

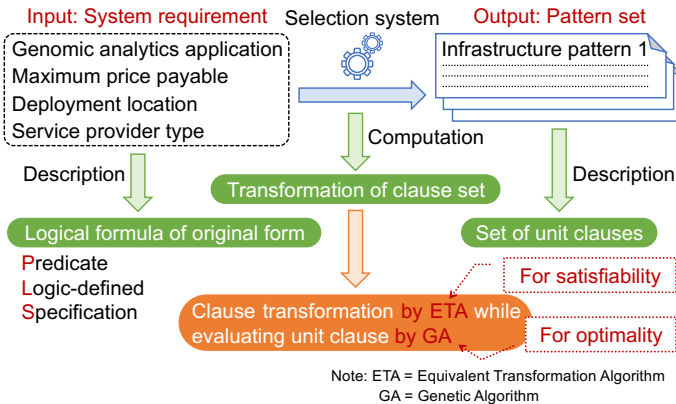
Cloud Resource Selection based on PLS Method for deploying Optimal Infrastructure for Genomic Analytics Application

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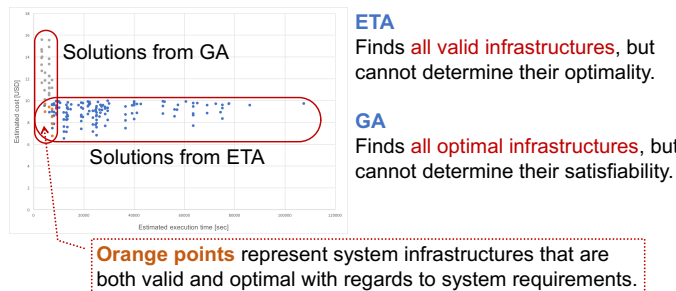
1. INTRODUCTION

We propose a method for determining cloud resources that concurrently meet **system infrastructure satisfiability and optimality requirements**.

Overview

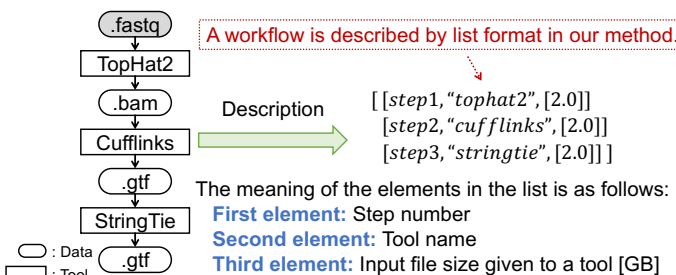


Importance of combining ETA and GA



2. GENOMIC ANALYTICS APPLICATION

A genomic analytics application is a **workflow composed of various tools** such as TopHat2, HISAT, and StringTie.



3. CLOUD RESOURCE SELECTION BASED ON PLS METHOD

The PLS method finds all feasible infrastructures that completely satisfy system requirements **by transforming clause sets**.

Cloud resource selection process

Step 1: Construct a PLS reflecting system requirements

A PLS has the following form:

$$\forall \bar{v} \{E_1 \wedge E_2 \wedge \dots \wedge E_n\},$$

where E is an atomic formula (atom), \bar{v} is a set of variables appearing in the formula.

Step 2: Generate a definite clause cl from the PLS

A definite clause cl has the following form:

$$ans(v_1, v_2, \dots, v_i) \leftarrow E_1, E_2, \dots, E_n,$$

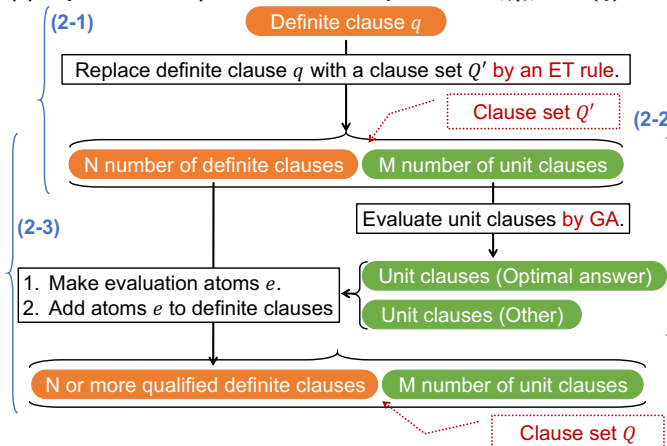
where the term of an ans atom is all variables in \bar{v} .

Step 3: Transform clause set \mathbb{D}

Clause set \mathbb{D} is transformed while preserving the declarative meaning of the initial state in accordance with the following procedure, where the **initial state is $\{cl\}$** .

(1) **Select a definite clause q from clause set \mathbb{D} , where $\mathbb{D} = \{q\} \cup D$.** A definite clause q is **randomly selected** from clause set \mathbb{D} . (NB: From the viewpoint of efficient computation, the definite clause selection method is important, but this method is left for future work.)

(2) **Replace clause q with a clause set Q , where $\mathcal{M}(\{q\}) = \mathcal{M}(Q)$.**



(3) **Make new clause set \mathbb{D} by joining clause sets D and Q .**

Termination condition of the transformation

- Clause set \mathbb{D} is a **unit clause set or an empty set**.
- A unit clause **determined to be optimal by GA** is obtained.

4. SYSTEM REQUIREMENTS AS INPUT DATA

To describe system requirements with respect to genomic analytics application deployment, **four (4) atoms** are utilized:

$$\forall \{*_structure\} \{Environment(*_structure, [[step1, "tophat2", [2.0]] [step2, "cufflinks", [2.0]] [step3, "stringtie", [2.0]]]) \wedge$$

$$Cost(*_structure, 10.0) \wedge$$

$$Location(*_structure, ["Virginia", "Tokyo"]) \wedge$$

$$Policy(*_structure, "public", "single") \}$$

- Environment atom:** Genomic analytics application to deploy
- Cost atom:** Maximum price payable
- Location atom:** Deployment location or region
- Policy atom:** Service type and number of cloud service providers

5. OPERATION EXPERIMENT OF PLS METHOD

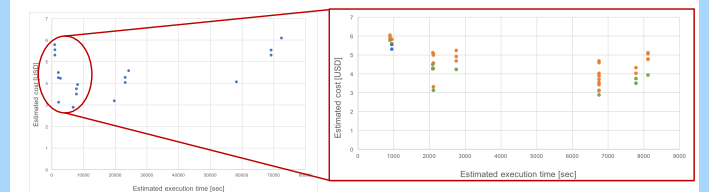
Purpose

Find optimal infrastructures that satisfy (1).

Computation

As a result of transformations of clause set \mathbb{D} , **sixty (60) definite clauses** and **nineteen (19) unit clauses** are obtained from the initial state.

In process (2-2) of Step 3, the unit clauses are evaluated by GA. **Green points** represent optimal solutions.



Example of an optimal solution

$$ans(["aws", "r4.16xlarge", "Virginia", [step1, "tophat2", [2.0], 892.49]], ["aws", "m4.xlarge", "Virginia", [step2, "cufflinks", [2.0], 1.0]], ["aws", "m4.xlarge", "Virginia", [step3, "stringtie", [2.0], 1.0]]) \leftarrow$$

6. CONCLUSION

- The proposed method makes it possible to find infrastructures that concurrently meet **system satisfiability and optimality requirements**.
- The proposed method can be extended to deploy other applications in addition to genomic analytics, **without modifying the resource selection algorithm**, simply by adding new atoms and their applicable ET rules.