



A novel distance between single valued neutrosophic sets and its application in pattern recognition

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Abstract

Among the most classic measures in single valued neutrosophic sets (*SVNS*) theory, the distance is an important tool to compare and calculate degree of difference between *SVNS*. Although there exist some types of distance for single valued neutrosophic sets, most of them lack of strictly axiomatic definition and exist counter-intuitive cases. In this paper, a novel distance between single valued neutrosophic sets based on matrix norm is given. Then we proved that the new distance satisfies the axiomatic definition of the metric. Finally, the distance is applied to pattern recognition and medical diagnoses.

Keywords Single valued neutrosophic sets · Distance measure · Pattern recognition · Medical diagnoses

1 Introduction

In the real world, uncertain, vague, inaccurate information can be found in many domains. The theory of fuzzy set had been proposed by Zadeh (1965), which had been a great success in dealing with uncertainty. Later, some extension of the fuzzy set has been studied by many scholars. For example, Atanassov (1986) presented intuitionistic fuzzy sets, which are characterized by a membership function and a non-membership function. Intuitionistic fuzzy sets are considered to be more effective method to handle uncertain, vague information. However, the intuitionistic fuzzy sets cannot better depict vague, indeterminate and inconsistent information. For example, three groups of experts evaluate the benefits of the fund, the first groups of experts believes that the probability of the fund will be profitable is 0.6, the second groups of experts believes that the probability of the fund will be loss is 0.3, the third groups of experts not sure whether the fund will be profitable is 0.3. In order to more comprehensive handle indeterminate and inconsistent information, Smarandache (1999, 2010) introduced the concept of neutrosophic set, which characterized by a true membership function $t_A(x)$, a false membership function $f_A(x)$ and an indetermi-

nate function $i_A(x)$, where $t_A(x), i_A(x), f_A(x) \in]0^-, 1^+[$. Ma et al. (2019) studied generalized neutrosophic extended triplet group. Since the nonstandard interval $]0^-, 1^+[$ may result in neutrosophic set is not easy to apply in practical problems. Smarandache (1998) and Wang et al. (2010) proposed single valued neutrosophic set (*SVNS*), which characterized by a true membership function $t_A(x)$, a false membership function $f_A(x)$ and an indeterminate function $i_A(x)$, where $t_A(x), i_A(x), f_A(x) \in [0, 1]$. In recent years, single-valued neutrosophic set has been applied in many fields including deductive filters (Borzooei et al. 2017), clustering analysis (Karaaslan 2017), decision-making problems and control theory (Liu and Li 2017; Liu 2016; Huang 2016; Luo et al. 2019; Zhang et al. 2018).

Distance measure is effective mathematical tool to measure the degree of difference between two objects. In recent years, the study of distance has attracted considerable attention of many researchers. As a pioneering research, (Broumi and Smarandache 2013b) presented Hausdorff distance between neutrosophic sets (NS). Later, the same author (Broumi and Smarandache 2015) extended Hausdorff distance for single valued neutrosophic sets and applied in medical diagnosis. The Hamming distance, Euclidean distance, normalized Hamming distance and normalized Euclidean distance between neutrosophic sets were studied by Majumdar (2015). The generalized weighted distance between neutrosophic sets was given by Ye (2013). Broumi et al. (2014) gave generalized weighted distance between interval neutrosophic sets. Huang (2016) defined a new

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distance measure, which consider the relationship of truth membership function, indeterminacy-membership function and falsity-membership function. In order to more precision, Garg Nancy (2017) proposed some new distance measures along with two parameters and applied them to pattern recognition and medical diagnoses. Besides, Samuel et al. introduced cosine logarithmic distance (Samuel and Narmadhagnanam 2018) between single valued neutrosophic sets. Ren et al. (2019) studied a Chi-square distance-based similarity measure between single valued neutrosophic sets and applied it to pattern recognition. Similarity measure can be used express the degree of similarity between two objects. Majumdar and Samanta (2014) studied some similarity measures between single valued neutrosophic sets. Broumi and Smarandache (2013b) given several similarity measures. Hamming distance, Euclidean distance and similarity measures between interval neutrosophic sets were studied by Ye (2014a). Ye (2014b) studied vector similarity measure for single valued neutrosophic sets and applied it to multi-criteria decision-making problems. Recently, some similarity measures between single valued neutrosophic sets have been put forward, such as similarity measures based on tangent function, cotangent function and generalized Dice measures (Ye and Fu 2016; Ye 2017b, 2015).

