Maximal Multi-layer Specification Synthesis

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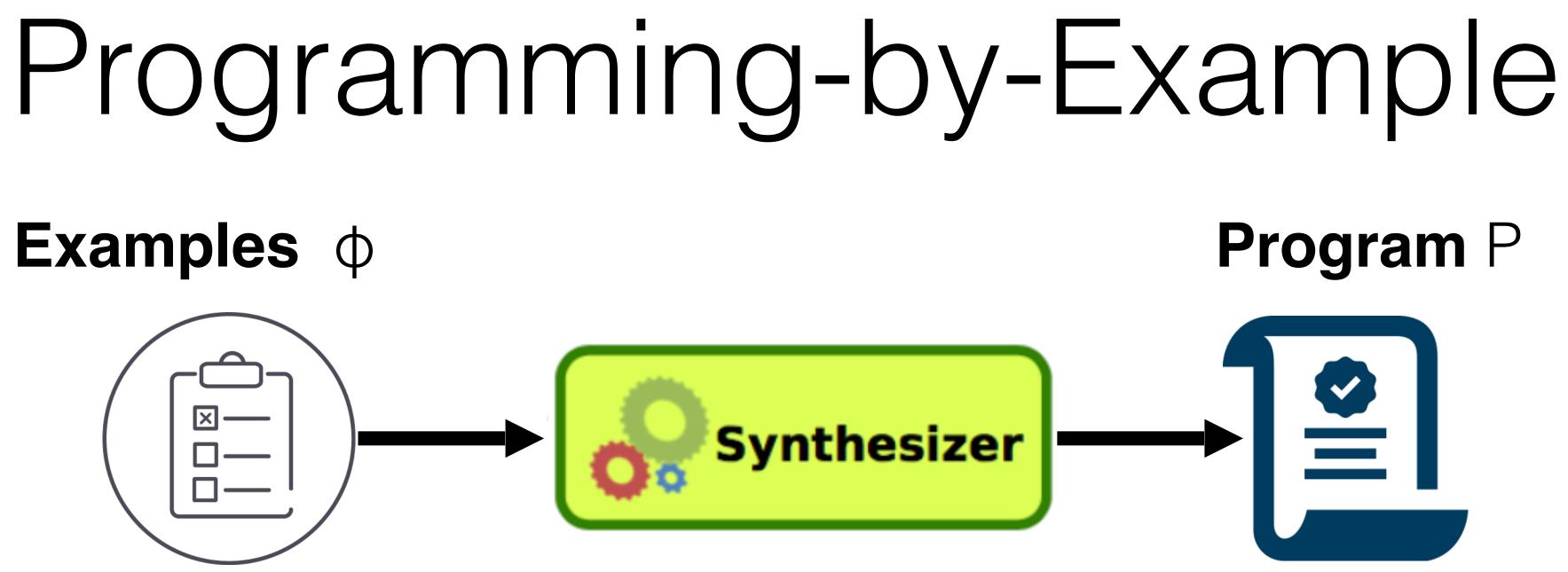
Specifications ϕ



$\exists P. \forall x. \phi(x, P(x))$

• Find a program P that for all inputs x meets the specification ϕ

Examples ϕ



• Find a program P that satisfies all input-output examples ϕ

Programming-by-Example

Input

Student	Grade	Score1	Score2	
Greg	A	75	76	
Greg	В	86	85	
Sally	А	85	86	
Sally	В	80	78	

Output

Student	B_Score1	B_Score2	A_Score1	A_Score
Greg	86	85	75	76
Sally	80	78	85	86

• Can we find a program that automatically **transforms tables** given input-output examples?



Programming-by-Example

Input

Student	Grade	Score1	Score2		Student	B_Score1	B_Score2	A_Score1	A_Score2
Greg	А	75	76		Greg	86	85	75	76
	D	96	0E		Sally	80	78	85	86
Greg	B	86	85	R	progra	m:			
Sally	А	85	86			r(input,S	core,Gr	ade,Score	e1,Score
Sally	В	80	78			(df1,AllSo read(df2,	•	•	•

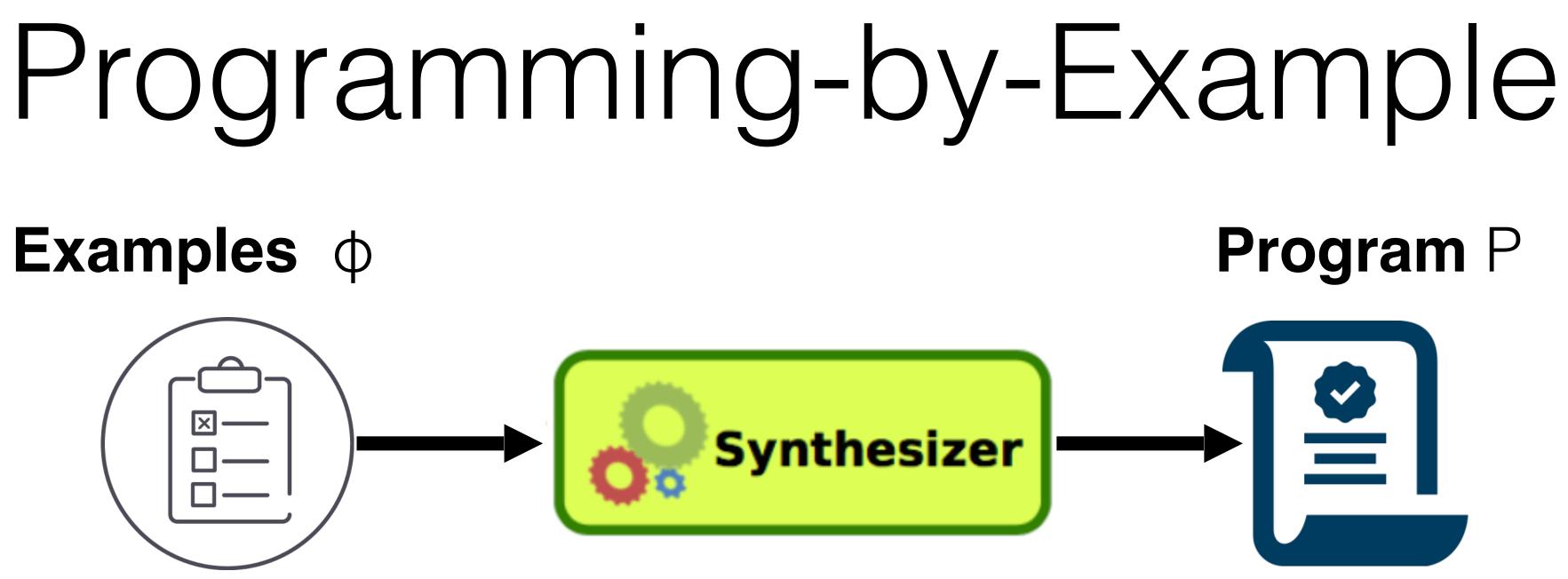
transformation tasks from examples. PLDI 2017

