field of study/research

APPLIED MATHEMATICS

Research subject code

2950

Title of proposed research

Quantitative Systems Biology Approach to Immunohematopoietic Regulation

List ten (10) key words that describe your proposed research.

Quantitative systems biology, Mathematical physiology, Mathematical modelling, Computational biology, Mechanistic modelling, Immune Response Model,

**PROPOSED LOCATION(S) OF TENURE (in order of preference)**

Institution/organization	Department	Program of study	Proposed supervisor
McGill,	Mathematics and Statistics	Applied Mathematics	Antony Humphries, Michael Mackey

Are any of your proposed programs of study:

Clinically-oriented?  Yes NoJoint programs with a professional degree (e.g., MD/PhD)?  Yes Yes No**SECTION TO BE COMPLETED BY PGS APPLICANTS ONLY**

Indicate the total number of months of graduate studies (master's and doctoral) you have completed as of December 31 of the year of application in the natural sciences and engineering.

25 months of full-time studies0 months of part-time studiesIndicate the number of months of studies you have completed, as of December 31 of the year of application, **in the program for which you are requesting funding.**5 months of full-time studies0 months of part-time studies

Indicate if you are attending university at the time of application.

 Attending full time Attending part time Not attending

If you are offered an award, do you plan to take it up at a foreign university?

 Yes NoIf you answered yes to the previous question, do you still want to be considered for an Alexander Graham Bell Canada Graduate Scholarship which is **tenable only in Canada?** Yes No



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<b>SCHOLARSHIPS AND OTHER AWARDS OFFERED (start with most recent and include NSERC awards)</b>					
Name of Award	Value (CDN\$)	Level Institutional, Provincial, National, International	Type Academic, Research, Leadership, Communication	Location of tenure	Period held (yyyy/mm - yyyy/mm)
CRM Applied Math	5,000	Provincial	Research	McGill University	2017/09 - 2018/09
CAMBAM Fellowship	5,000	Provincial	Research	McGill University	2017/09 - 2018/09
Graduate Excellence Fellowship	5,812	Institutional	Academic	McGill University	2017/09 - 2018/09
CRM Applied Math	5,000	Provincial	Research	McGill University	2016/09 - 2017/09
Graduate Excellence Fellowship	6,866	Institutional	Academic	McGill University	2016/09 - 2017/09
CAMBAM Fellowship	7,500	Provincial	Research	McGill University	2016/09 - 2017/09
James Lougheed Award of	15,000	Provincial	Research	McGill University	2015/09 - 2016/09
NSERC USRA	7,500	Institutional	Research	University of Alberta	2015/05 - 2015/09
NSERC USRA	7,500	Institutional	Research	University of Alberta	2014/05 - 2014/09

Type of Award <b>PGS D</b>	Personal identification no.(PIN) <b>Valid 439027</b>	Family name, given name and initial(s) of applicant <b>Cassidy, Tyler TT</b>
<b>THESIS COMPLETED OR IN PROGRESS</b>		
1. Degree <b>Doctor of Philosophy</b>	Supervisor <b>Antony Humphries and Michael Mackey</b>	Date degree requirements completed <b>12/2019</b>
Title of thesis <b>A Systems Biology Study of Immunity to Quantify Hematological Response</b>		
2. Degree	Supervisor	Date degree requirements completed <b>01/2018</b>
Title of thesis		
<b>SUMMARY OF THESIS MOST RECENTLY COMPLETED OR IN PROGRESS</b>		
<p>Do not reproduce abstract of thesis.</p> <p>In this work, I extend the quantitative system biology model of granulopoiesis derived by Craig et al. to include the immune response to bacterial infection. The physiology based model includes neutrophil recruitment to infected tissue, monocyte production, recruitment to infected tissue and differentiation into macrophages. The model reproduces the different time scales of immune response and can be used to understand local and systemic infection and to explore the pathology of chronic inflammation.</p>		

The hematopoietic (blood production) system is comprised of cells with lifespans varying from hours to years. The system creates new blood cells at a rate that reproduces the human body's own weight roughly every 10 years [1]. Homeostasis is maintained via complicated chains of positive and negative feedback loops that involve a legion of cytokines and hormones. Beyond maintaining homeostasis, the hematopoietic system must also respond to emergencies including blood loss and fungal, viral, and bacterial infections.

Bacterial infections lead to interactions between the immune and the hematopoietic systems. These interactions are controlled by a number of crosstalk relationships inherent to the immunohematological axis. Critical cytokines, including granulocyte colony stimulating factor (G-CSF), link the immune and hematopoietic systems and are amplified during infection. G-CSF induces white blood cell production and increases the release of mature white blood cells into the bloodstream [2]. Following resolution of the infection, the immune system decreases G-CSF production to signal the hematopoietic system to arrest the emergency production of white blood cells and return blood concentrations to homeostatic levels.

Experimental work has primarily focused on pairwise interactions between the cytokines governing immunohematologic interactions, as the number of cytokines involved in hematopoiesis renders systems level experiments intractable. Therefore, mathematical modelling offers an unparalleled opportunity for systems level characterization of the immunohematological response and to understand how systemic defects result in acute inflammation.