Although the existing distance takes into account the inter-relationship between elements, there are often unreasonable results in some cases (see Example 1). In order to deal with these cases, motivated by Luo and Zhao (2018) proposed distance measure between intuitionistic fuzzy sets, we give a novel distance between single valued neutrosophic sets based on matrix norm and a strictly monotonic real function.

The rest of this paper is organized as follows. In Sect. 2, we review some basic concepts for single valued neutrosophic sets and main results related to distance measure. In Sect. 3, a new distance between single valued neutrosophic sets is proposed. In Sect. 4, the effectiveness and feasibility of the novel distance measures are illustrated with some numerical examples of pattern recognition as well as medical diagnoses. The final section includes conclusion and further research.

2 Preliminaries

In this section, we review some basic concepts for single valued neutrosophic sets and main results related to distance measure, which will be applied to other sections of the paper. In this paper, $SVNS(X)$ stands for all single valued neutrosophic subsets on universal $X = \{x_1, x_2, \dots, x_n\}$.

Definition 1 (Wang et al. 2010; Smarandache 1998) A single valued neutrosophic set A on universal $X = \{x_1, x_2, \dots, x_n\}$ can be defined as follows

$$A = \{(t_A(x_j), i_A(x_j), f_A(x_j)) | x_j \in X\},$$

where $t_A(x_j), i_A(x_j), f_A(x_j) \in [0, 1]$ for each x_j in X , a truth-membership function $t_A(x_j)$, an indeterminacy-membership function $i_A(x_j)$, and a falsity-membership function $f_A(x_j)$, the sum of $t_A(x_j)$, $i_A(x_j)$ and $f_A(x_j)$ satisfies the condition $0 \leq t_A(x_j) + i_A(x_j) + f_A(x_j) \leq 3$.

Definition 2 (Wang et al. 2010) Let A, B be two single valued neutrosophic sets on universal $X = \{x_1, x_2, \dots, x_n\}$, the following relations are defined as follows:

- (1) $A \subseteq B$ if and only $t_A(x_j) \leq t_B(x_j)$, $i_A(x_j) \geq i_B(x_j)$ and $f_A(x_j) \geq f_B(x_j)$;
- (2) $A = B$ if and only $A \subseteq B$ and $B \subseteq A$;
- (3) $A \cap B = \langle \min(t_A(x_j), t_B(x_j)), \max(i_A(x_j), i_B(x_j)), \max(f_A(x_j), f_B(x_j)) \rangle$;
- (4) $A \cup B = \langle \max(t_A(x_j), t_B(x_j)), \min(i_A(x_j), i_B(x_j)), \min(f_A(x_j), f_B(x_j)) \rangle$;
- (5) $A^c = \{(f_A(x_j), 1 - i_A(x_j), t_A(x_j)) | x_j \in X\}$.

Definition 3 (Liu 2005) Let $M^{n \times n}$ be a matrix space, a mapping $\|U\|: M^{n \times n} \rightarrow R$ is called a matrix norm of U , if the following properties holds:

- (1) $\|U\| \geq 0$, for any $U \in M^{n \times n}$, $\|U\| = 0$ if and only if $U = 0$;
- (2) $\|aU\| = |a|\|U\|$, $a \in M$, for any $U \in M^{n \times n}$;
- (3) $\|U + V\| \leq \|U\| + \|V\|$, for any $U, V \in M^{n \times n}$;
- (4) $\|UV\| \leq \|U\|\|V\|$, for any $U, V \in M^{n \times n}$.