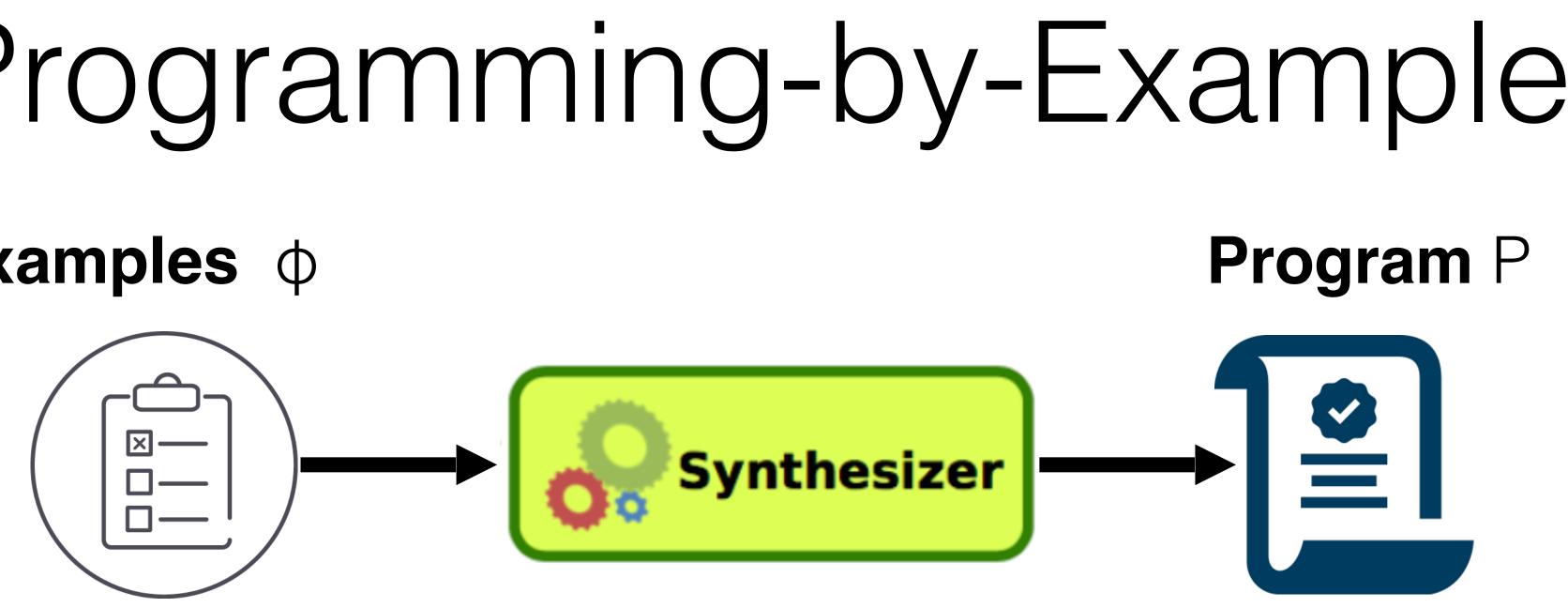
Output

Component-based synthesis of table consolidation and



e2)





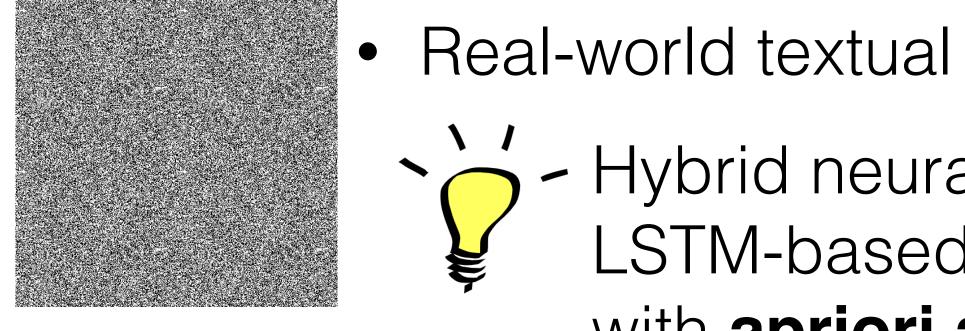


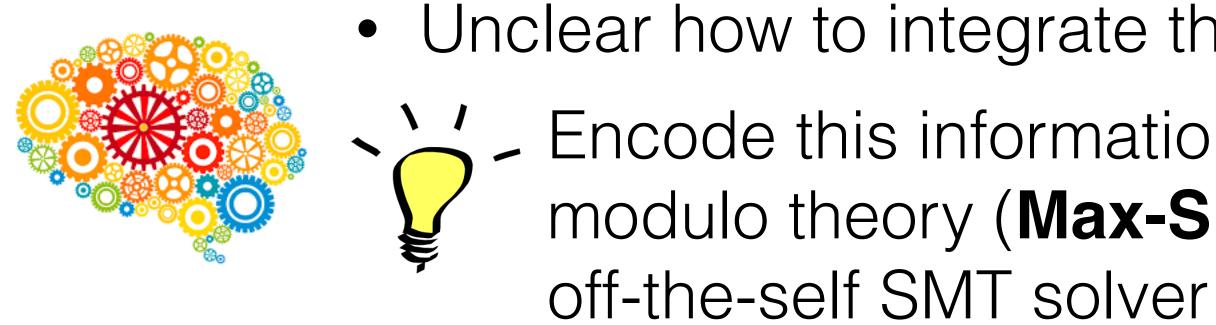


- Natural language
- Input-output examples

Capture the user intent with a combination of specifications:

Challenges





- Real-world textual information is noisy and ambiguous
 - Hybrid neural network architecture that combines
 LSTM-based sequence-to-sequence (seq2seq)
 with apriori algorithm for mining rules
- Unclear how to integrate this information with a synthesizer
 - Encode this information to maximum satisfiability modulo theory (Max-SMT) and solve it with an off-the-self SMT solver

Concrete

Input

Student	Grade	Score1	Score2	
Greg	А	75	76	
Greg	В	86	85	F
Sally	A	85	86	۲ ر
Sally	В	80	78	

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		(Dutput		
	Student	B_Score1	B_Score2	A_Score1	A_Score
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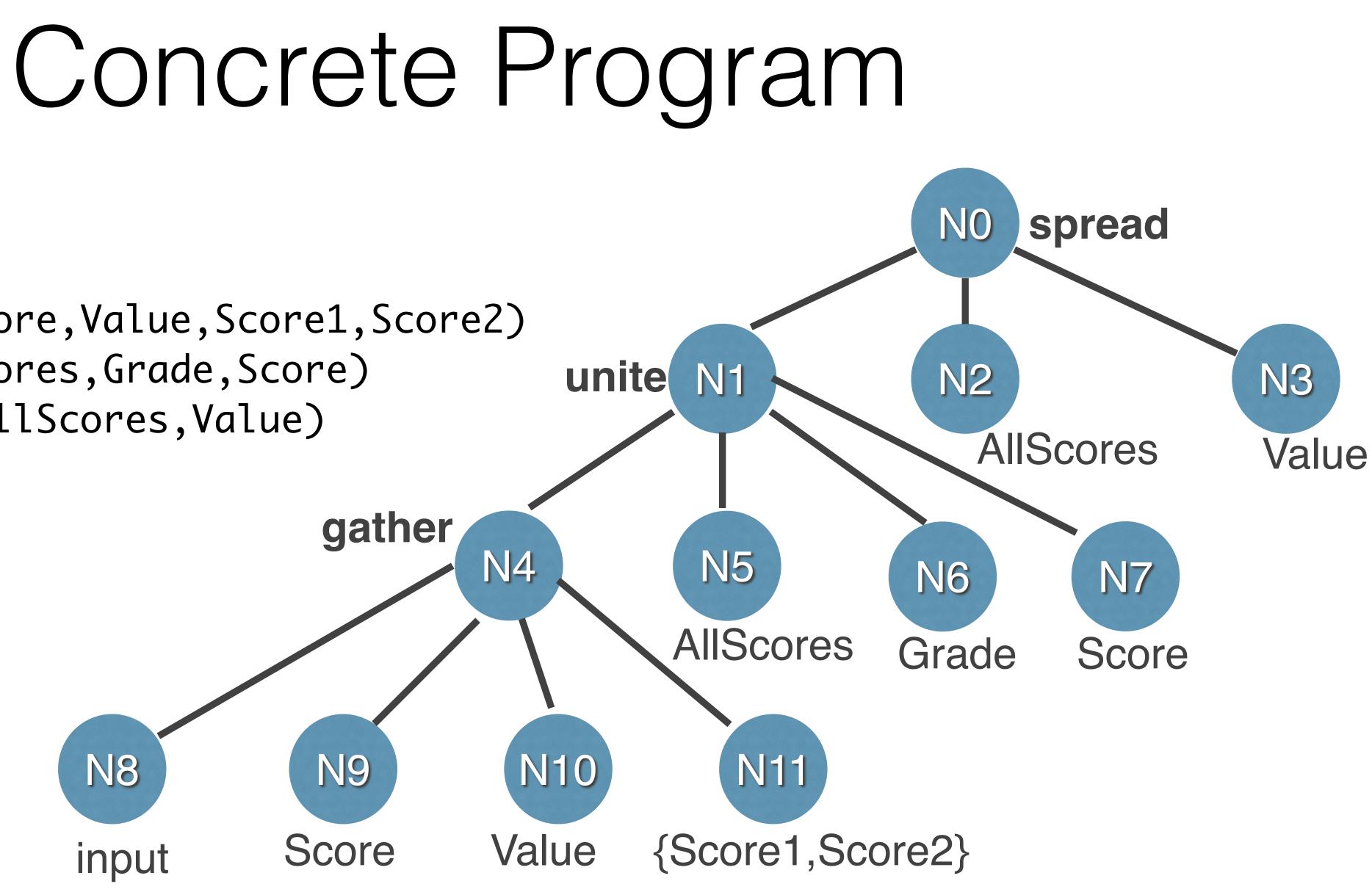
R program:

df1=gather(input,Score,Grade,Score1,Score2) df2=unite(df1,AllScores,Time,Score) output=spread(df2,AllScores,Grade)

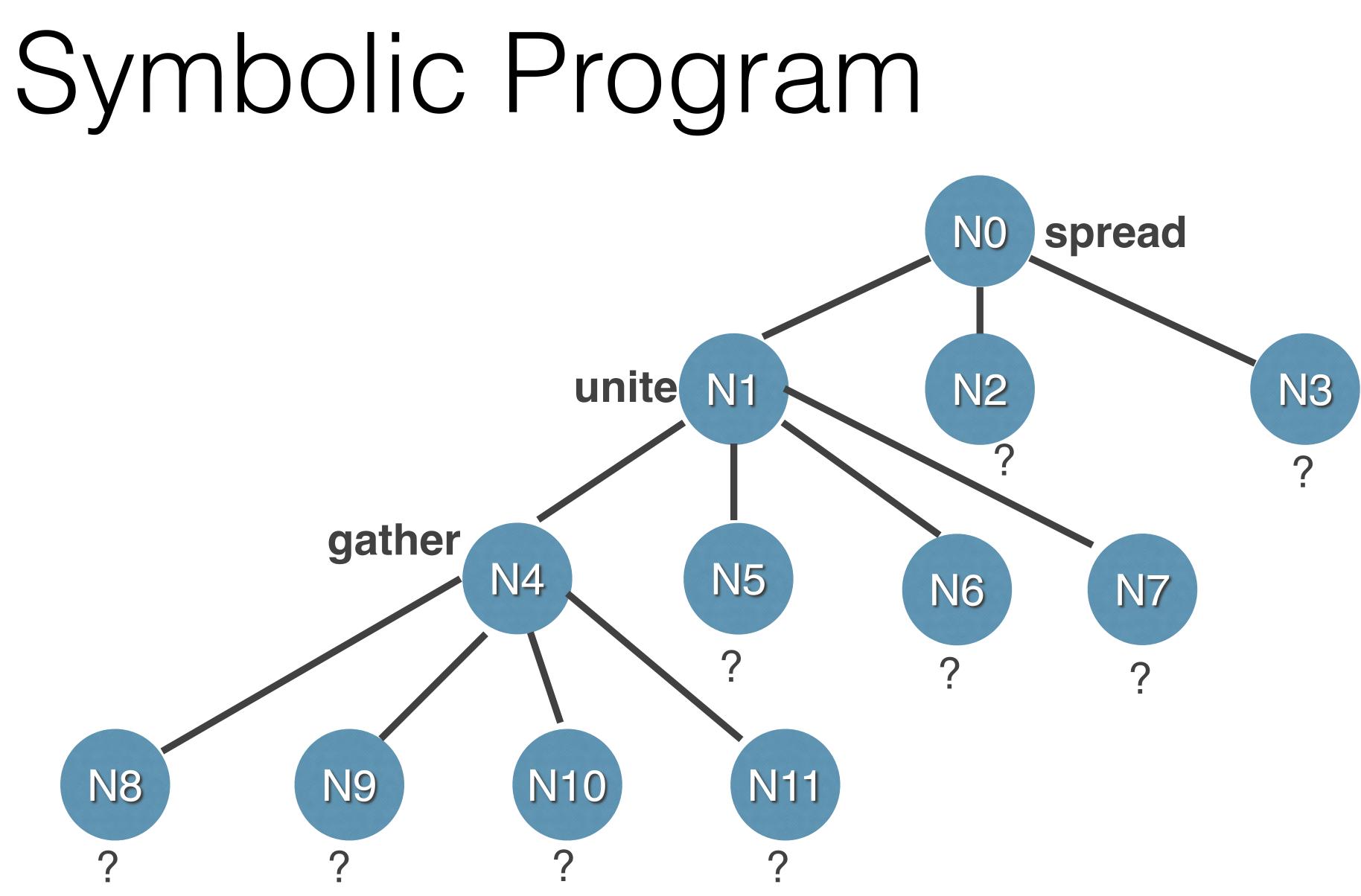


R program:

df1=gather(input,Score,Value,Score1,Score2) df2=unite(df1,AllScores,Grade,Score) output=spread(df2,AllScores,Value)



R program: df1=gather(??) df2=unite(??) output=spread(??)



r script to reshape and the columns within dataset
description
<pre> • • • • • • • • • • • • • • • • • • •</pre>
description
I need to reformal the data so that there is just one row per site wat i.e. in a given site name and data comparison with columns for the description refuse its species and

Problem Description

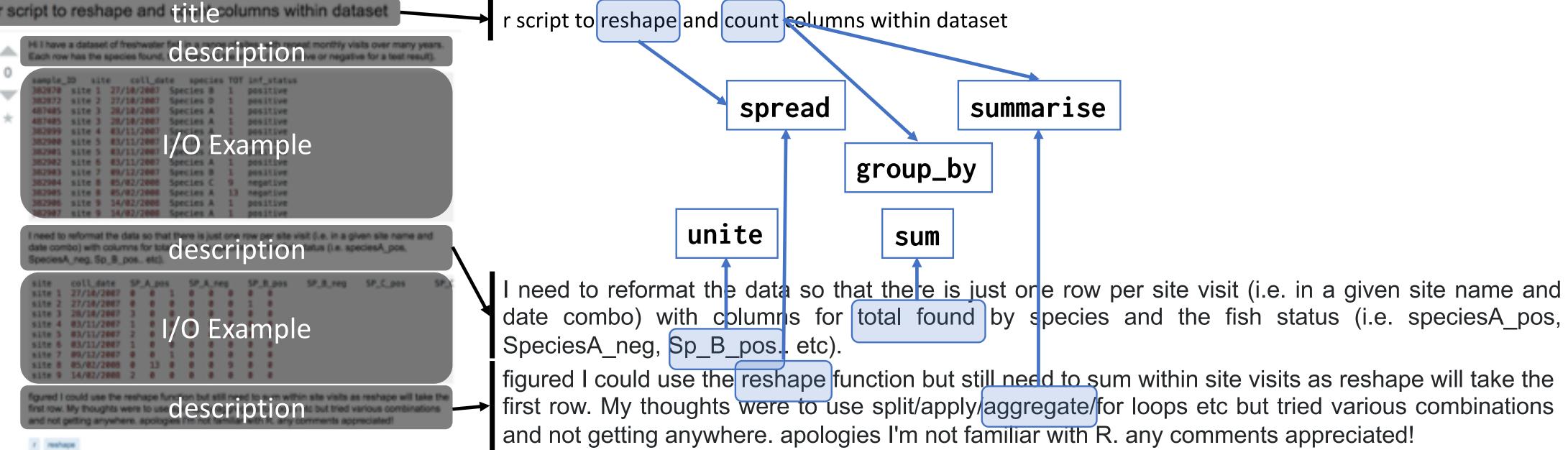
script to reshape and this columns within dataset description te coll_date species 12970 site 1 27/10/2007 Species 8 1 site 2 27/18/2007 Species 0 87485 site 3 28/18/2007 Species A 1 positive site 3 28/18/2007 Species A 1 positive 999 site 4 #3/11/3007 5 I/O Example 12983 site 7 89/12/2087 Species 8 2984 site 8 #5/82/2888 Seecies C site 8 #5/02/2008 Species A 13 negative ite 9 34/02/2008 Species A 1 positive description 8.pos SP_8_reg SP_C_pos I/O Example SpeciesA_neg, Sp_B_pos. etc). 118 6 03/11/2007 1 0 Lie 7 00/13/2007 0 Lte 8 05/02/2008 0 13 0 description r reshape

