

CAD Pseudo Code

```
*****
CAD:
    A function that finds the anomalies of a time series of random deviates from
    various normal distributions. (I need to clarify this further)

Calls:
    anomaly_finder

Called by:
    None

Input Parameters:
    time_series - the time series being searched for the anomalies
    delta - the value to be added to the value to the CUSUM parameter k; its
             default value is 3
    lambda - the minimum length of the anomalous subsequence the code should
             detect; the default is 5
    type - the type of the anomaly "upper" or "lower"
    number_of_windows - the number of windows used
    step_size - the distance between two consecutive windows
    training_set_window_length - the starting length of the training set searched
                               for within each window
    training_set_step_size - the size of the shift by which the
                             training_set_windows are moved down the sequence.
    starting_k_value - the starting value of the CUSUM parameter k
    starting_H_value - the starting value of the CUSUM parameter H

Returns:
    anomaly_indices - The indices of the anomalies
*****

FUNCTION    CAD(time_series, delta, lambda, type,
                number_of_windows, step_size, training_set_window_length,
                training_set_step_size)

    n <- LEN(time_series)
    window_length <- n - (number_of_windows*step_size) + 1
    INIT anomaly_indices <- NULL

    FOR i FROM 1 TO (number_of_windows*step_size) BY      step_size

        temp_var <- time_series[i TO (i+window_length -1)]
        temp_indices <- anomaly_finder(temp_var, delta, lambda,
                                     type, i, step_size,
                                     training_set_window_length,
                                     training_set_step_size)
        anomaly_indices <- CONCATENATE(anomaly_indices,
                                     temp_indices)

    ENDFOR

    RETURN(anomaly_indices)

ENDFUNCTION
```

```

*****
anomaly_finder:
    A function that finds the anomalies for a specific window
    of a time series

Calls:
    training_subsequence_finder
    evaluate_CUSUM_results

Called by:
    CAD

Input Parameters:
    subsqnce - a window length subsequence of the original
               time series.
    delta - the value to be added to the value to the CUSUM parameter k; its
            default value is 3
    lambda - the minimum length of the anomalous subsequence the code should
             detect; the default is 5
    type - the type of the anomaly "upper" or "lower"
    indx - the index of the first element of subsqnce within time_series.
    step_size - the distance between two consecutive windows
    training_set_window_length - the starting length of the training set searched
                                for within each window
    training_set_step_size - the size of the shift by which the
                             training_set_windows are moved down the sequence.
    starting_k_value - the starting value of the CUSUM parameter k
    starting_H_value - the starting value of the CUSUM parameter H

Returns:
    global_indices - The indices of the anomalies within the original time_series.
*****

FUNCTION anomaly_finder(subsqnce, delta, lambda, type, indx, step_size,
    training_window_length, training_set_step_size,
    starting_k_value, starting_H_value)

    training_set_obj <- training_subsequence_finder(subsqnce,
    training_window_length, training_set_step_size, starting_k_value,
    starting_H_value)

    IF training_set_obj[1] = NULL THEN
        WARNING("No training set was found.")
        RETURN(NULL)
    ELSE
        x_bar <- MEAN(training_set_obj[1])
        sigma <- STANDARD_DEVIATION(training_set_obj[1])
        k <- training_set_obj[2] + delta
        H <- training_set_obj[3]
        local_indices <- evaluate_CUSUM_results(subsqnce, x_bar, sigma, H, k,
            type, lambda)
        IF LEN(local_indices) != 0 THEN
            global_indices <- local_indices + indx -1
        ELSE
            global_indices <- NULL
        ENDIF
        RETURN(global_indices)
    ENDIF
ENDFUNCTION

```

training_subsequence_finder:

A function that searches for a subsequence of a time series that is both (close to) normal and in statistical control. This subsequence must have a minimal length of 30 and maximal length of **training_set_window_length**.