Lemma 1 (Xu and Zhang 2005) If $\|U\|$ is a matrix norm of U , then $\|-U\| = \|U\|$.

Definition 4 (Choudhary 1992) A metric space is an order pair (X, d) where X is a set and d is a metric on X , i.e., a real function $d: X \times X \rightarrow [0, +\infty)$ such that for any $x, y, z \in X$, the following holds:

- (d1) $d(x, y) \geq 0$;
- (d2) $d(x, y) = 0$, if and only $x = y$;
- (d3) $d(x, y) \leq d(x, z) + d(y, z)$.

The function d is called a distance.

We recall some existing distance between single valued neutrosophic sets on universal $X = \{x_1, x_2, \dots, x_n\}$, which are given as follows: The extended Hausdorff distance (Broumi and Smarandache 2013a):

$$d_H(A, B) = \frac{1}{n} \sum_{j=1}^n \max \{ |t_A(x_j) - t_B(x_j)|, |i_A(x_j) - i_B(x_j)|, |f_A(x_j) - f_B(x_j)| \}$$

The parametric normalized Hamming distance (Garg Nancy 2017):

$$d_{TH}(A, B) = \frac{1}{3n(2+t)} \sum_{j=1}^n \left\{ | -t|t_A(x_j) - t_B(x_j)| + |i_A(x_j) - i_B(x_j)| + |f_A(x_j) - f_B(x_j)| + ||t_A(x_j) - t_B(x_j)| - t|i_A(x_j) - i_B(x_j)| - |f_A(x_j) - f_B(x_j)|| + ||t_A(x_j) - t_B(x_j)| - |i_A(x_j) - i_B(x_j)| - t|f_A(x_j) - f_B(x_j)|| \right\}.$$

The normalized Hamming distance (Majumdar 2015):

$$d_{NH}(A, B) = \frac{1}{3n} \sum_{j=1}^n \{ |t_A(x_j) - t_B(x_j)| + |i_A(x_j) - i_B(x_j)| + |f_A(x_j) - f_B(x_j)| \}.$$

The normalized Euclidean distance (Majumdar 2015):

$$d_{NE}(A, B) = \left(\frac{1}{3n} \sum_{j=1}^n \{ (t_A(x_j) - t_B(x_j))^2 + (i_A(x_j) - i_B(x_j))^2 + (f_A(x_j) - f_B(x_j))^2 \} \right)^{\frac{1}{2}}.$$

The Gulfam Shahzadi's distance measure (Shahzadi et al. 2017):

$$d_{S_1}(A, B) = 1 - \frac{1}{n} \sum_{j=1}^n \{ \min(t_A(x_j), t_B(x_j)) + \min(i_A(x_j), i_B(x_j)) + \min(f_A(x_j), f_B(x_j)) \} / \{ \max(t_A(x_j), t_B(x_j)) + \max(i_A(x_j), i_B(x_j)) + \max(f_A(x_j), f_B(x_j)) \}.$$

$$d_{S_2}(A, B) = 1 - \left(\sum_{j=1}^n \{ \min(t_A(x_j), t_B(x_j)) + \min(i_A(x_j), i_B(x_j)) + \min(f_A(x_j), f_B(x_j)) \} \right) / \left(\sum_{j=1}^n \{ \max(t_A(x_j), t_B(x_j)) + \max(i_A(x_j), i_B(x_j)) + \max(f_A(x_j), f_B(x_j)) \} \right).$$

$$d_{S_3}(A, B)$$

$$= 1 - \frac{1}{n} \left[\sum_{j=1}^n \left(1 - \frac{1}{3} \{ |t_A(x_j) - t_B(x_j)| + |i_A(x_j) - i_B(x_j)| + |f_A(x_j) - f_B(x_j)| \} \right) \right].$$

$$d_{S_4}(A, B) = \sum_{j=1}^n \{ |t_A(x_j) - t_B(x_j)| + |i_A(x_j) - i_B(x_j)| + |f_A(x_j) - f_B(x_j)| \} / \sum_{j=1}^n \{ |t_A(x_j) + t_B(x_j)| + |i_A(x_j) + i_B(x_j)| + |f_A(x_j) + f_B(x_j)| \}.$$

