

# demo document

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## 1 Equations

This is equation section.  $Studentcount = \mu ni + \sum \frac{A_k}{B_k}$  is an example for inline equation.  $Student_{count} = \mu ni + \sum \frac{A_k}{B_k}$  is another example. Equation-2 is an example for numbered equation.

$$a = 10b = 90 \tag{1}$$

$$a = 09 + 89 - 67$$

$$\frac{\sum \frac{a}{\varpi b}}{b - 90} \tag{2}$$

In the content based analysis, the amount of  $T$  contained by  $R$  is quantified. The respective contents of  $T$  present in each of the partitioned zones are symbolized by  $l^s$ ,  $l^a$ ,  $l^p$ ,  $l^{ua}$  and  $l^{us}$  as listed above. The relative contents  $\frac{l^s}{l^t}$ ,  $\frac{l^a}{l^t}$ ,  $\frac{l^p}{l^t}$ ,  $\frac{l^{ua}}{l^t}$  and  $\frac{l^{us}}{l^t}$  are defined. They are multiplied by suitable weightage factors, which are so chosen to properly project the relative contents. The set of weightage factors ( $w_a, w_b, w_c, w_d, w_e$ ) used for content based computations are given in table-??. The expression for similarity due to the content component in the  $k^{th}$  feature is given by

$$S_c(R_k, T_k) = \frac{l^s}{l^t} w_a + \frac{l^a}{l^t} w_b + \frac{l^p}{l^t} w_c + \frac{l^{ua}}{l^t} w_d + \frac{l^{us}}{l^t} w_e \tag{3}$$

The expression for similarity due to the content component in the  $k^{th}$  feature is given by:

$$D_c(R_k, T_k) = 1 - S_c(R_k, T_k) \tag{4}$$

## 2 Tables

This section shows the syntax for inserting tables.

The different examples are as follows.

val1	val2	val3	val4
10	10	30	40

Table 1: This is a demo table1

Table 2: This is a demo table2

val1	val2	val3	val4
10	10	30	40

Table 3: Interpretation of the values in feature-interaction table(FIT). FIT data  $Dj_i$  indicates data from row-j and column-i

FIT data	Feature inter-action value	Common Heading		
		Inferred from previously observed features' values	Inferred from present feature value	Net/overall affiliation
D1i	0	healthy	healthy	healthy
D4i	1	PD	PD	PD

Table 4: Proposed Range of distribution zones for different distribution patterns.  $\mu_{ni}$  represents mean value of  $i^{th}$  feature in distribution of healthy samples,  $\sigma_{ni}$  represents the standard deviation.  $\mu_{ai}$  and  $\sigma_{ai}$  represent the same in PD samples.

Distribution Pattern	Safe (Wi=00.1)	Acceptable (Wi=0.1)	Permissible (Wi=1)	Unacceptable (Wi=10)	Unsafe (Wi=100)
1	$\leq (\mu_{ni} - 2\sigma_{ni})$	$(\mu_{ni} - 2\sigma_{ni})$ to $(\mu_{ni} - \sigma_{ni})$	$(\mu_{ni} - \sigma_{ni})$ to $(\mu_{ni} + \sigma_{ni})$	$(\mu_{ni} + \sigma_{ni})$ to $(\mu_{ni} + 2\sigma_{ni})$	$\geq (\mu_{ni} + 2\sigma_{ni})$
2	-	$\leq min_{ai}$	$min_{ai}$ to $max_{ai}$	$\geq max_{ai}$	-
		$\geq max_{ai}$	$max_{ai}$	$\leq min_{ni}$	

Table 5: Regions of partitions and corresponding weightage factors.

	Region of $T^k = (\underline{T}_k, \bar{T}_k)$	Weightage factors for analysis based on	
		Position	Content
Case 1	$< Z_s$ (pattern 1) or $> Z_s$ (pattern 4)	$W_1 = 0.1$	$W_a = 1.0$
Case 2	$[Z_s, Z_a]$ (patterns 1 and 2) or $[Z_a, Z_s]$ (patterns 3 and 4)	$W_2 = 0.3$	$W_b = 0.8$