## COMP90025 ASSIGNMENT 02A WRITTEN REPORT

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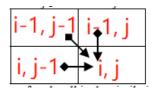


Figure 1: Matrix cell dependency

As we have the cell calculation dependencies shown in figure 1.

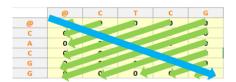


Figure 2: Anti-diagonal matrix traverse

First intuitive optimization is to traverse the matrix in an anti-diagonal manner in figure 2 and parallel each diagonal rather than calculate 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.

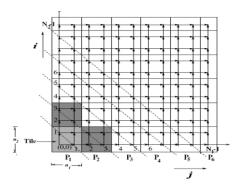


Figure 3: Tiled matrix & Diagonal tile parallelism

Sathe and Shrimankar (2011) 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 as shown in figure 3. Then

we apply the anti-diagonal parallelism technique on tiles and calculate cells sequentially in each tile to reduce cache missing. Implementation can be found in figure 5. 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 in figure 4.
- 2. delete unnecessary "memset (dp[0], 0, size);" in given code.
- 3. delete "delete[] dp[0]; delete[] dp;" to speedup.
- 4. no need to do last branching condition as it needs to execute that step in figure 6.
- 5. 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

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

## References

S. R. Sathe and D. D. Shrimankar. Parallelization of DNA sequence alignment using OpenMP. *ACM International Conference Proceeding Series*, pages 200–203, 2011. doi: 10.1145/1947940.1947983.

```
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