The cosine distance measure (Ye 2015a):

$$d_{CS_1}(A, B) = 1 - \frac{1}{n} \sum_{j=1}^n \cos \left[\frac{\pi}{2} \max \{ |t_A(x_j) - t_B(x_j)|, |i_A(x_j) - i_B(x_j)|, |f_A(x_j) - f_B(x_j)| \} \right].$$

$$d_{CS_2}(A, B) = 1 - \frac{1}{n} \sum_{j=1}^n \cos \left[\frac{\pi}{6} \{ |t_A(x_j) - t_B(x_j)| + |i_A(x_j) - i_B(x_j)| + |f_A(x_j) - f_B(x_j)| \} \right].$$

The tangent distance measure (Ye and Fu 2016):

$$d_{T_1}(A, B) = \frac{1}{n} \sum_{j=1}^n \tan \left[\frac{\pi}{4} \max \{ |t_A(x_j) - t_B(x_j)|, |i_A(x_j) - i_B(x_j)|, |f_A(x_j) - f_B(x_j)| \} \right]$$

$$d_{T_2}(A, B) = \frac{1}{n} \sum_{j=1}^n \tan \left[\frac{\pi}{12} \{ |t_A(x_j) - t_B(x_j)| + |i_A(x_j) - i_B(x_j)| + |f_A(x_j) - f_B(x_j)| \} \right]$$

The vector distance measure (Ye 2014b):

$$d_V(A, B) = 1 - \frac{1}{n} \sum_{j=1}^n \{ (t_A(x_j)t_B(x_j) + i_A(x_j)i_B(x_j) + f_A(x_j)f_B(x_j)) / \sqrt{t_A^2(x_j) + i_A^2(x_j) + f_A^2(x_j)} \cdot \sqrt{t_B^2(x_j) + i_B^2(x_j) + f_B^2(x_j)} \}.$$

3 A new distance between single valued neutrosophic sets

In this section, we present a new distance between single valued neutrosophic sets on universal $X = \{x_1, x_2, \dots, x_n\}$.

Theorem 1 Let A, B be two single valued neutrosophic sets on universal $X = \{x_1, x_2, \dots, x_n\}$. A function $g: [0, 1] \times [0, 1] \rightarrow [0, 1]$ is a strictly monotonic binary function. The mapping $d_g: SVN S(X) \times SVN S(X) \rightarrow [0, 1]$ can be defined as follows

$$d_g(A, B; g) = \frac{1}{3n} (\|\Pi(t_A) - \Pi(t_B)\| + \|\Pi(i_A) - \Pi(i_B)\| + \|\Pi(f_A) - \Pi(f_B)\|)$$

where

$$\begin{aligned} \Pi(t_A) &= \begin{bmatrix} g(t_A(x_1), t_A(x_1)) & g(t_A(x_1), t_A(x_2)) & \dots & g(t_A(x_1), t_A(x_n)) \\ g(t_A(x_2), t_A(x_1)) & g(t_A(x_2), t_A(x_2)) & \dots & g(t_A(x_2), t_A(x_n)) \\ \vdots & \vdots & \dots & \vdots \\ g(t_A(x_n), t_A(x_1)) & g(t_A(x_n), t_A(x_2)) & \dots & g(t_A(x_n), t_A(x_n)) \end{bmatrix}, \\ \Pi(i_A) &= \begin{bmatrix} g(i_A(x_1), i_A(x_1)) & g(i_A(x_1), i_A(x_2)) & \dots & g(i_A(x_1), i_A(x_n)) \\ g(i_A(x_2), i_A(x_1)) & g(i_A(x_2), i_A(x_2)) & \dots & g(i_A(x_2), i_A(x_n)) \\ \vdots & \vdots & \dots & \vdots \\ g(i_A(x_n), i_A(x_1)) & g(i_A(x_n), i_A(x_2)) & \dots & g(i_A(x_n), i_A(x_n)) \end{bmatrix}, \\ \Pi(f_A) &= \begin{bmatrix} g(f_A(x_1), f_A(x_1)) & g(f_A(x_1), f_A(x_2)) & \dots & g(f_A(x_1), f_A(x_n)) \\ g(f_A(x_2), f_A(x_1)) & g(f_A(x_2), f_A(x_2)) & \dots & g(f_A(x_2), f_A(x_n)) \\ \vdots & \vdots & \dots & \vdots \\ g(f_A(x_n), f_A(x_1)) & g(f_A(x_n), f_A(x_2)) & \dots & g(f_A(x_n), f_A(x_n)) \end{bmatrix}, \end{aligned}$$

