

BACHELOR THESIS
OPTIMIZED PATTERN MATCHING IN GENOMIC DATA

Report

MARTIN WESTH PETERSEN - MQT967
KASPER MYRTUE - VKL275

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1 Analysis

1.1 Scan For Matches

In order to understand what is good from Scan for matches we needed to understand some code segments and the overall structure of the program, simply put we came to understand

B1
($m + 1$)

A
(1)

C
($2m - 1$)

B2
($m + 1$)

this flow in the code: