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Verification, Model Checking, and Abstract Interpretation

18th International Conference, VMCAI 2017 Paris, France, January 15–17, 2017 Proceedings

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Preface

This volume contains the papers presented at VMCAI 2017, the 18th International Conference on Verification, Model Checking, and Abstract Interpretation, held during January 15–17, 2017, in Paris, France, co-located with POPL 2017 (the annual ACM SIGPLAN/SIGACT Symposium on Principles of Programming Languages). Previous meetings were held in Port Jefferson (1997), Pisa (1998), Venice (2002), New York (2003), Venice (2004), Paris (2005), Charleston (2006), Nice (2007), San Francisco (2008), Savannah (2009), Madrid (2010), Austin (2011), Philadelphia (2012), Rome (2013), San Diego (2014), Mumbai (2015), and St. Petersburg, Florida (2016).

VMCAI provides a forum for researchers from the communities of verification, model checking, and abstract interpretation, facilitating interaction, cross-fertilization, and advancement of hybrid methods that combine these and related areas. VMCAI topics include: program verification, model checking, abstract interpretation and abstract domains, program synthesis, static analysis, type systems, deductive methods, program certification, debugging techniques, program transformation, optimization, hybrid and cyber-physical systems.

This year the conference attracted 60 submissions. Each submission was reviewed by at least three Program Committee members. The committee decided to accept 27 papers. The principal selection criteria were relevance, quality, and originality. We are pleased to include in the proceedings the contributions of three invited keynote speakers: Ernie Cohen (Amazon Web Services), Pascal Cuoq (Trust in Soft), and Jasmin Fisher (Microsoft Research). We warmly thank them for their participation and for their contributions.

We would like also to thank the members of the Program Committee and the external reviewers for their excellent work. We also thanks the members of the Steering Committee, and in particular Andreas Podelski and Lenore Zuck, for their helpful advice, assistance, and support. We also thank Laure Gonnord for her invaluable help in all aspects related to the organization of the conference. We thank Annabel Satin for the help in coordinating the events co-located with POPL 2017, and we thank the POPL 2017 Organizing Committee for providing all the logistics for organizing VMCAI. We are also indebted to EasyChair for providing us with an excellent conference management system.

Finally, we would like to thank our generous sponsors: AdaCore, Amazon Web Services, Facebook, and Microsoft Research.

December 2016

Ahmed Bouajjani
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Abstracts of Invited Talks

Bringing LTL Model Checking to Biologists

Zara Ahmed[1], David Benque[2], Sergey Berezin[3],
Anna Caroline E. Dahl[4], Jasmin Fisher[1,5], Benjamin A. Hall[6],
Samin Ishtiaq[1], Jay Nanavati[1], Nir Piterman[7],
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Abstract. The BioModelAnalyzer (BMA) is a web based tool for the development of discrete models of biological systems. Through a graphical user interface, it allows rapid development of complex models of gene and protein interaction networks and stability analysis without requiring users to be proficient computer programmers. Whilst stability is a useful specification for testing many systems, testing temporal specifications in BMA presently requires the user to perform simulations. Here we describe the LTL module, which includes a graphical and natural language interfaces to testing LTL queries. The graphical interface allows for graphical construction of the queries and presents results visually in keeping with the current style of BMA. The Natural language interface complements the graphical interface by allowing a gentler introduction to formal logic and exposing educational resources.

Verified Concurrent Code: Tricks of the Trade

Ernie Cohen

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Abstract. Modular code verification, suitably extended with shared atomic objects, supports a number of useful verification idioms and semantic models, without further logical extension.

Keywords: Real-time · Hybrid systems · Probability · Stopping failures · Weak memory · Cryptography · Ownership · Permissions · Simulation · Knowledge · Behavioral polymorphism · Device drivers · Concurrent data structures · Transactions · Linearizability · Deductive verification · VCC

Detecting Strict Aliasing Violations in the Wild

Pascal Cuoq[1], Loïc Runarvot[1], and Alexander Cherepanov[2,3]

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Abstract. Type-based alias analyses allow C compilers to infer that memory locations of distinct types do not alias. Idiomatic reliance on pointers on the one hand, and separate compilation on the other hand, together make it impossible to get this aliasing information any other way. As a consequence, most modern optimizing C compilers implement some sort of type-based alias analysis. Unfortunately, pointer conversions, another pervasive idiom to achieve code reuse in C, can interact badly with type-based alias analyses. This article investigate the fine line between the allowable uses of low-level constructs (pointer conversions, unions) that should never cause the predictions of a standard-compliant type-based alias analysis to be wrong, and the dangerous uses that can result in bugs in the generated binary. A sound and precise analyzer for strict aliasing violations is briefly described.

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