

Here is the exact pseudocode from the PDF for each section, followed by the explanation in the **Input** → **Process** → **Output** format.

4.1 Scan

This algorithm calculates the **prefix sums** (running totals) of an array.

The Pseudocode (Figure 2)

```
scanUp(A, L, f) =  
    if (|A| = 1) then return A[0]  
    else  
        n ← |A|;  
        m ← n/2;  
        l ← scanUp(A[0 : m], L[0 : m - 1], f)  
        ||  
        r ← scanUp(A[m : n], L[m : n - 1],  
f);  
        L[m - 1] ← l;  
        return f(l, r)
```

```

scanDown(R, L, f, s) =
    if (|R| = 1) then R[0] = s; return;
    else
        n ← |A|; // Note: Context implies
this uses the size of the current slice
        m ← |R|/2;
        scanDown(R[0 : m], L[0 : m - 1], s) ||
        scanDown(R[m : n], L[m : n - 1], f(s,
L[m - 1]));
    return

```

```

scan(A, f, I) =
    L ← array[|A| - 1];
    R ← array[|A|];
    total ← scanUp(A, L, f);
    scanDown(R, L, f, I);
    return <R, total>;

```

Explanation

1. Input

- **A** : An array of numbers, e.g., [1, 2, 3, 4].
- **f** : A function to combine them (usually addition +).

- **I** : An identity element (usually **0** for addition).

2. Process

This approach uses a logical tree structure.

- **Step 1 (scanUp)**: The "Summing Phase."
 - The algorithm recursively splits the array in half until it hits single numbers.
 - It adds pairs of results together ($f(l, r)$).
 - It saves partial sums in the array **L** (think of **L** as the internal nodes of a tree).
- **Step 2 (scanDown)**: The "Passing Phase."
 - It takes a value **s** (the sum coming from the left/parent) and pushes it down.
 - **Left Child**: Inherits **s** directly (because it's the start of the sequence).
 - **Right Child**: Inherits **s** + the sum of the Left Child (stored in **L**).

3. Output

- **R** : An array of prefix sums. For input $[1, 2, 3, 4]$, **R** is $[0, 1, 3, 6]$.
- **total** : The sum of the whole array (10).

4.2 Filter

This algorithm removes elements that don't fit a certain rule.

The Pseudocode (Figure 3)

```
FILTER(A, p) =  
  n ← |A|;  
  F ← array[n];  
  parfor i in [0 : n]  
    F[i] ← p(A[i])  
  <X, m> ← plusScan(F);  
  R ← array[m];  
  parfor i in [0 : n]  
    if (F[i]) then R[X[i]] ← A[i]  
  return R
```

Explanation

1. Input

- A : A list of items.
- p : A "predicate" (a true/false question).

Example: "Is the number odd?"

2. Process

- **Line 4-5 (parfor):** Every processor checks one item. If $p(A[i])$ is true, they write 1 into array F . If false, they write 0.
- **Line 6 (plusScan):** We run a scan on F . This calculates the *destination index* for every surviving item.
 - Example $F : [1, 0, 1]$.
 - Scan $X : [0, 1, 1]$. (Item 0 goes to index 0. Item 2 goes to index 1).
- **Line 8-9 (parfor):** Processors look at the flags again. If the flag is 1, they copy the item from A into the new array R at the position calculated by the scan $X[i]$.

3. Output

- R : A condensed array containing only the items that returned true.

4.2 Flatten

This algorithm turns a "list of lists" into one simple list.

The Pseudocode (Figure 4)

```
FLATTEN(A) =  
    sizes ← array(|A|);  
    parfor i in [0 : |A|]  
        sizes[i] ← |A[i]|  
    <X, m> ← plusScan(sizes);  
    R ← array(m);  
    parfor i in [0 : |A|]  
        o ← X[i];  
        parfor j in [0 : |A[i]|]  
            R[o + j] ← A[i][j]  
    return R
```

Explanation

1. Input

- A : A nested array, e.g., `[[10, 20], [30], [40, 50]]`.

