# Van-Giang Trinh: Curriculum Vitae

## 1 Personal

• Name: Van-Giang Trinh (Trinh Văn Giang in Vietnamese)

• Address: 10 Rue Granon, 13004, Marseille, France

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• Citizenship: Vietnamese



## 2 Employment History and Education

## 2.1 Employment History

- Postdoc, Laboratoire d'Informatique & Systèmes, Aix-Marseille University
   04/2022–current
- Researcher, School of Information Science, Japan Advanced Institute of Science and Technology 12/2021–03/2022
- Research Student, School of Information Science, Japan Advanced Institute of Science and Technology
   10/2017–09/2018
- Research Assistant, Faculty of Computer Science and Engineering, Ho Chi Minh City University of Technology
   06/2016–06/2017
- Internship, School of Information Science, Japan Advanced Institute of Science and Technology 10/2015–11/2015

#### 2.2 Education

 Ph.D., Information Science, Japan Advanced Institute of Science and Technology, Japan 10/2018– 12/2021

Supervisor: Kunihiko Hiraishi

- M.S., Computer Science, Ho Chi Minh City University of Technology, Vietnam 08/2014–04/2017
   Supervisor: Quan Thanh Tho
- B.S., Computer Science, Ho Chi Minh City University of Technology, Vietnam 09/2009–04/2014
- High School, Le Loi High School, Tho Xuan, Thanh Hoa, Vietnam

09/2006-05/2009

## 3 Research

#### 3.1 Research Interests

My research interests include theoretical computer science, artificial intelligence, and computational systems biology. In particular, I focus on Boolean networks, Petri nets, answer set programming, and their applications to modeling, analysis, and control of biological systems.

## 3.2 Awards

- 1. Outstanding Performance Award, Japan Advanced Institute of Science and Technology 12/2021
- Japanese Government (Monbukagakusho: MEXT) Scholarship, Japan Advanced Institute of Science and Technology
   10/2017–09/2021
- 3. JAIST President Award, Japan Advanced Institute of Science and Technology 09/2019

#### 3.3 Publications

#### 3.3.1 In preparation

- 1. **Van-Giang Trinh**, Belaid Benhamou, & Koichi Kobayashi. (2023). Attractor-based framework for scalable synchronization of temporal Boolean networks. *Journal*. (in preparation)
- 2. Van-Giang Trinh, Belaid Benhamou, Samuel Pastva, Jordan C Rozum, & Kyu Hyong Park. (2023). Positive and negative feedback loops enable fast state space mapping and control. *Journal*. (in preparation)
- 3. **Van-Giang Trinh**, Belaid Benhamou, & Sylvain Sene. (2024). Feedback loops and reduced dynamics in random order asynchoronous Boolean networks. In *Conference*. (in preparation)

#### 3.3.2 Submitted

- 1. **Van-Giang Trinh**, Belaid Benhamou, & Loïc Paulevé. (2023). mpbn: a simple tool for efficient edition and analysis of elementary properties of Boolean networks. *Journal*. (major revision)
- 2. Van-Giang Trinh, Belaid Benhamou, Tarek Khaled, & Kunihiko Hiraishi. (2023). Computing attractors of large-scale asynchronous Boolean networks using minimal trap spaces. *Journal*. (under review)
- 3. Van-Giang Trinh, Belaid Benhamou, Samuel Pastva, & Sylvain Soliman. (2024). Scalable enumeration of trap spaces in Boolean networks via answer set programming. In *Conference*. (under review)

#### 3.3.3 Peer-Reviewed Journals (top venues in bold)

- Van-Giang Trinh, Belaid Benhamou, & Sylvain Soliman. (2023b, September). Trap spaces of Boolean networks are conflict-free siphons of their Petri net encoding. *Theoretical Computer Science*, 971, 114073. (IF 1.1, CORE Rank A) <a href="https://doi.org/10.1016/j.tcs.2023.114073">https://doi.org/10.1016/j.tcs.2023.114073</a>
- Tarek Khaled, Belaid Benhamou, & Van-Giang Trinh. (2023, July). Using answer set programming to deal with Boolean networks and attractor computation: Application to gene regulatory networks of cells. Annals of Mathematics and Artificial Intelligence, 1–38. (IF 1.2) https://doi.org/10.1007/s10472-023-09886-7
- 3. **Van-Giang Trinh**, Belaid Benhamou, Thomas Henzinger, & Samuel Pastva. (2023a, June). Trap spaces of multi-valued networks: Definition, computation, and applications. *Oxford Bioinformatics*, 39(Supplement\_1), i513–i522. (IF 5.8, oral presentation at ISMB/ECCB 2023) https://doi.org/10.1093/bioinformatics/btad262
- Van-Giang Trinh, & Kunihiko Hiraishi. (2020b). On attractor detection and optimal control of deterministic generalized asynchronous random Boolean networks. *IEEE/ACM Transactions on Computational Biology and Bioinformatics*, 19(3), 1794–1806. (IF 4.5) https://doi.org/10.1109/ TCBB.2020.3043785

- Van-Giang Trinh, Tatsuya Akutsu, & Kunihiko Hiraishi. (2020). An FVS-based approach to attractor detection in asynchronous random Boolean networks. *IEEE/ACM Transactions on Computational Biology and Bioinformatics*, 19(2), 806–818. (IF 4.5) https://doi.org/10.1109/TCBB.2020.3028862
- Van-Giang Trinh, & Kunihiko Hiraishi. (2020c). A study on attractors of generalized asynchronous random Boolean networks. *IEICE TRANSACTIONS on Fundamentals of Electronics, Communications and Computer Sciences*, 103(8), 987–994. (IF 0.5) https://doi.org/10.1587/transfun.2019EAP1163
- 7. **Van-Giang Trinh**, Le Ngoc Kim Khanh, Bang Ngoc Bao Tam, Tram Loi Quan, Bui Hoai Thang, & Quan Thanh Tho. (2016). Modelling and congestion detection of wireless sensor networks: A concurrent-based approach using coloured Petri nets. *International Journal of Applied Information Systems*, 11(7), 1–9. https://doi.org/10.5120/ijais2016451629
- 8. Pham Hong Long, **Van-Giang Trinh**, Dinh Hoang Mai, Mai Phuong Nam, Quan Thanh Tho, & Ngo Quang Hung. (2014). Assisting students in finding bugs and their locations in programming solutions. *International Journal of Quality Assurance in Engineering and Technology Education* (*IJQAETE*), 3(2), 12–27. https://doi.org/10.4018/ijqaete.2014040102

#### 3.3.4 Peer-Reviewed Conferences (top venues in bold)

- 1. **Van-Giang Trinh**, Belaid Benhamou, & Sylvain Soliman. (2023a, August). Efficient enumeration of fixed points in complex Boolean networks using answer set programming. In *International Conference on Principles and Practice of Constraint Programming* (pp. 35:1–35:19). (main track, the primary conference of the Association for Constraint Programming, AR = 40%, CORE Rank A) https://doi.org/10.4230/LIPIcs.CP.2023.35
- Van-Giang Trinh, Belaid Benhamou, Thomas Henzinger, & Samuel Pastva. (2023b, July). Trap spaces of multi-valued networks: Definition, computation, and applications. In *The 31st Annual Intelligent Systems For Molecular Biology and the 22nd Annual European Conference on Computational Biology*. Oxford University Press. (proceedings track, the flagship meeting of the International Society for Computational Biology, AR 17.9%, CORE Rank A) https://doi.org/10.1093/bioinformatics/btad262
- 3. Van-Giang Trinh, Kunihiko Hiraishi, & Belaid Benhamou. (2022, August). Computing attractors of large-scale asynchronous Boolean networks using minimal trap spaces. In *ACM International Conference on Bioinformatics, Computational Biology and Health Informatics* (pp. 1–10). ACM. (the flagship conference of the ACM SIGBio, AR 29%) https://doi.org/10.1145/3535508.3545520
- 4. **Van-Giang Trinh**, Belaid Benhamou, Kunihiko Hiraishi, & Sylvain Soliman. (2022, August). Minimal trap spaces of logical models are maximal siphons of their Petri net encoding. In *International Conference on Computational Methods in Systems Biology* (pp. 158–176). Springer. (AR 65%) https://doi.org/10.1007/978-3-031-15034-0\_8
- Van-Giang Trinh, & Kunihiko Hiraishi. (2021, October). An improved method for finding attractors of large-scale asynchronous Boolean networks. In *IEEE International Conference on Computational Intelligence in Bioinformatics and Computational Biology* (pp. 1–9). IEEE. (AR 51%) https://doi.org/10.1109/cibcb49929.2021.9562947
- 6. Van-Giang Trinh, & Kunihiko Hiraishi. (2020a, December). An efficient method for approximating attractors in large-scale asynchronous Boolean models. In *IEEE International Conference on Bioinformatics and Biomedicine* (pp. 1820–1826). IEEE. (workshop paper, AR 60%) https://doi.org/10.1109/bibm49941.2020.9313230

