COMP90025 ASSIGNMENT 02A WRITTEN REPORT

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September 12, 2020

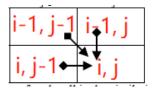


Figure 1: Matrix cell dependency

As we have the cell calculation dependencies (figure 1) for the dynamic programming (dp) approach in sequential algorithm.

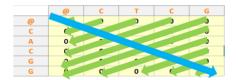


Figure 2: Anti-diagonal matrix traverse

Instead of changing to a new algorithm, first intuitive optimization is to traverse the dp matrix in an anti-diagonal manner (figure 2). Sequentially iterate through each anti-diagonal and parallel dp cell calculation on each anti-diagonal rather than calculating cells in sequential row by row manner. However, this achieves no speedup because there can be a lot of cache missing and false sharing due to the way we traverse the matrix.

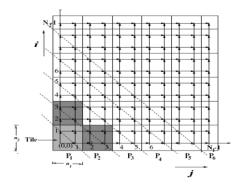


Figure 3: Tiled matrix & Diagonal tile parallelism

Sathe et al. (2011) [1] describes a tiling parallel algorithm. Rather than parallel each cell diagonally, we parallel **tiles** diagonally. Each tile has $\frac{matrix\ width}{p}*\frac{matrix\ height}{p}$ cells where p =#processors (figure 3). Then we apply the anti-diagonal parallelism technique on tiles and calculate cells sequentially in each tile (figure 5) to reduce cache missing. On spartan, I found that 22 threads performs best with this algorithm.

Other speedup techniques:

- 1. add "#pragma omp parallel for" for the two matrix initialization for loops without implicit barriers (figure 4).
- 2. no need to do last branching condition (figure 6).
- 3. Rather than choose p = #processors, choose $p = \frac{4}{3} * \#$ processors to maximize threads usage.

```
#pragma omp parallel
{
    #pragma omp for nowait
    for (i = 0; i <= m; i++) {
        dp[i][0] = i * pgap;
    }
    #pragma omp for
    for (i = 1; i <= n; i++) {
        dp[0][i] = i * pgap;
    }
}</pre>
```

Figure 4: Matrix initialization parallelism

```
int n_parallel = n_threads + 7;
            int tile_width = (int) ceil((1.0*m) / n_parallel), tile_length = (int) ceil((1.0*n) / n_parallel);
            int num_tile_in_width = (int) ceil((1.0*m) / tile_width); int num_tile_in_length = (int) ceil((1.0*n) / tile_length);;
            for (int line = 1; line <= (num_tile_in_width + num_tile_in_length - 1); line++) {
                int start_col = max(0, line - num_tile_in_width);
                int count = min(line, min((num_tile_in_length - start_col), num_tile_in_width));
                #pragma omp parallel for
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                     int tile_i start = (min(num tile_in_width, line)-z-1)*tile_width +1,
tile_j_start = (start_col+z)*tile_length +1;
                     for (int i = tile_i_start; i < min(tile_i_start + tile_width, row); i++) {</pre>
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                          for (int j = tile_j_start; j < min(tile_j_start + tile_length, col); j++) {</pre>
                              if (x[i - 1] == y[j - 1]) {
| dp[i][j] = dp[i - 1][j - 1];
                              } else {
                                  dp[i][j - 1] + pgap);
```

Figure 5: Anti-diagonal tile parallelism & sequential traverse in tile

References

[1] S. R. Sathe and D. D. Shrimankar. Parallelization of DNA sequence alignment using OpenMP. *ACM International Conference Proceeding Series*, pages 200–203, 2011.

```
while (!(i == 0 || j == 0)) {{
    if (x[i - 1] == y[j - 1]) {
        xans[xpos--] = (int) x[i - 1];
        yans[ypos--] = (int) y[j - 1];
        i--;
        j--;
    } else if (dp[i - 1][j - 1] + pxy == dp[i][j]) {
        xans[xpos--] = (int) x[i - 1];
        yans[ypos--] = (int) y[j - 1];
        i--;
        j--;
} else if (dp[i - 1][j] + pgap == dp[i][j]) {
        xans[xpos--] = (int) x[i - 1];
        yans[ypos--] = (int) '_';
        i--;
        // } else if (dp[i][j - 1] + pgap == dp[i][j]) {
        else {
            xans[xpos--] = (int) '_';
            yans[ypos--] = (int) y[j - 1];
            j--;
        }
}
```

Figure 6: Backtracking optimization