Problem Description

r script to reshape and count columns within dataset

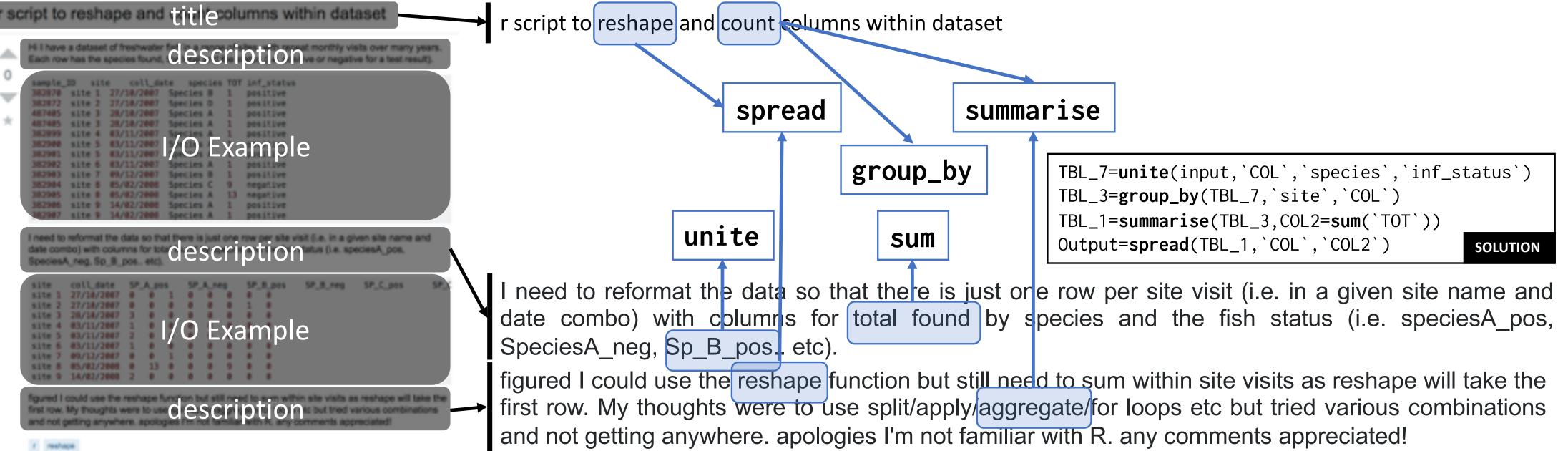
I need to reformat the data so that there is just one row per site visit (i.e. in a given site name and date combo) with columns for total found by species and the fish status (i.e. speciesA_pos,

figured I could use the reshape function but still need to sum within site visits as reshape will take the first row. My thoughts were to use split/apply/aggregate/for loops etc but tried various combinations and not getting anywhere. apologies I'm not familiar with R. any comments appreciated!

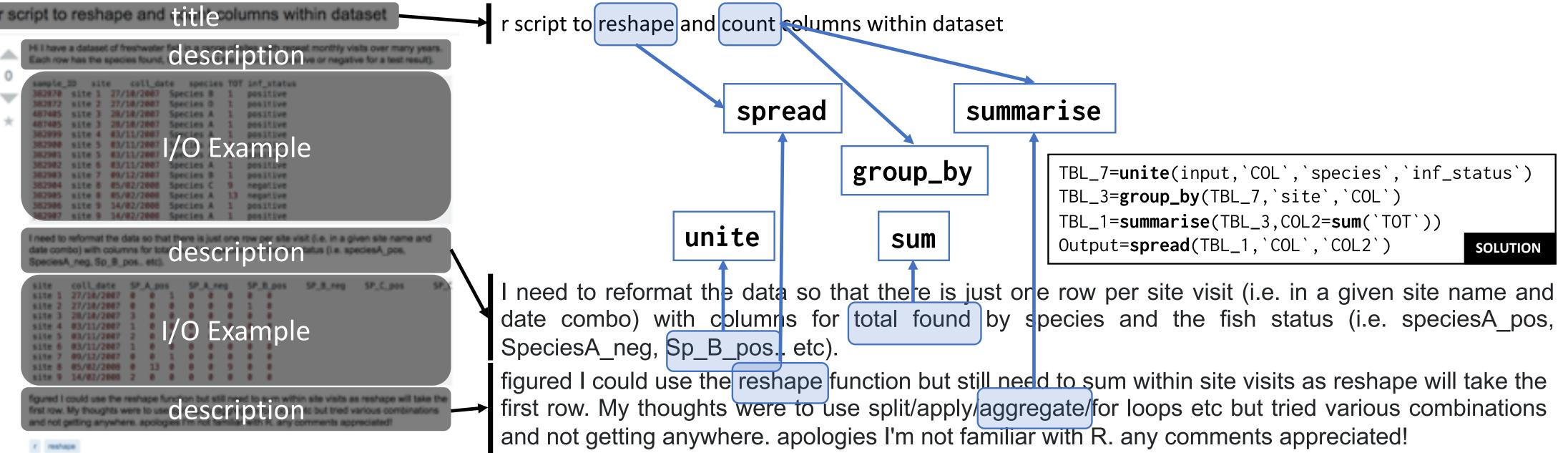


Problem Description

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Problem Description



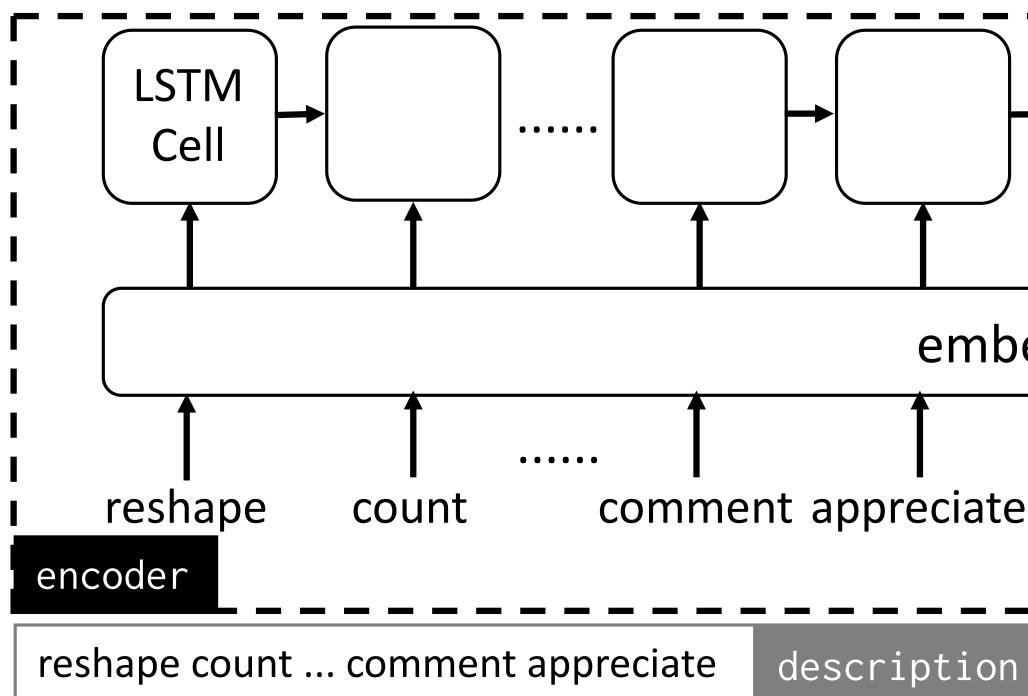
• To capture hints from natural language, we model description and symbolic programs relationships using seq2seq model