Calls:

normality_finder
parameter_finder

Called by:

anomaly_finder

Input Parameters:

x - a time series; it must have length greater than 30.

training_set_window_length - the starting length of the training set searched for within each window

training_set_step_size - the size of the shift by which the training_set_windows are moved down the sequence.

starting_k_value - the starting value of the CUSUM parameter k

starting_H_value - the starting value of the CUSUM parameter H

Returns:

training_set_obj - a list/object of length 3, where

training_set_obj[1] - the subsequence of **x** that is to be used as the training data set

training_set_obj[2] - the CUSUM k value of **training_set_obj[1]**

training_set_obj[3] - the CUSUM H value of **training_set_obj[1]**

FUNCTION training_subsequence_finder(x, training_set_window_length, training_set_step_size, starting_k_value, starting_H_value)

INIT segment_found <- FALSE

INIT cant_be_done <- FALSE

INIT first_time_through <- TRUE

INIT parameters_plus <- NULL

n <- LEN(x)

WHILE (NOT segment_found) AND (NOT cant_be_done)

FOR i FROM 1 TO (n - training_set_window_length +1) BY training_set_step_size

temp <- x[i TO (i+training_set_window_length-1)]

temp_obj <- normality_finder(temp)

IF temp_obj[1]> 0.05 THEN

p_value <- temp_obj[1]

skew <- temp_obj[2]

kurt <- temp_obj[3]

parameters_obj <- parameter_finder(temp, starting_k_value, starting_H_value)

IF (parameters_obj[1] != NULL) THEN

the_J <- ABS(skew) + ABS(3 - kurt)+ ABS(1-p_value)+

(100*parameters_obj[1])+(100*parameters_obj[2])

dummy_var <- [temp, p_value, skew, parameters_obj[1], parameters_obj[2],the_J]

IF first_time_through THEN

parameters_plus <- dummy_var

first_time_through <- FALSE

```

ELSE
    IF parameters_plus[6] > the_J
        parameters_plus <- dummy_var
    ENDIF
ENDIF
ENDIF
ENDIF
ENDFOR

IF parameters_plus != NULL THEN
    segment_found <- TRUE
ELSE
    IF training_set_window_length > 30 THEN
        training_set_window_length <-
            training_set_window_length -1
    ELSE
        cant_be_done <- TRUE
    ENDIF
ENDIF
ENDWHILE

IF cant_be_done THEN
    return_obj <- [NULL, NULL, NULL]
ELSE
    return_obj <- [parameters_plus[1], parameters_plus[4],
parameters_plus[5]]
ENDIF

RETURN(return_obj)

ENDFUNCTION

```

```

*****
normality_finder:
    A function that finds Skewness, Kurtosis and Shapiro-Wilk normality test p-
    value for a time series (the input.)

Calls:
    SKEWNESS
    KURTOSIS
    SHAPIRO_WILK_P_VALUE

Called by:
    training_subsequence_finder

Input Parameters:
    x - a time series; it must have length of at least 30.

Returns:
    normality_obj - a list/obj of length 3, where
        normality_obj[1] - skewness of x
        normality_obj[2] - kurtosis of x
        normality_obj[3] - Shapiro-Wilk p-value of x
*****

```

```

FUNCTION normality_finder(x)

    temp1 <- SKEWNESS(x)
    temp2 <- KURTOSIS(x)
    temp3 <- SHAPIRO_WILK_P_VALUE(x)

    return([temp1, temp2, temp3])

ENDFUNCTION

```

parameter_finder:

A function takes a time series *x* of length at least 30 that has a (nearly) normal distribution and selects the smallest CUSUM parameters *k* and *H* for which the time series is in statistical control

Calls:

CUSUM

Called by:

training_subsequence_finder

Input Parameters:

x - a time series of length of at least 30 with (nearly) normal distribution
starting_k_value - the starting value of the CUSUM parameter *k*, default is 3
starting_H_value - the starting value of the CUSUM parameter *H*, default is 5

