Where:

* *Ci*,*j*​ is the value in the *i*th feature column and *j*th row.
* *n* is the total number of feature columns.
* *n*0​ is the number of rows in Normal Group 0 (where the predicted anomaly score value is below or equal to the threshold).
* *n*1​ is the number of rows in Outlier Group 1 (where the predicted anomaly score value is above the threshold).
* Min\_Value*i*​ and Max Value Max\_Value*i*​ are the minimum and maximum values in the *i*th column, respectively.

This formula calculates the separation distance between normalized values between the two groups (defined by the predicted anomaly score column: outlier and normal) across all n columns.