- 7. **Van-Giang Trinh**, & Kunihiko Hiraishi. (2019, June). Algorithms for finding attractors of generalized asynchronous random Boolean networks. In *12th Asian Control Conference* (pp. 67–72). IEEE. Retrieved from http://ieeexplore.ieee.org/document/8765169 (AR 73%)
- 8. Le Ngoc Kim Khanh, **Van-Giang Trinh**, Bui Hoai Thang, & Quan Thanh Tho. (2017, April). Probabilistic modelling for congestion detection on wireless sensor networks. In *International Conference on Control, Decision and Information Technologies* (pp. 0190–0195). IEEE. (AR 48%) https://doi.org/10.1109/CoDIT.2017.8102589
- 9. **Van-Giang Trinh**, Nguyen Duc Khoan, Nguyen Duy Khuong, Vu Phu Thuc, & Quan Thanh Tho. (2016, September). Fast-and-Fit: An intelligent auto-pricing system for airlines travel agencies. In *SAI Intelligent Systems Conference* (pp. 853–865). Springer. https://doi.org/10.1007/978-3-319 -56994-9 58
- 10. **Van-Giang Trinh**, Kunihiko Hiraishi, & Quan Thanh Tho. (2016, July). Modeling and analysing Boolean networks by coloured Petri nets. *IEICE Proceedings Series*, 61(4447). https://doi.org/10.34385/proc.61.4447
- 11. Bao Trung Pham Duy, **Van-Giang Trinh**, Le Dinh Thuan, & Quan Thanh Tho. (2015, October). Reusing symbolic observation graph for efficient model checking. In *International Conference on Knowledge and Systems Engineering* (pp. 250–255). IEEE. https://doi.org/10.1109/kse.2015.44

#### 3.3.5 Other Publications

1. F. Kordon, H. Garavel, L. M. Hillah, F. Hulin-Hubard and G. Chiardo, A. Hamez, L. Jezequel, A. Miner, J. Meijer, E. Paviot-Adet, D. Racordon, C. Rodriguez, C. Rohr, J. Srba, Y. Thierry-Mieg, Van-Giang Trinh, & K. Wolf. (2016, June). *Complete Results for the 2016 Edition of the Model Checking Contest*. Retrieved from https://mcc.lip6.fr/2016/index.php

#### 3.4 Talks and Seminars

- 1. LIRICA Seminar (LIS), Efficient Enumeration of Fixed Points in Complex Boolean Networks using Answer Set Programming (20/11/2023)
- 2. Demi Journées du Pôle Calcul on Artificial Intelligence (LIS), Efficient Enumeration of Fixed Points in Complex Boolean Networks using Answer Set Programming (15/06/2023)
- 3. CANA Seminar (LIS), Trap Spaces of Boolean Networks are Conflict-Free Siphons of Their Petri Net Encoding (30/05/2023)
- 4. Journées BioLogique BIOSS/CAVIAR, An Approach Based on ASP and Petri Nets for the Calculation of Attractors in Boolean Networks (25/05/2023)
- 5. Journées Scientifiques du LIS, Efficient Enumeration of Minimal Trap Spaces in Large-Scale Boolean Networks of Gene Networks (23/05/2023)
- 6. LIRICA Seminar (LIS), Minimal Trap Spaces of Boolean Models are Maximal Siphons of Their Petri Net Encoding (17/10/2022)
- 7. LIRICA Seminar (LIS), An FVS-based Approach to Attractor Detection in Asynchronous Boolean Networks (28/06/2021)

# 4 Teaching

#### 4.1 Courses

- Exercises on Graph Theory (graduate course, teaching assistant)
  - Japan Advanced Institute of Science and Technology: 06/2021–08/2021

- Functional Programming (graduate course, teaching assistant)
  - Japan Advanced Institute of Science and Technology: 10/2020-12/2020
- Introduction to Computer Programming (undergraduate course, visiting lecturer)
  - Ho Chi Minh City University of Technology: 01/2017-06/2017
- Principles of Programming Languages (undergraduate course, teaching assistant)
  - Ho Chi Minh City University of Technology: 09/2013–01/2014, 09/2014–01/2015, 09/2015– 01/2016
- Object-Oriented Programming (undergraduate course, teaching assistant)
  - Ho Chi Minh City University of Technology: 09/2015-01/2016
- Principles of Programming Languages (undergraduate course, visiting lecturer)
  - Ho Chi Minh City University of Natural Resources and Environment: 01/2015-05/2015
- Data Structures and Algorithms (undergraduate course, teaching assistant)
  - Ho Chi Minh City University of Technology: 09/2012–01/2013, 01/2015–05/2015
- Programming Fundamentals (undergraduate course, teaching assistant)
  - Ho Chi Minh City University of Technology: 01/2015-05/2015

## 5 Professional Service

### 5.1 Journals Reviewers

- SIAM Journal on Applied Dynamical Systems, 1 paper, 2023
- IEEE/ACM Transactions on Computational Biology and Bioinformatics, 1 paper, 2023
- IEEE/ACM Transactions on Computational Biology and Bioinformatics, 1 paper, 2022

## 6 References

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