Problem Description

- Question-solution pair (D,S):
 - Question (D) is a user description composed by word tokens D = (d1, d)d2, ..., dn):
 - D = ("r", "script", "to", "reshape", "and", "count",...)
 - Solution (S) is a symbolic program composed by a sequence of functions S = (s1, s2, ..., sn)
 - S = ("unite", "group_by", "summarise", "spread",...)
 - seq2seq model is used to estimate the probability of P(SID)

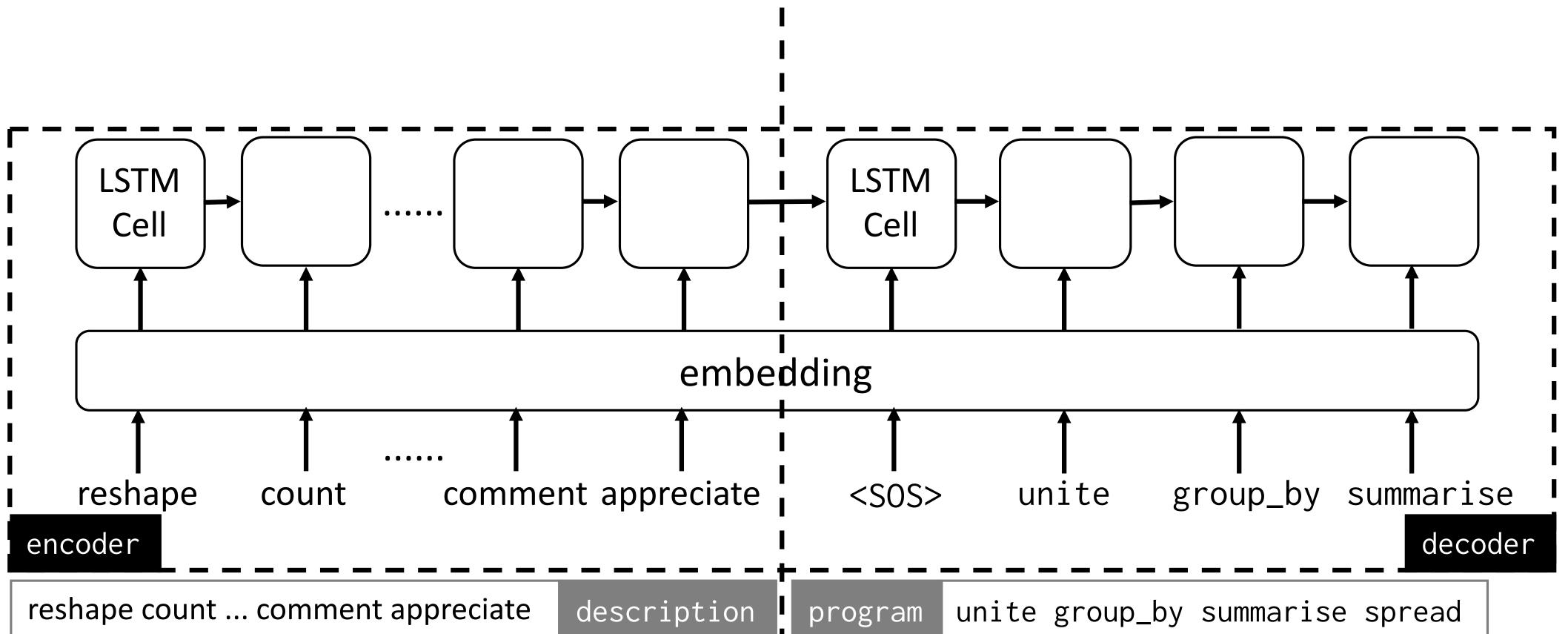
Neural

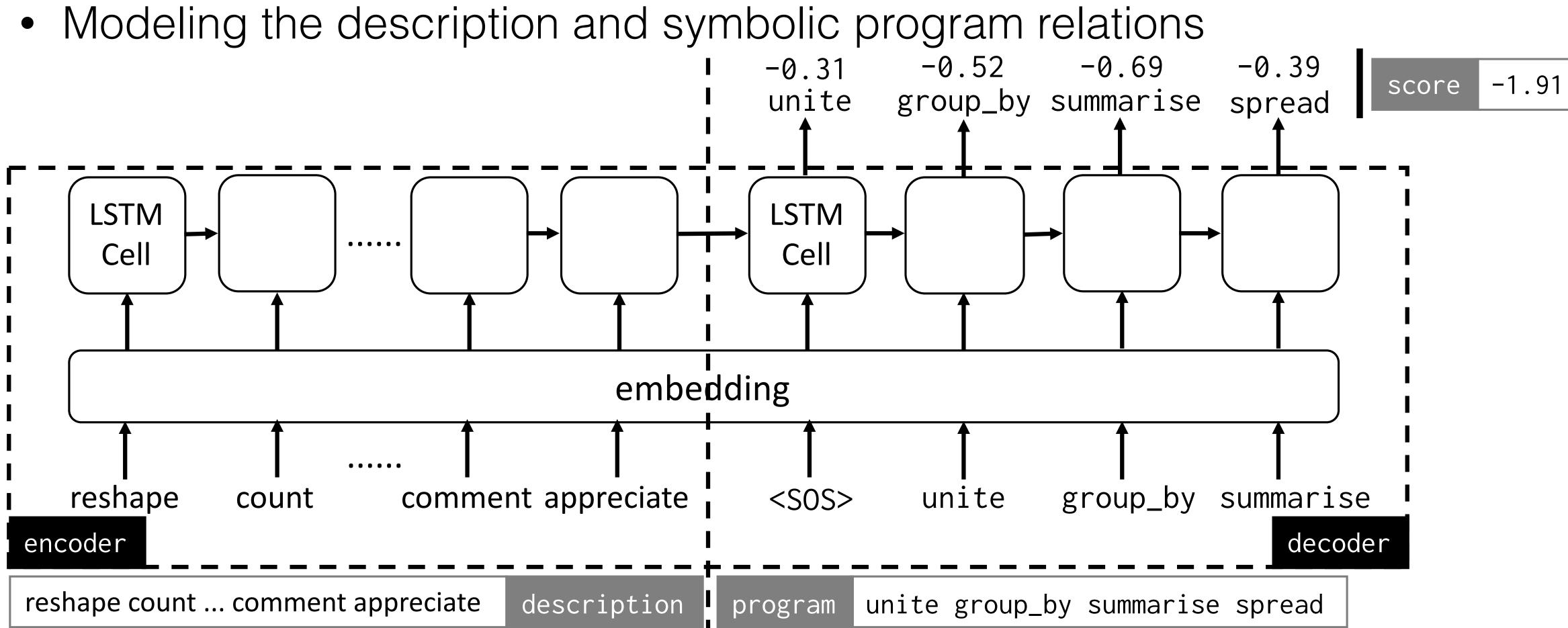
Modeling the description and system



Network	
ymbolic program relations	
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edding	

Modeling the description and symbolic program relations







- Use beam search to generate a ranked list (P, w):
 - P Symbolic Program
 - w likelihood of being part of the solution
- Score is computed by summing log likelihood in every step
- 1. {mutate, group_by, summarise, spread}(92) 2. {group_by, summarise, mutate, select}(91)
- **130.** {mutate, group_by, summarise, spread}(79)



. . .

. . .