Previous mathematical models have focused on characterizing the interaction between immune and bacterial cells [4, 5]. Existing work accurately reproduces murine data by accounting for the staggered arrival of neutrophils and macrophages without explicitly modelling the complex physiology leading to immune cell production. However, without explicitly accounting for the precise bacterial-immune cell interactions that engender increased cytokine production and the physiological differences in neutrophil and macrophage production that account for their staggered arrival, the potential of mathematical models in immunoregulation remains unfulfilled.

My doctoral research will focus on mathematically modelling the immunohematopoietic response to bacterial infection. I will extend the mechanistic state dependent delay differential equation (DDE) model of granulopoiesis given by Craig et al. [3] to include the innate immunohematopoietic pathways.

I will construct a mechanistic mathematical model of bacterial infection by including bacteria proliferation and phagocyte mediated death. As G-CSF production is dependent on the severity of infection, I will introduce a bacteria-dependent G-CSF production rate. Linking G-CSF production to bacterial load will quantify how bacteria inoculum initiates immune response. I will derive a delay differential equation for monocyte production using partial differential equations, and I will model the evolution of circulating monocytes into macrophages during immune stress. Finally, I will use infection driven cytokine production to govern immune cell migration into infected tissues. The accumulation of immune and dead cells in infected tissue will then be used as a cipher for immune mediated inflammation.

To calibrate the resulting mathematical model, I will use sophisticated parameter estimation techniques and global sensitivity analysis. Analysis of the resulting model will combine techniques from infinite dimension dynamical systems and numerical analysis.

Extending Craig et al.'s mechanistic model of granulopoiesis [3] to include the innate immune response will allow for quantification of the interaction between the immune and hematopoietic systems. The goal of my work is to elucidate the mechanisms and interactions that characterize the immunohematopoietic chain. Beyond acute infection, this work is critical to understanding the mechanisms behind immunoregulation and how flaws in the immunohematopoietic chain lead to chronic inflammatory diseases, like rheumatoid arthritis and atherosclerosis, and will improve both clinical and experimental investigations of acute infections and chronic inflammation.

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[1] Dale, D.C. and Mackey, M.C. (2015) *Bull. Math. Biol.* **77**(5), 739–757.

[2] Roberts, A. (2005) *Growth Factors* **23**(1), 33–41.

[3] Craig, M., Humphries, A., and Mackey, M. (2016) *Bull. Math. Biol.* **78**(12), 2304–2357

[4] Smith, A., McCullers, J., and Adler, F. (2011) *J. Theor. Biol.* **276**(1), 106–116.

[5] Schirm, S., Ahnert, P., Wienhold, S., Mueller-Redetzky, H., Nouailles-Kursar, G., Loeffler, M., Witzentrath, M., and Scholz, M. (2016) *PLoS One* **11**(5), 1–22.

## **I Contributions to research and development**

### **a) Articles published or accepted in peer-reviewed journals**

1) **Cassidy, T.**, Gaudreau, P. and Safouhi, H.,(2017) On the Computation of Eigenvalues of the Anharmonic Coulombic Potential, *Journal of Mathematical Chemistry*, DOI 10.1007/s10910-017-0801-5 (B.Sc work)

### **b) Articles submitted to peer-reviewed journals**

1) De Souza, D.C, Craig, M., **Cassidy, T.**, Li,J.,0 Nekka, F., Humphries, A.R., Transit and lifespan in neutrophil production: implications for drug intervention, *Journal of Pharmacokinetics and Pharmacodynamics*, JOPA-D-17-00062. Submitted 05/22/2017. (Ph.D Work)

### **d) Non-peer-reviewed contributions**

1) **Cassidy, T.\***, Humphries, A.R., and Mackey, M.C, Mathematical Modelling of Cyclic Neutropenia, CAMBAM Seminar (National Oral Presentation), 2017 (M.Sc work)

2) **Cassidy, T.\***, Humphries, A.R., and Mackey, M.C, Understanding, Treating and Avoiding Hematological Disease: Better Medicine Through Mathematics, SIAM Life Sciences Meeting 2016 (International Oral Presentation), 2016 (M.Sc Work)

3) **Cassidy, T.\***, Humphries, A.R., and Mackey, M.C, Mathematical Modelling of Cyclic Neutropenia, SIAM General Meeting 2016 (International Poster), 2016 (M.Sc work)

4) **Cassidy, T.**, Gaudreau, P., and Safouhi, H., The Use of the DESCIM to Produce Numerical Solutions to the Schrödinger Equation (National Oral Presentation), PIMS Young Researchers Conference, 2015 (B.Sc work)

5) **Cassidy,T.\*** and Safouhi, H. The Importance of Delays in a Mathematical Model of Cyclical Neutropenia (Institutional Oral Presentation), Apprentis Chercheurs- University of Alberta, 2015 (B.Sc work)

6) **Cassidy, T.**, Gaudreau, P., and Safouhi, H., Efficient Computation of Schrödinger Energy Eigenvalues for Potentials having  $n$ th Order Singularities (Technical Report), 2015, (B.Sc work)