Returns:

parameter_obj - a list/object of length 2, where
parameter_obj[1] - the smallest CUSUM *k* value for which *x* is in statistical control
parameter_obj[2] - the smallest CUSUM *H* value for which *x* is in statistical control



NOTE: In the current version of the code, *H* stays fixed at 5 and it is not modified at all. It might need to be messed with in future versions of CAD. But, *H*, as it stands now, can be left out of the code.

```

FUNCTION parameter_finder(x, starting_k_value = 3, starting_H_value = 5)
  INIT parameters_found <- FALSE
  x_bar <- MEAN(x)
  sigma <- STANDARD_DEVIATION(x)

  WHILE (NOT parameters_found)
    temp_obj <- CUSUM(x, x_bar, sigma, H, k)
    IF (temp_obj[3] = NULL AND temp_obj[4] = NULL ) THEN
      parameters_found <- TRUE
    ELSE
      IF k > 20 THEN
        WARNING("Something is off in CUSUM land")
        k <- NULL
        parameters_found <- TRUE
      ELSE
        k <- k +1
      ENDIF
    ENDIF
  ENDWHILE

  RETURN ([k, H])
ENDFUNCTION

```

evaluate_CUSUM_results:

A function that finds the indices of anomalies, if there are any, of the time series **x**, given its mean, standard deviation, and the minimum CUSUM parameters H and k for which **x** should be in statistical control.

Calls:

turning_points_finder
interval_finder
CUSUM

Called by:

anomaly_finder

Input Parameters:

x - a time series length at least 30
x_bar - the mean of **x**
sigma - the standard deviation of **x**
k - CUSUM parameter k
H - CUSUM parameter H
type - the type of the anomaly "upper" or "lower"
lambda - the minimum length of the anomalous subsequence the code should detect; the default is 5

Returns:

indices - The indices of the anomalies within **x**.

```
FUNCTION evaluate_CUSUM_results(x, x_bar, sigma, H=5, k, type, lambda)
```

```
INIT hi_sum_indices <- NULL
INIT low_sum_indices <- NULL
INIT indices <- NULL
```

```
CUSUM_obj <- CUSUM(x, x_bar, sigma, H, k)
low_sums<-CUSUM_obj[1]
hi_sums<-CUSUM_obj[2]
upper_viol_index<-CUSUM_obj[3]
lower_viol_index<-CUSUM_obj[4]
lower_viol<-low_sums[lower_viol_index]
upper_viol<-hi_sums[upper_viol_index]
```

```
IF (LEN(upper_viol_index) > 0) THEN
  high_sum_turning_pts <- turning_points_finder(hi_sums)
  IF (high_sum_turning_pts[1] != 1) THEN
    high_sum_turning_pts <- CONCATENATE(1, high_sum_turning_pts)
  ENDIF
  hi_sum_df <- interval_finder(hi_sums, high_sum_turning_pts, type)
ENDIF
```

```
IF ( LEN(lower_viol_index) > 0) THEN
  low_sum_turning_pts <- turning_points_finder(low_sums)
  IF (low_sum_turning_pts[1] != 1) THEN
    low_sum_turning_pts <- CONCAT(1, low_sum_turning_pts)
  ENDIF
  low_sum_df <- interval_finder(low_sums, low_sum_turning_pts, type)
ENDIF
```

```
IF (type = "lower") THEN
##### Finding "lower" anomalies #####
```

```

#finding the indices of the decreasing terms for the upper violations seq.
IF (LEN(upper_viol_index) > 0) THEN
    dummy_df <-
    SELECT
        "left_index", "right_index"
    FROM
        hi_sum_df
    WHERE
        (sign = "decreasing") AND ((left_index-right_index)>lambda)

    IF dummy_df != NULL THEN
        n <- NUMBER_OF_ROWS(dummy_df)
        FOR i FROM 1 TO n
            IF (left_index[i] != NULL AND right_index != NULL) THEN
                interval <- [FROM left_index[i] TO right_index[i]]
                interval <- interval INTERSECT upper_viol_index
                IF (LEN(interval) > lambda) THEN
                    hi_sum_indices <- CONCATENATE(hi_sum_indices,
                                                    interval)
                ENDIF
            ENDIF
        ENDFOR
    ENDIF
ENDIF

#finding the indices of the decreasing terms for the lower violations seq.
IF (LEN(lower_viol_index) > 0) THEN
    dummy_df <-

    SELECT
        "left_index", "right_index"
    FROM
        low_sum_df
    WHERE
        (sign = "decreasing") AND ((right_index - left_index) > lambda)

    IF dummy_df != NULL THEN
        n <- NUMBER_OF_ROWS(dummy_df)
        FOR i FROM 1 TO n
            IF (left_index[i] != NULL AND right_index != NULL) THEN
                interval <- [FROM left_index[i] TO right_index[i]]
                interval <- interval INTERSECT lower_viol_index
                IF (LEN(interval) > lambda) THEN
                    low_sum_indices <- CONCATENATE(low_sum_indices,
                                                    interval)
                ENDIF
            ENDIF
        ENDFOR
    ENDIF
ENDIF

indices <- CONCATENATE(hi_sum_indices, low_sum_indices)

ELSE
##### Finding "upper" anomalies #####

```



```

#finding the indices of the increasing terms for the upper violations seq.
IF (LEN(upper_viol_index) > 0) THEN
    dummy_df <-

    SELECT
        "left_index","right_index"
    FROM
        hi_sum_df
    WHERE
        (sign = "increasing") AND ((right_index - left_index) > lambda)

    IF dummy_df != NULL THEN
        n <- NUMBER_OF_ROWS(dummy_df)
        FOR i FROM 1 TO n
            IF (left_index[i] != NULL AND right_index != NULL) THEN
                interval <- [FROM left_index[i] TO right_index[i]]
                interval <- interval INTERSECT upper_viol_index
                IF (LEN(interval) > lambda) THEN
                    hi_sum_indices <- CONCATENATE(hi_sum_indices,
                                                    interval)
                ENDIF
            ENDIF
        ENDFOR
    ENDIF
ENDIF

#finding the indices of the increasing terms for the lower violations seq.
IF (LEN(lower_viol_index) > 0) THEN
    dummy_df <-

    SELECT
        "left_index","right_index"
    FROM
        low_sum_df
    WHERE
        (sign = "increasing") AND ((right_index - left_index) > lambda)

    IF dummy_df != NULL THEN
        n <- NUMBER_OF_ROWS(dummy_df)
        FOR i FROM 1 TO n
            IF (left_index[i] != NULL AND right_index != NULL) THEN
                interval <- [FROM left_index[i] TO right_index[i]]
                interval <- interval INTERSECT lower_viol_index
                IF (LEN(interval) > lambda) THEN
                    low_sum_indices <- CONCATENATE(low_sum_indices,
                                                    interval)
                ENDIF
            ENDIF
        ENDFOR
    ENDIF
ENDIF

indices <- CONCATENATE(hi_sum_indices, low_sum_indices)
ENDIF

RETURN(indices)

ENDFUNCTION

```

turning_points_finder:

A recursively defined function that takes a sequence of numbers, **x**, and finds the indices of the values at which the sequence turns from increasing to decreasing or vice-versa

Calls:

SIGN

Called by:

evaluate_CUSUM_results

Input Parameters:

x - a sequence of numbers longer than 1
turning_points - recursive variable, set to **NULL** when calling the function
sgn - recursive variable, set to **0** when calling the function
index - recursive variable, set to **1** when calling the function