Mining Association Rules

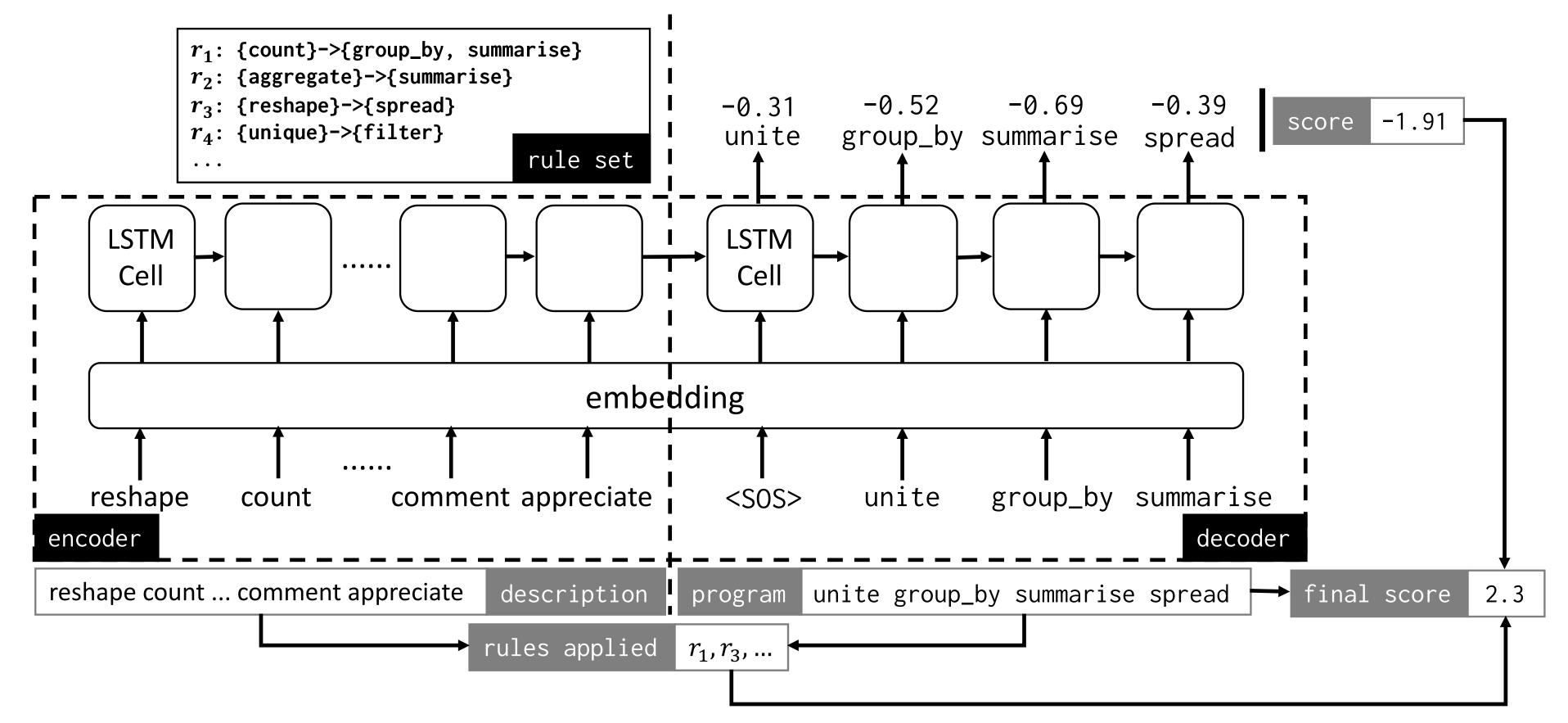
Association rules

. . .

. . .

- **X** => **Y** ; **X**: keywords ; **Y**: functions
- Unsupervised learning using the apriori algorithm
- Example of association rules: {"reshape", "count"} => {spread} Ranking further improves • {","reshape"} => {unite}
- 1. {unite, group_by, summarise, spread}(109)
- **30.** {mutate, group_by, summarise, spread}(94)

Hybrid Neural Network



• Use the learned association rule set to adjust the ranking scores on the fly



Program Synthesizers





Enumerative Search Constraint Solving





Stochastic Search

Most synthesizers do not combine different techniques in an **unified** way

Synthesis using Statistical Models

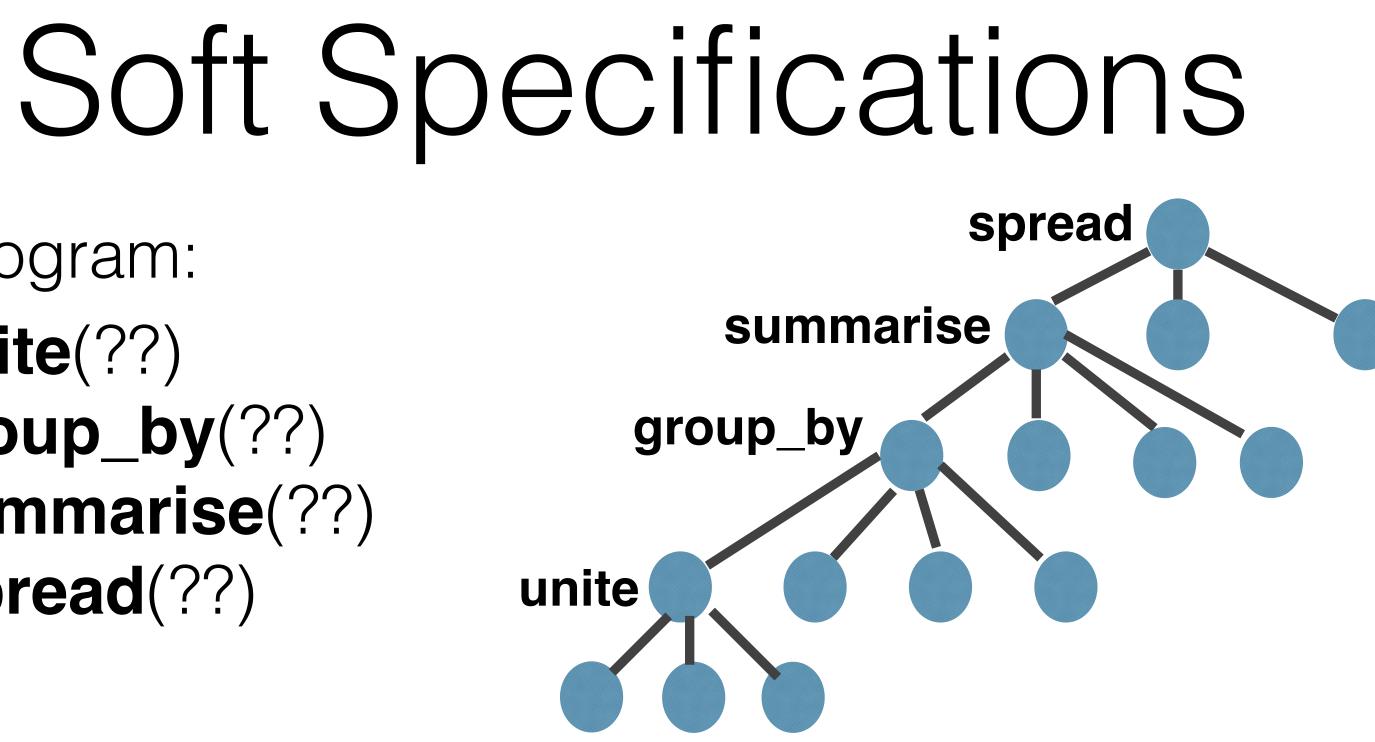
- Hard specifications (must be satisfied)
 - Input-output examples
 - Program type-checks
- **Soft specifications** (may be satisfied)
 - User preference in the form of natural language
- Satisfy all hard specifications while maximizing the sum of the weights of the **satisfied soft** specifications

Satisfiability Modulo Theories

- Satisfiability Modulo Theories (SMT) problem is a decision problem for formulas that are composed with multiple theories
- We can encode the enumeration of concrete programs into SMT using Linear Integer Arithmetic (LIA)
- Hard specifications:
 - 1. Assign functions to the root node that are consistent with input-output examples
 - 2. Only constants and inputs can be assigned to the leaf nodes 3. The concrete program is well formed, i.e. type-checks



- Symbolic program: $TBL_7 = unite(??)$ $TBL_3 = group_by(??)$ TBL_1 = summarise(??) Output = **spread**(??)
- Predicates:
 - (**occurs**(s), w): s occurs in the solution with likelihood w: \bullet
 - Example: (occurs(summarise), 109)
 - (hasChild(s, s'),w): s is a parent of s' in the solution with likelihood w: Example: (hasChild(group_by,unite), 109)