### **f) Contributions resulting from your participation in industrially relevant RD activities**

1) **Cassidy,T.\***, Rieger, T. and Baraldi, R., Modelling Lipoprotein Dynamics during PCSK-9 Inhibition, Pfizer Internal Medicine Research Unit Quantitative Research Group Seminar (Institutional Oral Presentation and technical report), 2017

## **II Most significant contributions to research and development**

1) During my undergraduate, I worked as a research assistant in numerical analysis during two summers. I performed asymptotic analysis of solutions to the Schrödinger equation with a potential involving singularities of arbitrary order. I constructed an adaptation to the Sinc collocation method (SCM) that converges exponentially to the energy eigenvalues of the potential. I wrote MATLAB code implementing this numerical method. I introduced a numerical scaling technique that increases numerical stability of the algorithm and proved the existence of a maximal scaling factor that increases convergence speed. The combination of the scaling factor and the SCM is capable of finding arbitrarily high energy eigenvalues to machine precision. I co-wrote the article detailing the application of the algorithm to the Coulombic potential with Philippe Gaudreau as well as a technical report detailing the general case. The resulting article was published in the *Journal of Mathematical Chemistry* in 2017.

2) During my M.Sc, I utilized the delay differential equation model for neutrophil production presented by Craig et al. (Craig 2016) to model a dynamical haematological disease, cyclic neutropenia. I identified a Hopf-bifurcation in the model that reproduces the quantitative behaviour of the disease. Translating the position of the Hopf bifurcation to mechanistic changes confirms one of the two accepted explanation for the origins of cyclic neutropenia. Typical data from patients with cyclic neutropenia is sparse in time, highly oscillatory and therefore ill-suited to typical parameter fitting routines. I utilized Lomb



periodograms and sparse data analysis techniques to extract useful details from the data. I defined an error functional and parameter fitting technique that reproduces the defining characteristics of cyclic neutropenia independently of data quality. Utilizing the parameter fitting routine reproduces the dynamical behaviour of the disease and provides a mathematical model of the disease. I incorporated the standard treatment technique into the model. Successfully reproducing the impact of treatment raised possible avenues of further experimental work to optimize treatment strategies.

3) During a summer internship at Pfizer Inc. I extended a mechanistic model of circulating lipoprotein dynamics to include liver production and internalization of blood lipoproteins and triglycerides. The extension of the model included modelling the production of lipoprotein receptors on the liver and the receptor antagonist. This model was capable of reproducing healthy dynamics and pathologies resulting from mechanistic defects, including familial cholesterolaemia. Finally, we explicitly modelled a monoclonal antibody and statin treatment to understand the mechanism leading to their synergistic function and impact on familial cholesterolaemia. The model's predictions agreed with published clinical trials.

### **Part III Applicant's statement**

#### **Research experience**

During my B.Sc, I performed research in numerical analysis and learned to analyse solutions to differential equations via their asymptotic behaviour. This work led to multiple presentations in English and French and a publication in the *Journal of Mathematical Chemistry*.

My senior year project introduced me to delay differential equations and the theory regarding their stability by studying the importance of delays in a mathematical model of hematopoiesis. I attended two summer schools in mathematical biology where I learned to apply mathematical methods to biological problems.

My graduate research has involved sparse data analysis techniques, analytic and numerical methods for dynamical systems for infinite dimensional systems and mechanistic modelling of the immune system using delay differential equations. During a research internship at Pfizer, I developed a physiology based model of a monoclonal antibody. The model reproduced published antibody mono- and combination therapy data. I performed global sensitivity analysis to understand interpatient treatment response variability.

#### **Relevant activities**

**Teaching:** I was the teaching assistant for *Calcul élémentaire I* (2013), *Calcul élémentaire* (2013,2014), *Calcul élémentaire II* (2013,2014), *Algebre linéaire I* (2012,2013,2014) and *Statistiques I* (2012) at the University of Alberta. These tutorials were given in French. At McGill, I was the teaching assistant for Calculus for Management and Calculus II. My responsibilities include preparing and leading tutorial sessions, giving office hours and marking exams and assignments.

**Organizational:** I organized a series of undergraduate research seminars at the University of Alberta-Campus Saint Jean. Undergraduate researchers presented their work to a multi-disciplinary audience during lunch. The seminar series included professional development presentations and social activities to promote interdisciplinary communication.

At McGill, I organize a Montreal wide student's only computational biology seminar. The seminar is geared towards a scientific audience and involves presentations from a variety of quantitative biology sub-fields. We host weekly research presentations and social events.

**Volunteer:** I was elected to the McGill mathematics graduate student government (2017). I was a governor of the Francophone Sport Association of Alberta (2012-2014). I was a *Chef de Mission* at the provincial francophone games (2012-2014) and an track and field coach at the Canadian Francophone games (2014,2017). I organized training sessions and seasonal training plans for blind athletes through Alberta Sport and Recreation for the Blind (2013-2015). I was a member of the Canadian Ski Patrol(2014-2015).