## **Binding-Time Analysis for MINIKANREN**

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MINIKANREN<sup>1</sup> is a family of domain-specific languages for pure *relational* programming which core can be implemented in a few lines of the host language [2]. A program in MINIKANREN can be run in different directions: given some of its arguments, it computes all the possible values of the rest of the arguments. When the relation is run with only the last argument unknown, we say it is run in the *forward direction*, otherwise we call it running in the *backward direction*. A single relational specification creates a multitude of directions, each solving distinctive problems.

The search employed in MINIKANREN is complete, so all possible answers will be computed eventually, albeit it may take a long time. In reality, the running time depends on the direction and is highly unpredictable. This is where the promise of MINIKANREN falls short: one has to write the relational specification with the specific direction in mind for it to be efficient. There are several attempts to tackle this problem in an automatic fashion: one way is to use a *divergence test* to stop execution of definitely diverging computations [4], the other is to employ *specialization* [3], while the one we have been working recently is to converse a relational specification into a function which runs fast in the given direction.

Given a relational specification equipped with a certain direction, the functional conversion constructs a function in which unifications are mapped either into a pattern matching or are used in letbindings to compute the intermediate results. Disjunctions give rise to branches in pattern matchings within a function definition, while conjunctions mean sequential computations. The nondeterministic computations are modeled with lists. The most nontrivial part of the functional conversion is to determine the order in which to bind each variable within the function body.

It has been shown in [3] that an online conjunctive partial deduction is capable of improving running time of a program in a given direction, but we believe that offline specialization may provide better results. Offline specializers employ a static analysis before the specialization step. The most prominent of them is the *binding-time analysis*. It is used to determine which data is available statically, and which — dynamically. Having this annotations, a specialization algorithm can ignore dynamic data and only symbolically execute a program with respect to the static data. This usually leads to more precise and powerful specialization.

We believe that a variation of a binding-time analysis can serve as a solution to figuring out the order of variable bindings necessary for functional conversion as well as be a part of an offline specializer. The goal of a binding-time analysis is to identify at which step of a computation each variable gets its value. This process is called annotations: each variable gets an annotation from the domain of *binding-time*. Existing binding-time analyses for pure PROLOG [1] and MERCURY [6] only distinguish between static and dynamic variables, which does not help to work out the order. Natural numbers serve as annotations in [5] and are better suited to our problem.

The binding-time analysis for MINIKANREN should annotate a given goal as well as determine suitable directions for all relation calls involved in the result computation. We start by marking some free

<sup>&</sup>lt;sup>1</sup>MINIKANREN language web site: http://minikanren.org

variables within the goal with 0 which denotes they are the *input* or *static* variables, while all others are annotated with *undef*. This is the way we specify a direction for the goal.

We assume that all goals are in *normal form*: a goal is a disjunction of conjunctions of either relation calls or unifications; all free variables are brought up into scope by the *fresh* operator on the top level. Relation calls and constructors can only have variables as their arguments. Unification should have a single variable on the left-hand side with the exception of unifying two constructors of arity 0.

$$G = \underline{fresh} \ V_1 \dots V_n \bigvee \bigwedge (\underline{call} \ R^k \ V_1 \dots V_k \ | \ V \equiv T \ | \ \underline{cons} \ C^0 \equiv \underline{cons} \ C^0)$$

The algorithm is to execute the following steps until a fixpoint is reached. Since disjuncts do not influence each other, we treat every disjunct in isolation. A conjunct is either a unification or a relation call. At each step, a unification suitable for annotation is selected. If there are no unifications to annotate, then we select a relation call. Whenever a new annotation is assigned to a variable, all other uses of the same variable within the conjunction get the same annotation.

Unifications are annotated if there is enough information. There are several possible cases described below. We write annotations in superscripts;  $t[x_0, \dots, x_k]$  denotes a term with free variables  $x_0, \dots, x_k$ .

- The unification of an unannotated variable with a term in which all free variables are annotated with numbers:  $x^{undef} \equiv t[y_0^{i_0}, \dots, y_k^{i_k}]$ . In this case we annotate x with  $1 + max\{i_0, \dots, i_k\}$ .
- The unification of an annotated variable with a term, in which some variables are not annotated:  $x^n \equiv t[y_0^{i_0}, \dots, y_k^{i_k}]$ . Here all unannotated variables of the term are annotated with 1+n.

If a unification does not conform to any of these cases, it should not be annotated at the current step. Relation calls are only annotated when all unifications have been considered. A relation call in which all variables have not been annotated (are marked with *unde f*) as well as those already annotated with some positive numbers do not need to be considered. Thus relation calls in which only some of the variables are annotated with positive numbers are to be annotated.

We say that two calls to the same relation have a consistent direction if their free variables which are annotated with numbers are ordered in the same way. More formally: relation call  $R^n v_1^{i_1} \dots v_k^{i_k}$  has a direction consistent with  $R^n u_1^{j_1} \dots u_k^{j_k}$ , if all variables with numerical annotations are ordered in the same way:  $v_{l_1}^{i_{l_1}} \leq \cdots \leq v_{l_m}^{i_{l_m}}$  and  $u_{l_1}^{j_{l_1}} \leq \cdots \leq u_{l_m}^{j_{l_m}}$ .

If the current conjunct has a consistent direction with some relation call annotated before, then we

use that relation call to annotate the current.

Only a relation call with the direction which has not been encountered before should be annotated. It is done by considering its body: variables annotated within the call keep their annotations in the body, which is annotated by the described algorithm.

This algorithm terminates since there are finitely many relation calls to consider, and each relation call has finitely many possible annotations.

## References

- [1] Stephen-John Craig, John P Gallagher, Michael Leuschel & Kim S Henriksen (2004): Fully automatic binding-time analysis for Prolog. In: International Symposium on Logic-Based Program Synthesis and Transformation, Springer, pp. 53-68.
- [2] Jason Hemann & Daniel P. Friedman (2013): uKanren: A Minimal Functional Core for Relational Programming.

- [3] Petr Lozov, Ekaterina Verbitskaia & Dmitry Boulytchev (2019): *Relational Interpreters for Search Problems*. In: *Relational Programming Workshop*, p. 43.
- [4] Dmitri Rozplokhas & Dmitri Boulytchev (2018): *Improving Refutational Completeness of Relational Search via Divergence Test*. In: Proceedings of the 20th International Symposium on Principles and Practice of Declarative Programming, pp. 1–13.
- [5] Peter Thiemann (1997): A Unified Framework for Binding-Time Analysis. In: TAPSOFT.
- [6] Wim Vanhoof, Maurice Bruynooghe & Michael Leuschel (2004): *Binding-time analysis for Mercury*. In: *Program Development in Computational Logic*, Springer, pp. 189–232.