- **seq2seq** neural network:
 - PyTorch framework

 - Embedding layer:
 - Maps 25,004 words and 14 functions
 - To vectors of the dimension 256
- Association rule mining:
 - Efficient-Apriori package
 - Filter out association rules with low confidence

Setup

• Hyperparameters are obtained through a simple grid search

Data Collection and Preparation

- 80 benchmarks for data wrangling tasks using R libraries (*tidyr*, *dplyr*)
- Collected 20,640 pages from StackOverflow:
 - Testing benchmarks excluded
 - Each page contains a single question and multiple answers
- Removing duplicate and questions with no solutions: 16,459 question-solution pairs used to train seq2seq
- Extract descriptions from answers and their corresponding solutions: • 37,748 transactions as input of the Apriori algorithm

 - Learned 187 association rules

• Morpheus:

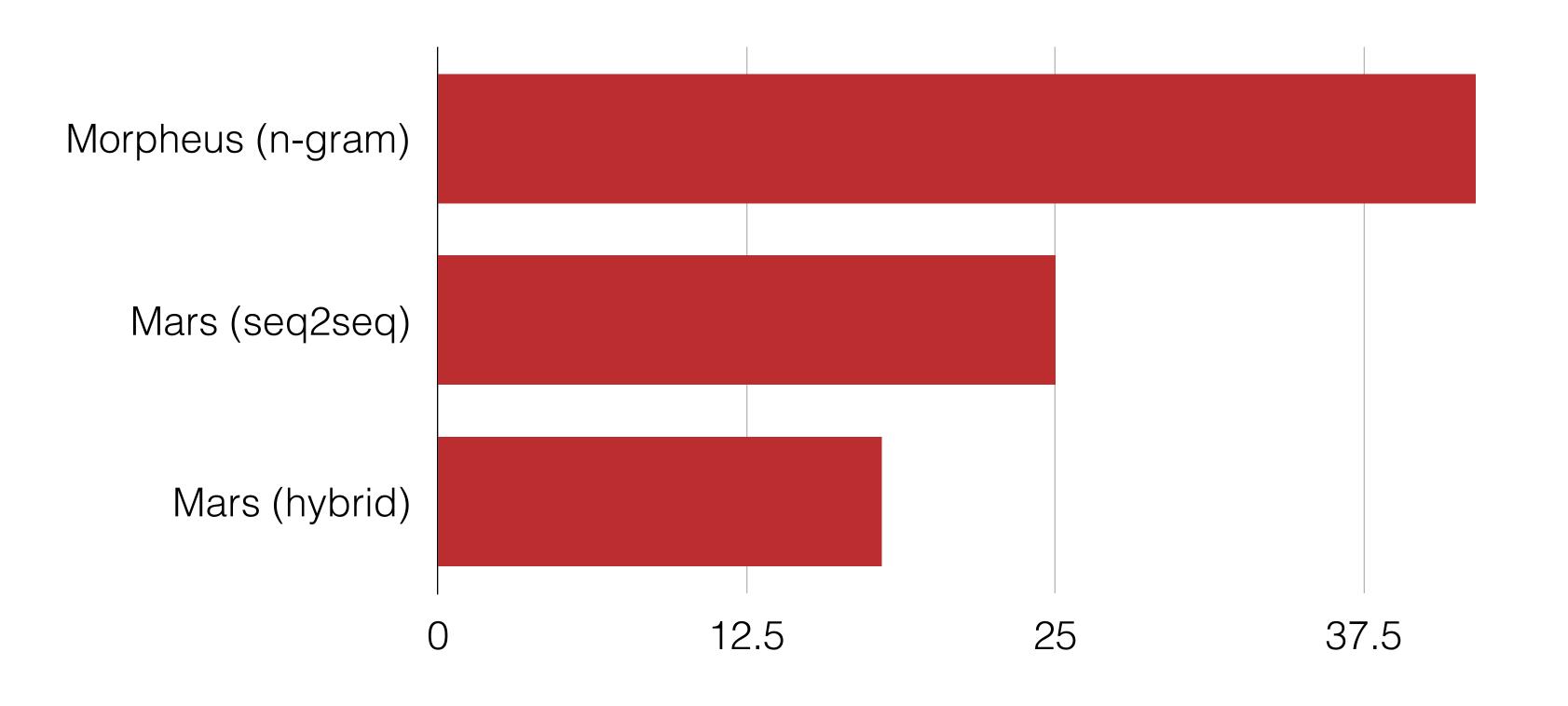
- n-gram model
- Mars:
 - Considers the user description for each task
 - seq2seq
 - Hybrid (seq2seq + apriori algorithm)

Evaluation — Data Wrangling

Global ranking that does not consider user description

Quality of Suggested Candidates

solution is found? (smaller is better)