$\|\Pi\| = \sqrt{\lambda_{\max}}$, λ is the largest non-negative eigenvalue of positive definite matrix $\Pi^T \Pi$ (Π^T is the transpose of matrix Π) (Baker 2001). Then, d_g is a metric on $SVNS(X)$. ($SVNS(X), d_g$) is called a metric space and d_g is called a distance on $SVNS(X)$.

Proof (d1) Obviously, $d_g \geq 0$, we obtain $0 \leq g(t_A(x_i), t_A(x_j)) \leq 1$ and $0 \leq g(t_B(x_i), t_B(x_j)) \leq 1$, thus $\|\Pi(t_A) - \Pi(t_B)\| \leq n$, $\|\Pi(i_A) - \Pi(i_B)\| \leq n$ and $\|\Pi(f_A) - \Pi(f_B)\| \leq n$, then $\frac{\|\Pi(t_A) - \Pi(t_B)\|}{3n} + \frac{\|\Pi(i_A) - \Pi(i_B)\|}{3n} + \frac{\|\Pi(f_A) - \Pi(f_B)\|}{3n} \leq 1$. Therefore, $0 \leq d_g \leq 1$.

(d2) If $d_g(A, B; g) = 0$, then $\|\Pi(t_A) - \Pi(t_B)\| + \|\Pi(i_A) - \Pi(i_B)\| + \|\Pi(f_A) - \Pi(f_B)\| = 0$, by Definition 3 (1), we have $\|\Pi(t_A) - \Pi(t_B)\| = 0$, $\|\Pi(i_A) - \Pi(i_B)\| = 0$ and $\|\Pi(f_A) - \Pi(f_B)\| = 0$, then $\|\Pi(i_A)\| = \|\Pi(i_B)\|$, $\|\Pi(i_A)\| = \|\Pi(i_B)\|$ and $\|\Pi(f_A)\| = \|\Pi(f_B)\|$. Thus, we obtain $g(t_A(x_i), t_A(x_j)) = g(t_B(x_i), t_B(x_j))$, $g(i_A(x_i), i_A(x_j)) = g(i_B(x_i), i_B(x_j))$ and $g(f_A(x_i), f_A(x_j)) = g(f_B(x_i), f_B(x_j))$. Because g is a strictly monotonic binary function for each argument, so we obtain

$t_A(x_i) = t_B(x_i)$, $i_A(x_i) = i_B(x_i)$ and $f_A(x_i) = f_B(x_i)$, i.e., $A = B$.

If $A = B$, then we obtain $t_A(x_i) = t_B(x_i)$, $i_A(x_i) = i_B(x_i)$ and $f_A(x_i) = f_B(x_i)$, because g is a strictly monotonic binary function for each argument, thus $g(t_A(x_i), t_A(x_j)) = g(t_B(x_i), t_B(x_j))$, $g(i_A(x_i), i_A(x_j)) = g(i_B(x_i), i_B(x_j))$ and $g(f_A(x_i), f_A(x_j)) = g(f_B(x_i), f_B(x_j))$, then $\|\Pi(t_A) - \Pi(t_B)\| = 0$, $\|\Pi(i_A) - \Pi(i_B)\| = 0$ and $\|\Pi(f_A) - \Pi(f_B)\| = 0$, so $d_g(A, B; g) = 0$.

Therefore, $d_g(A, B; g) = 0$ if and only if $A = B$.

(d3)

$$\begin{aligned} d_g(A, B; g) &= \frac{\|\Pi(t_A) - \Pi(t_B)\| + \|\Pi(i_A) - \Pi(i_B)\| + \|\Pi(f_A) - \Pi(f_B)\|}{3n} \\ &= \frac{\|\Pi(t_A) - \Pi(t_C) + \Pi(t_C) - \Pi(t_B)\| + \|\Pi(i_A) - \Pi(i_C) + \Pi(i_C) - \Pi(i_B)\| + \|\Pi(f_A) - \Pi(f_C) + \Pi(f_C) - \Pi(f_B)\|}{3n} \\ &\leq \frac{\|\Pi(t_A) - \Pi(t_C)\| + \|\Pi(t_C) - \Pi(t_B)\| + \|\Pi(i_A) - \Pi(i_C)\| + \|\Pi(i_C) - \Pi(i_B)\| + \|\Pi(f_A) - \Pi(f_C)\| + \|\Pi(f_C) - \Pi(f_B)\|}{3n} \\ &= \frac{\|\Pi(t_A) - \Pi(t_C)\| + \|\Pi(i_A) - \Pi(i_C)\| + \|\Pi(f_A) - \Pi(f_C)\|}{3n} + \frac{\|\Pi(t_C) - \Pi(t_B)\| + \|\Pi(i_C) - \Pi(i_B)\| + \|\Pi(f_C) - \Pi(f_B)\|}{3n} \\ &= d_g(A, C; g) + d_g(B, C; g) \quad (\text{by Definition 3 (3)}) \end{aligned}$$

In short, $d_g(A, B; g) \leq d_g(A, C; g) + d_g(B, C; g)$. Therefore, d_g is a metric on $SVNS(X)$. \square

4 Application in pattern recognition

In this section, let $g(a, b) = 1 - a - b$. In order to demonstrate the effectiveness and feasibility of the proposed distance, we give a numerical example to compare the proposed distance measure with existing distance measure in Example 1. Moreover, we present a algorithm for pattern classification and medical diagnosis in Example 2–5 to illustrate the advantage of the proposed distance.

Table 1 Comparison of different distance measure in terms of counter-intuitive case (Counter-intuitive case denoted by bold)

	1	2	3	4
A_i	$\langle 0.9, 0.2, 0.3 \rangle$	$\langle 0.9, 0.2, 0.3 \rangle$	$\langle 0.8, 0.3, 0.2 \rangle$	$\langle 0.8, 0.3, 0.2 \rangle$
B_i	$\langle 0.7, 0.1, 0.2 \rangle$	$\langle 0.7, 0.1, 0.2 \rangle$	$\langle 0.7, 0.1, 0.2 \rangle$	$\langle 0.7, 0.1, 0.2 \rangle$
	$\langle 0.7, 0.2, 0.5 \rangle$	$\langle 1.0, 0.3, 0.1 \rangle$	$\langle 0.7, 0.3, 0.2 \rangle$	$\langle 0.9, 0.2, 0.1 \rangle$
	$\langle 0.6, 0.2, 0.4 \rangle$	$\langle 0.6, 0.3, 0.3 \rangle$	$\langle 0.6, 0.2, 0.3 \rangle$	$\langle 0.8, 0.2, 0.3 \rangle$
d_H (Broumi and Smarandache 2013a)	0.200	0.200	0.100	0.100
d_{NH} (Majumdar 2015)	0.133	0.133	0.067	0.100
d_{NE} (Majumdar 2015)	0.153	0.141	0.092	0.092
d_{S_1} (Shahzadi et al. 2017)	0.279	0.279	0.192	0.223
d_{S_2} (Shahzadi et al. 2017)	0.205	0.205	0.192	0.223
d_{S_3} (Shahzadi et al. 2017)	0.133	0.133	0.067	0.100
d_{S_4} (Shahzadi et al. 2017)	0.160	0.160	0.087	0.125
d_{CS_1} (Ye 2015a)	0.049	0.049	0.012	0.012
d_{CS_2} (Ye 2015a)	0.022	0.022	0.007	0.012
d_{T_1} (Ye and Fu 2016)	0.158	0.158	0.079	0.079
d_{T_2} (Ye and Fu 2016)	0.105	0.105	0.159	0.240
d_g	0.252	0.283	0.164	0.254