Returns:

turning_points - The indices of the turning points of **x**

```

FUNCTION turning_points_finder(x, turning_points = NULL, sgn = 0, index = 1)
  n <- LEN(turning_points)

  IF ((LEN(x) = 1) OR (LEN(x) = 0)) THEN
    IF (LEN(turning_points) > 0) THEN
      turning_points <- CONCATENATE(turning_points, index)
    ENDIF
    RETURN(turning_points)
  ENDIF

  diff <- x[2] - x[1]

  IF (diff = 0) THEN
    IF (sgn != 0) THEN
      turning_points <- c(turning_points, index)
    ENDIF
    sgn <- 0
    index <- index + 1
    RETURN(turning_points_finder( x[FROM 2 TO LEN(x)], turning_points, sgn,
      index))
  ELSE
    IF (SIGN(diff) = sgn) THEN
      index <- index + 1
      RETURN(turning_points_finder( x[FROM 2 TO LEN(x)], turning_points, sgn,
        index))
    ELSE
      sgn <- SIGN(diff)
      turning_points <- CONCATENATE( turning_points, index)
      index <- index + 1
      RETURN(turning_points_finder( x[FROM 2 TO LEN(x)], turning_points, sgn,
        index))
    ENDIF
  ENDIF
ENDIF
ENDFUNCTION

```

```

*****
interval_finder:
    Given a sequence, x, this function finds the endpoints of the monotone
    increasing, monotone decreasing and constant subsequences it is composed of.

Calls:
    sign_finder

Called by:
    evaluate_CUSUM_results

Input Parameters:
    x - a sequence of numbers longer than 1
    tp_x - the indices of the turning points of x

Returns:
    df - A table with the following schema:
        right_index: right indices of the intervals
        left_index: left indices of the intervals
        right_value: right endpoint values
        left_value: left endpoint values
        sign: "increasing", "decreasing", "constant"
*****

```

```

FUNCTION interval_finder(x, tp_x)

    IF ((LEN(x) <= 1) OR LEN(tp_x) <= 1) THEN
        RETURN(NULL)
    ENDIF

    INIT sign <- NULL

    n <- LEN(tp_x)
    left_endpt_indices <- tp_x[ FROM 1 TO (n - 1)]
    right_endpt_indices <- tp_x[ FROM 2 TO n]

    tp_values1 <- x[left_endpt_indices]
    shifted <- left_endpt_indices + 1    #1 is added to each element of left_endpt_indices
    pt_value_after_tp <- x[shifted]

    FOR i FROM 1 TO (n-1)
        sign[i] <- sign_finder(tp_values1[i], pt_value_after_tp[i])
    ENDFOR

    CREATE TABLE df
        left_endpt_indices
        right_endpt_indices
        sign
    END CREATE TABLE

    RETURN(df)

ENDFUNCTION

```



I had some safety checks ensuring that the values of the input sequence between the turning points were indeed monotone increasing, monotone decreasing or constant. I excluded that from the code here.

sign_finder:

Given two values, **a1** and **a2**, it returns "increasing" if $(a2 - a1) > 0$
returns "decreasing" if $(a2 - a1) < 0$, return "constant" if $a2 = a1$.

Calls:

none

Called by:

interval_finder

Input Parameters:

a1 - a numeric value

a2 - a numeric value

Returns:

sign - a factor variable with "increasing", "decreasing" or "constant" as
possible values

```
FUNCTION sign_finder(x1, x2)
  IF (x2 - x1) > 0 THEN
    RETURN("increasing")
  ELSE
    IF (x2 - x1) < 0 THEN
      RETURN("decreasing")
    ELSE
      RETURN("constant")
   ENDIF
  ENDIF
ENDFUNCTION
```

SIGN:

Given a value **x**, it returns 1 if **x** is positive, -1 if **x** is negative and 0 if **x** =0.

Calls:

none

Called by:

turning_points_finder

Input Parameters:

x - a numeric value

Returns:

s - a numeric value of -1, 0 or 1

```
FUNCTION SIGN(x)
  IF (x < 0) THEN
    RETURN(-1)
  ELSE
    IF (x > 0) THEN
      RETURN(1)
    ELSE
      RETURN(0)
    ENDIF
  ENDIF
ENDFUNCTION
```