E-matching \subseteq **Relational Join** Simpler, faster, and optimal

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MAPPING E-GRAPHS TO RELATIONS



An example e-graph. Each solid box denotes an e-node and each dashed box denotes an e-class, which represents a set of equivalent terms. Labels at top-left corner denotes the e-class id. Represented terms include f(1,g(1)), f(1,g(2)), f(2,g(1)), etc. $(O(N^2)$ total).

eclass-id	child₁	child₂
cf	1	cg
cf	2	cg
cf	Ν	cg

eclass-id	child₁
c <i>g</i>	1
c <i>g</i>	2
c <i>g</i>	N

 R_f : relation representing *f* (left). R_q : relation representing g (right).

E-GRAPH & E-MATCHING

E-graph is a data structure that efficiently represents sets of congruent terms. E-graph has wide applications in automated-theorem proving and program optimization.

E-matching is a fundamental query of e-graphs that searches for a pattern modulo congruence. Existing backtracking-based e-matching algorithms rely on depth-first search over the e-graph and fail to take equality constraints over the pattern into consideration during query planning.

CQS & GENERIC JOIN

• **Conjunctive query** (CQ) is a restricted class of relational queries that only involve joins of relations.

Generic join is an algorithm proposed by Ngo et al. that computes CQs in worst-case optimal time with respect to the output size. > Has great performance especially when the CQ is complex (e.g., cyclic).

REDUCING E-MATCHING TO CONJUNCTIVE QUERIES

 $f(\boldsymbol{\alpha}, g(\boldsymbol{\alpha}))$

- An e-matching pattern that matches all expressions where
- 1. the 1st argument to f is g and
- 2. the 2^{nd} argument of f and the 1^{st} argument of g refer to the same eclass.



 $Q(root, \alpha) := R_f(root, \alpha, x), R_g(x, \alpha)$

The conjunctive guery derived from the pattern. Nested functions are flattened by introducing auxiliary variables (x).

Enumerated terms by backtracking-based e-matching ($O(N^2)$ many)

 $f(1,g(1))\checkmark$ f(2, g(1)) $f(2,g(2))\checkmark$ f(3, g(1))f(3, g(2)) $f(3,g(3))\checkmark$

Tuples visited by relational e-matching (O(N) many)

 $\begin{pmatrix} c_f, 1, c_g \end{pmatrix} \quad \begin{pmatrix} c_g, 1 \end{pmatrix} \checkmark \\ \begin{pmatrix} c_f, 2, c_a \end{pmatrix} \quad \begin{pmatrix} c_g, 2 \end{pmatrix} \checkmark$...

RELATIONAL E-MATCHING

• We propose **relational e-matching**, which reduces e-matching to CQs over a **relational representation** of e-graphs.

> The CQ form of e-matching fully exploits the equality constraints over the pattern, compared to existing backtracking-based algorithms where only the structural constraints are considered during query planning.

> To solve the complex CQs generated by relational e-matching, we use generic joins as our solver subroutine.

• Relational e-matching preserves the worst-case optimality of generic joins: Fix a pattern p, let M(p, E) be the set of substitutions yielded by e-matching on e-graph E with size n, relational e-matching runs in time $\tilde{O}(\max_E(|M(p,E)|)).$

BENCHMARK & RESULTS

We benchmarked with three representative e-matching queries, collected from the test suite for mathematical expressions of egg, a state-of-the-art e-graph framework. > On e-matching queries with equality constraints (the cyclic and the non-linear acyclic cases), relational ematching achieve **asymptotic speed-ups up to 426**× over the baseline e-matching algorithm by exploiting the equality constraints during query planning. On e-matching queries without equality constraints (the linear case), relational e-matching achieves similar performance as the baseline e-matching.

Speed-ups over backtracking-based e-matching algorithm