4.1 Numerical comparisons

Example 1 Let A, B be single valued neutrosophic sets on X , the results of the proposed distance measure compared with some existing distance measures are shown in Table 1.

For the results of Table 1, by comparing A_1, A_2 and B_1, B_2 , when $A_1 = A_2, B_1 \neq B_2$, we can see that the distance d_H (Broumi and Smarandache 2013a), d_{NH} (Majumdar 2015), $d_{S_1}, d_{S_2}, d_{S_3}$ and d_{S_4} (Shahzadi et al. 2017), d_{CS_1} and d_{CS_2} (Ye 2015a), d_{T_1} and d_{T_2} (Ye and Fu 2016) inconsistent with our intuition. Similarity, by comparing A_3, A_4 and B_3, B_4 , we can see that the distance measures d_H (Broumi and Smarandache 2013a), d_{NE} (Majumdar 2015), d_{S_1} (Shahzadi et al. 2017), d_{CS_1} (Ye 2015a) and d_{T_1} (Ye and Fu 2016) inconsistent with our intuition. Obviously, the proposed distance is not exist counter-intuitive cases. The results show that the most effective method is our proposed the new distance.

4.2 Algorithm and applications

4.2.1 Algorithm for pattern recognition

Given universe $X = \{x_1, x_2, \dots, x_n\}$, m patterns $P_j = \{\langle t_{P_j}(x_i), i_{P_j}(x_i), f_{P_j}(x_i) \rangle | x_i \in X\}$ ($j = 1, 2, \dots, m$) and a text sample $S = \{\langle t_S(x_i), i_S(x_i), f_S(x_i) \rangle | x_i \in X\}$. Which pattern does S belong to? The recognition process is as follows:

Step 1. Compute the distance measure $d(P_j, S)$ ($j = 1, 2, \dots, m$), between P_j and S .

Step 2. Select the smallest $d(P_{j_0}, S)$ from $d(P_j, S)$ ($j = 1, 2, \dots, m$), i.e., $d(P_{j_0}, S) = \min_{1 \leq j \leq m} d(P_j, S)$ and then classify the test sample S to pattern P_{j_0} .

Step 3. Compute the degree of confidence (DoC), $DoC^{(j_0)} = \sum_{j=1, j \neq j_0}^m |d(P_j, S) - d(P_{j_0}, S)|$ (Luo and Zhao 2018). If $DoC^{(j_0)}$ is greater, then the recognition result using this distance is more believable.

4.2.2 Applications in pattern recognition

Example 2 (Garg Nancy 2017) Given patterns P_1, P_2, P_3 and the test sample S as presented in Table 2. Which pattern does S belong to?

We obtain the classification results of the distances and the confidence degree of each distance in Table 3. According to Table 3, the results show that $d_g(P_2, S) \leq d_g(P_3, S) \leq d_g(P_1, S)$, i.e., the minimum distance is $d_g(P_2, S)$. Thus, it is shown that the test sample S belongs to P_2 , where the result is the same as Garg Nancy (2017). In addition, it is easy to find that d_g with the highest degree of confidence than the remaining distances in Table 3. For the results of Table 3, the distance d_g not only can classify the test sample S to pattern P_2 but also obtain a much higher degree of confidence. Therefore, the distance d_g can be seen as a reasonable application to pattern recognition.

Example 3 Given patterns P_1, P_2, P_3 and the test sample S as presented in Table 4. Which pattern does S belong to?

