ars20gramopteng

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Algorithmisch Rekursive Sequenzanalyse 2.0

[]: Example for optimizing grammar

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[]: ## Problem statement and solution approach

Problem

Given is an empirical sequence of terminal characters obtained from a specificular anguage model.

come as close as possible to the terminal characters occurring in the empirical $_{\!\!\!\!\perp}$ -sequence.

Procedure

generate different possible sequences with different probabilities.

grammar, and the relative frequency of each terminal character is calculated.

- correlation coefficient serves as a measure of agreement. Additionally, a p-value is calculated to check the significance of the correlation.
- 4. **Optimization by adjusting probabilities**: If the correlation is low, the probabilities within the grammar are iteratively adjusted to better

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match the generated sequences to the empirical sequence. This ensures that the probabilities are normalized to 1 after each adjustment.

5. **Stopping criterion**: The optimization process ends when a significantly high correlation is achieved or the maximum number of iterations is reached.

### Result

After the optimization process, the adjusted grammar is output, along with the best correlation coefficient and its significance level (p-value). Au significantly high correlation means that the grammar has been successfully adapted to the empirical sequence and can serve as a model for its generation.
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[4]: import numpy as np
     from scipy.stats import spearmanr
     from collections import Counter
     # Beispielhafte empirische Sequenz
     empirical_sequence = ['KBG', 'VBG', 'KBBd', 'VBA', 'KBBd', 'VBA', 'KBA', 'VBA',
                           'KBBd', 'VBBd', 'KBBd', 'VBBd', 'KBA', 'VBA', 'KBA', L

    'VBA',
                           'KAA', 'VAA', 'KAV', 'VAV']
     # Grammatik mit Gewichtungen
     grammar = {
         "<Start>": [["<Begrüßung>", "<Bedarf>", "<Abschluss>", "<Verabschiedung>", "
         "<Begrüßung>": [["KBG", "VBG", 1.0]],
         "<Bedarf>": [["<BedarfSegment>", "<Bedarf>", 0.8], ["<BedarfSegment>", 0.
         "<BedarfSegment>": [["KBBd", "VBBd", 0.4], ["KBBd", "VBA", 0.3], ["KBA", 1.1]
      \hookrightarrow"VBA", 0.3]],
         "<Abschluss>": [["KAA", "VAA", 0.6], ["VAA", "KAA", 0.4]],
         "<Verabschiedung>": [["KAV", "VAV", 0.7], ["VAV", "KAV", 0.3]]
     }
     # Funktion zur Generierung einer Terminalsequenz basierend auf der Grammatik
     def generate_terminal_sequence(grammar):
         sequence = []
         production_stack = ["<Start>"]
         while production_stack:
             production = production_stack.pop()
             if production in grammar:
                 rules = grammar[production]
                 options = [rule[:-1] for rule in rules] # Die Produktionsregeln⊔
      ⇔ohne die Wahrscheinlichkeit
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probabilities = [rule[-1] for rule in rules] # Die_
 \hookrightarrow Wahrscheinlichkeiten
            selected_rule = np.random.choice(len(options), p=probabilities) #__
 →Index der gewählten Regel
            production_stack.extend(reversed(options[selected_rule]))
        else:
            sequence.append(production)
    return sequence
# Berechne die Häufigkeit der Terminalsymbole in einer Liste von Sequenzen
def calculate_frequencies(sequences):
    flat sequence = [symbol for sequence in sequences for symbol in sequence]
    total_count = len(flat_sequence)
    frequencies = {symbol: count / total_count for symbol, count in_{LL}
 →Counter(flat_sequence).items()}
    return frequencies
# Häufigkeiten der empirischen Sequenz
empirical_frequencies = calculate_frequencies([empirical_sequence])
# Optimierungsprozess
best_correlation = -1
best grammar = grammar
best_p_value = None
for _ in range(100): # Maximale Anzahl an Iterationen
    # Generiere mehrere Sequenzen und berechne die Häufigkeit der
 → Terminalzeichen
    generated_sequences = [generate_terminal_sequence(grammar) for _ in__
 →range(1000)]
    generated_frequencies = calculate_frequencies(generated_sequences)
    # Konvertiere Frequenzdaten für Korrelationstest
    empirical_values = np.array([empirical_frequencies.get(symbol, 0) for_
 ⇔symbol in empirical_frequencies])
    generated_values = np.array([generated_frequencies.get(symbol, 0) for__
 ⇔symbol in empirical_frequencies])
    # Berechne die Korrelation zur empirischen Sequenz
    spearman corr, spearman p = spearmanr(empirical values, generated values)
    # Aktualisiere das beste Ergebnis, falls signifikant und besser
    if spearman_p < 0.05 and spearman_corr > best_correlation:
        best_correlation = spearman_corr
        best_grammar = grammar
        best_p_value = spearman_p
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# Wenn die Korrelation akzeptabel ist, beende die Schleife
    if spearman corr > 0.8: # Setze einen gewünschten Korrelationswert fest
        break
    # Andernfalls passe die Wahrscheinlichkeiten leicht an
    for key in grammar:
        for i, rule in enumerate(grammar[key]):
            new_prob = rule[-1] + np.random.uniform(-0.05, 0.05)
            grammar[key][i][-1] = max(0, min(new_prob, 1)) # Stelle sicher,
 ⇔dass die Wahrscheinlichkeiten im Bereich [0, 1] bleiben
        # Normiere die Wahrscheinlichkeiten neu, damit ihre Summe 1 ist
        total_prob = sum(rule[-1] for rule in grammar[key])
        for rule in grammar[key]:
            rule[-1] /= total_prob
# Ausgabe der optimierten Grammatik, Korrelation und Signifikanz
print("Optimized Grammar:", best_grammar)
print("Best Spearman Correlation:", best_correlation)
print("Significance (p-value):", best_p_value)
if best p value < 0.05:
    print("The correlation is statistically significant.")
else:
    print("The correlation is not statistically significant.")
Optimized Grammar: {'<Start>': [['<Begrüßung>', '<Bedarf>', '<Abschluss>',
'<Verabschiedung>', 1.0]], '<Begrüßung>': [['KBG', 'VBG', 1.0]], '<Bedarf>':
[['<BedarfSegment>', '<Bedarf>', 0.8], ['<BedarfSegment>', 0.2]],
'<BedarfSegment>': [['KBBd', 'VBBd', 0.4], ['KBBd', 'VBA', 0.3], ['KBA', 'VBA',
0.3]], '<Abschluss>': [['KAA', 'VAA', 0.6], ['VAA', 'KAA', 0.4]],
'<Verabschiedung>': [['KAV', 'VAV', 0.7], ['VAV', 'KAV', 0.3]]}
Best Spearman Correlation: 0.9692307692307693
Significance (p-value): 3.778488151361357e-06
The correlation is statistically significant.
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