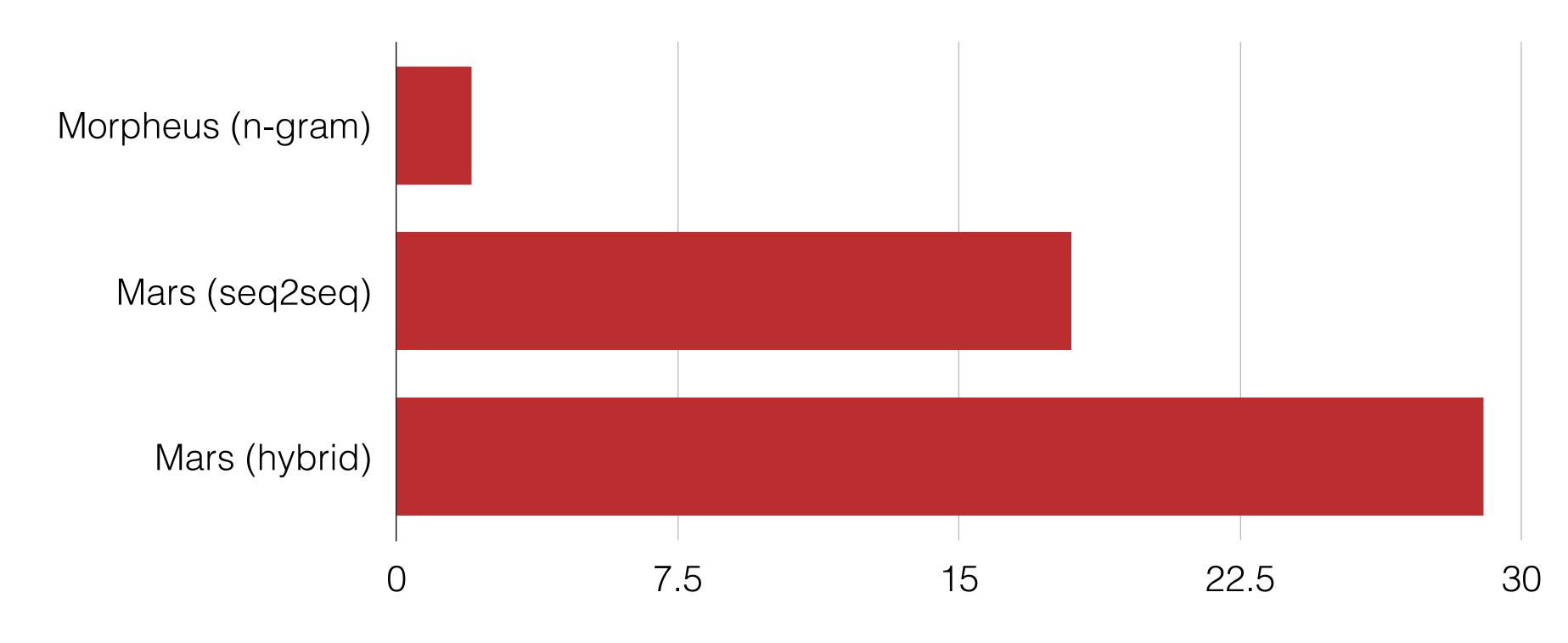


How many symbolic programs need to be enumerated until a

50

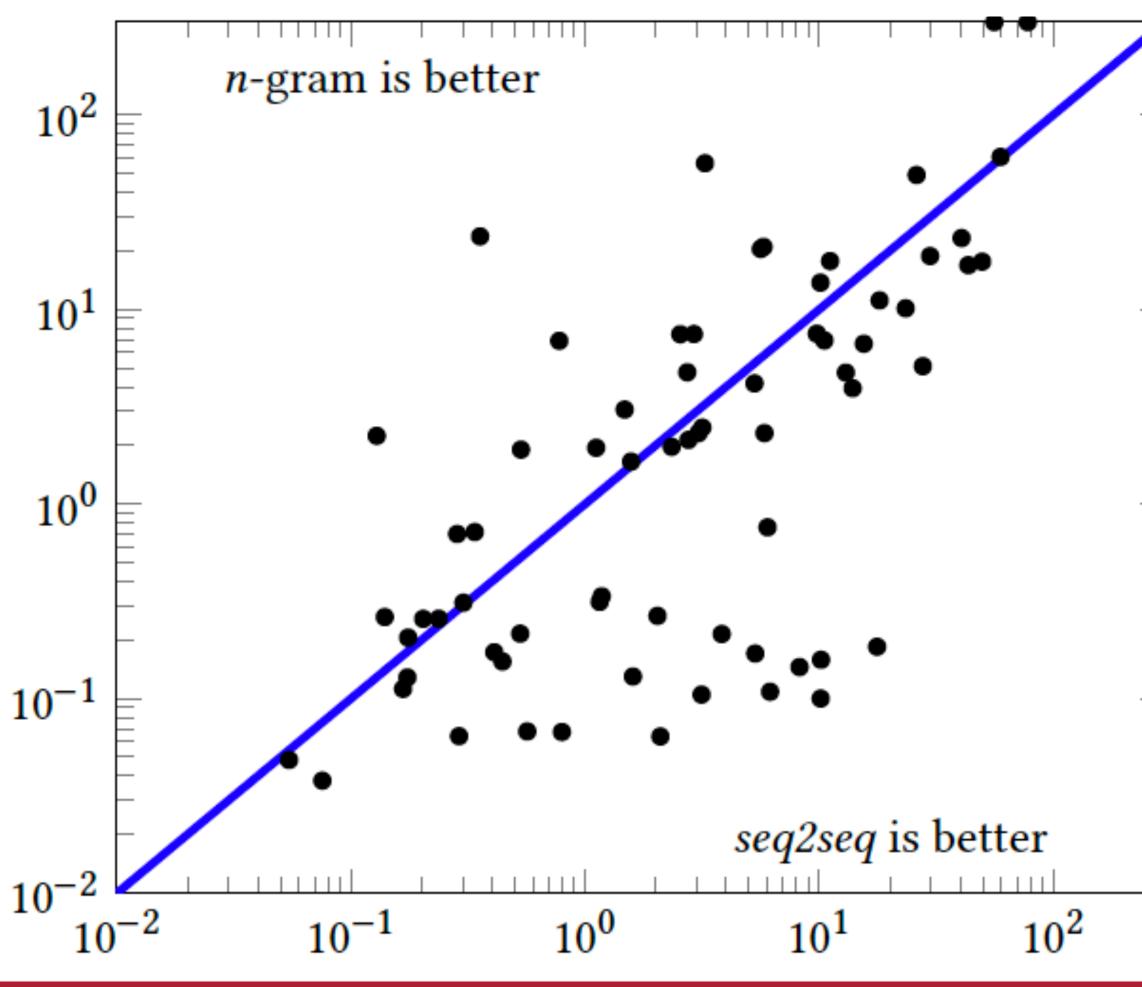
Quality of Suggested Candidates

(larger is better)



How often is the correct symbolic program among the first three?

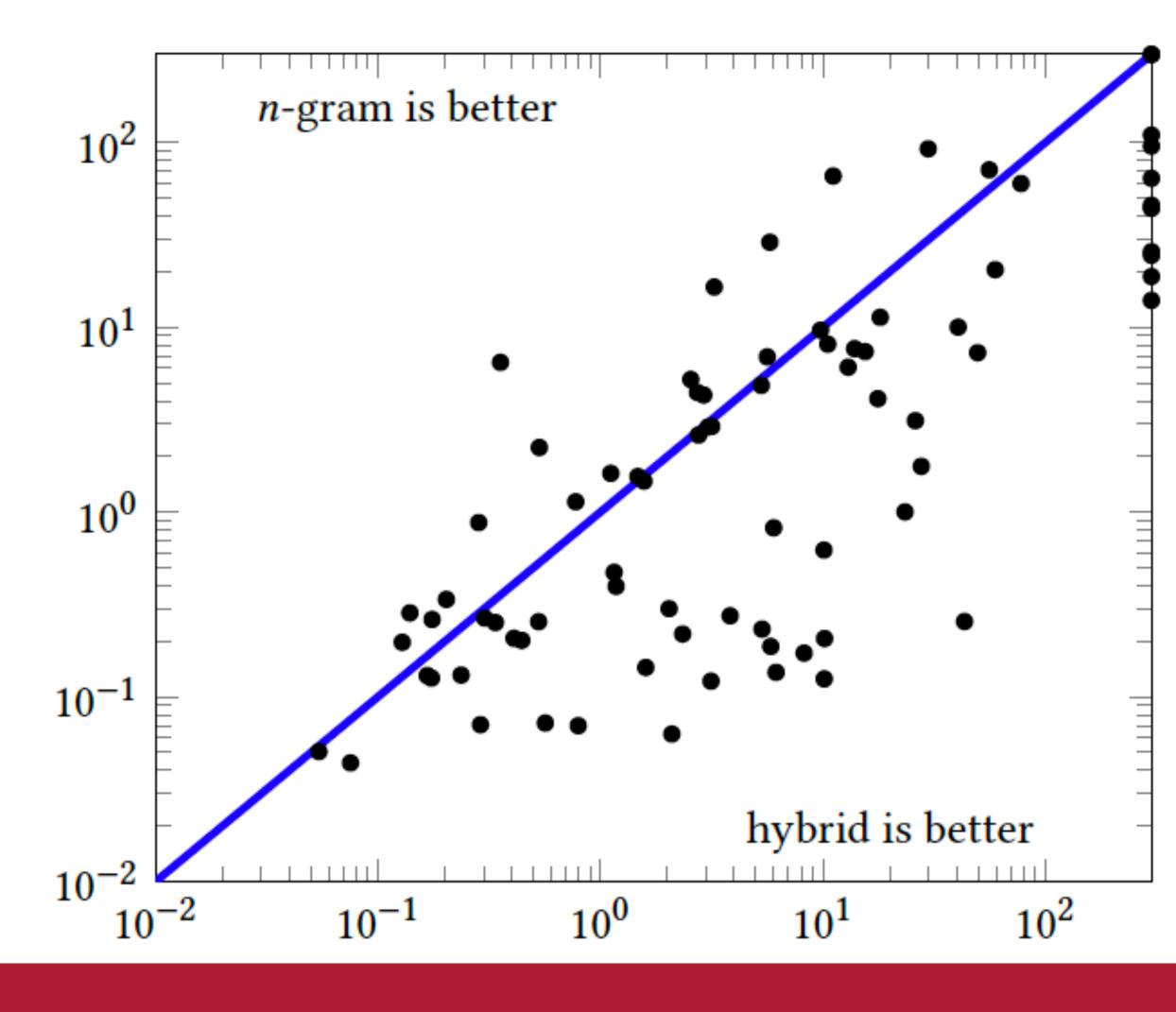
Performance Improvement



Morpheus (n-gram) vs Mars (seq2seq)

	Avg. Speedup	#timeouts
Morpheus (n-gram)	1x	11
Mars (seq2seq)	6x	8

Performance Improvement



Morpheus (n-gram) vs Mars (hybrid)

	Avg. Speedup	#timeouts
Morpheus (n-gram)	1x	11
Mars (hybrid)	15x	2

Takeaway



- seq2seq requires a lot of data:
 - Association rules can find hidden connections between keywords and functions
 - Hybrid neural network model achieves better accuracy
- We can encode multiple specifications as a Maximum Satisfiability Modulo Theory problem (Max-SMT)

Multiple specifications accurately capture the user intent