2. Process

- **Lines 2-4:** Calculate how big each sub-list is.
Result: `[2, 1, 2]` .
- **Line 5 (`plusScan`):** Calculate starting positions (offsets). Result `X` : `[0, 2, 3]` .
 - Sub-list 1 starts at 0.
 - Sub-list 2 starts at 2.
- **Lines 7-10:** A double loop. The outer loop `i` runs for each sub-list. The inner loop `j` runs for each item in that sub-list.
 - It copies the item `A[i][j]` to `R[o + j]` (Offset + current position).

3. Output

- `R` : One single array `[10, 20, 30, 40, 50]` .
-

4.3 Search

This finds where a value belongs in a sorted list.

The Pseudocode (Figure 5)

```
findBlock(A, v, k) =
  s ← |A|/k;
```

```

    r ← k;
    parfor i in [0 : k]
        if (A[i * s] < v and A[(i + 1) * s] >
v)
            then r ← i
    return (A[r * s, (r + 1) * s], i * s)

search(A, v, k) =
    (B, o) = findBlock(A, v, min(|A|, k));
    if (|A| ≤ k) then return o
    else return o + search(B, v, k);

```

Explanation

1. Input

- **A** : A sorted array.
- **v** : The value we are looking for.
- **k** : The number of splits (how many processors we can use).

2. Process

- **findBlock** : Instead of checking just the middle (like Binary Search), we check **k** positions evenly spaced out.

- The `parfor` checks all `k` spots at the same time.
- It identifies which "block" (slice) of the array holds the value `v`.
- `search` : This is the main loop.
 - It calls `findBlock` to narrow down the array to a smaller slice `B`.
 - It adds the offset `o` to keep track of the total position.
 - It recurses (calls itself) on the smaller slice `B`.

3. Output

- An integer index representing where `v` is (or should be) in `A`.
-

4.4 Merge

This combines two sorted arrays into one.

The Pseudocode (Figure 7)

```

MERGE(A, B, R) =
  case (|A|, |B|) of
    (0, _)  $\Rightarrow$  copy B to R; return;
    (_, 0)  $\Rightarrow$  copy A to R; return;
    otherwise  $\Rightarrow$ 
      m  $\leftarrow$  |R|/2;
      (ma, mb) = KTH(A, B, m);
      MERGE(A[0 : ma], B[0 : mb], R[0 : m])
  ||
      MERGE(A[ma : |A|], B[mb : |B|], R[m :
|R|]);
  return

```

Explanation

1. Input

- **A** and **B** : Two arrays that are already sorted.
- **R** : An empty result array.

2. Process

- **Line 6 (m)**: We pick the midpoint of the final result array.
- **Line 7 (KTH)**: We calculate how many elements from **A** (**ma**) and how many from **B** (**mb**) are

needed to fill the first half of R . (This KTH helper function is complex, but essentially it finds split points).

- **Lines 8-9 (`||`):**
 - **Left Processor:** Merges the small halves of A and B into the left half of R .
 - **Right Processor:** Merges the big halves of A and B into the right half of R .
 - These run at the exact same time.

3. Output

- R : The final array containing all elements from A and B , perfectly sorted.

4.5 K-th Smallest

This finds the element that would be at index k if the array were sorted.

The Pseudocode (Figure 9)

```
kthSmallest(A, k) =  
    p ← selectPivot(A);
```

```
L ← filter(A, \x.(x < p));
G ← filter(A, \x.(x > p));
if (k < |L|) then
    return kthSmallest(L, k)
else if (k > |A| - |G|) then
    return kthSmallest(G, k - (|A| - |G|))
else return p
```

Explanation

1. Input

- **A** : A scrambled (unsorted) array.
- **k** : The rank we want (e.g., 0 for min, $|A|/2$ for median).

2. Process

- **Line 2:** Pick a "Pivot" number **p** (usually random).
- **Lines 3-4:** Use the **filter** function (from 4.2) to create two new arrays:
 - **L** : All numbers smaller than **p**.
 - **G** : All numbers bigger than **p**.
- **Lines 5-9:** Logic check:

- If k is smaller than the size of L , the answer must be inside L . We recurse on L .
- If k is very large (larger than everything except G), the answer is in G . We recurse on G (subtracting the count of the numbers we skipped).
- Otherwise, the answer is the pivot p itself.

3. Output

- The single value that represents the k -th smallest number.