We obtain the classification results of the distances and the confidence degree of each distance in Table 5. According

Table 2 Patterns and test sample

	x_1	x_2	x_3	x_4
Pattern P_1	$\langle 0.7, 0.0, 0.1 \rangle$	$\langle 0.6, 0.1, 0.2 \rangle$	$\langle 0.8, 0.7, 0.6 \rangle$	$\langle 0.5, 0.2, 0.3 \rangle$
Pattern P_2	$\langle 0.4, 0.2, 0.3 \rangle$	$\langle 0.7, 0.1, 0.0 \rangle$	$\langle 0.1, 0.1, 0.6 \rangle$	$\langle 0.5, 0.3, 0.6 \rangle$
Pattern P_3	$\langle 0.5, 0.2, 0.2 \rangle$	$\langle 0.4, 0.1, 0.2 \rangle$	$\langle 0.1, 0.1, 0.4 \rangle$	$\langle 0.4, 0.1, 0.2 \rangle$
Test sample S	$\langle 0.4, 0.1, 0.4 \rangle$	$\langle 0.6, 0.1, 0.1 \rangle$	$\langle 0.1, 0.0, 0.4 \rangle$	$\langle 0.4, 0.4, 0.7 \rangle$

Table 3 The results of distance measure application in pattern recognition

Distances	$d(P_1, S)$	$d(P_2, S)$	$d(P_3, S)$	Classification results	DoC
d_H (Broumi and Smarandache 2013a)	0.375	0.250	0.125	P_2	0.375
d_{NH} (Majumdar 2015)	0.258	0.083	0.133	P_2	0.225
d_{NE} (Majumdar 2015)	0.345	0.100	0.196	P_2	0.341
d_{CS_1} (Ye 2015a)	0.215	0.021	0.101	P_2	0.274
d_{CS_2} (Ye 2015a)	0.116	0.007	0.030	P_2	0.132
d_{T_1} (Ye and Fu 2016)	0.314	0.099	0.202	P_2	0.318
d_{T_2} (Ye and Fu 2016)	0.146	0.066	0.106	P_2	0.120
d_{S_1} (Shahzadi et al. 2017)	0.473	0.246	0.349	P_2	0.330
d_{S_2} (Shahzadi et al. 2017)	0.415	0.233	0.390	P_2	0.339
d_{S_3} (Shahzadi et al. 2017)	0.258	0.083	0.133	P_2	0.225
d_{S_4} (Shahzadi et al. 2017)	0.365	0.132	0.246	P_2	0.347
d_V (Ye 2014b)	0.176	0.019	0.085	P_2	0.223
d_g	0.243	0.080	0.321	P_2	0.404

Table 4 Patterns and test sample

	x_1	x_2	x_3
Pattern P_1	$\langle 0.7, 0.5, 0.2 \rangle$	$\langle 0.8, 0.3, 0.5 \rangle$	$\langle 0.9, 0.1, 0.1 \rangle$
Pattern P_2	$\langle 0.8, 0.4, 0.1 \rangle$	$\langle 0.5, 0.3, 0.5 \rangle$	$\langle 0.7, 0.3, 0.4 \rangle$
Pattern P_3	$\langle 0.5, 0.4, 0.3 \rangle$	$\langle 0.6, 0.4, 0.6 \rangle$	$\langle 0.6, 0.2, 0.1 \rangle$
Test sample S	$\langle 0.6, 0.3, 0.2 \rangle$	$\langle 0.7, 0.2, 0.6 \rangle$	$\langle 0.8, 0.3, 0.1 \rangle$

to Table 5, d_H (Broumi and Smarandache 2013a), d_{NH} and d_{NE} (Majumdar 2015), d_{S_3} (Shahzadi et al. 2017), d_{CS_1} and d_{CS_2} (Ye 2015a), d_{T_1} and d_{T_2} (Ye and Fu 2016) cannot be determined the classification results. And the results show that $d_g(P_1, S) \leq d_g(P_3, S) \leq d_g(P_2, S)$, i.e., the minimum distance is $d_g(P_1, S)$. Thus, it is shown that the test sample S belongs to P_1 . In addition, it is easy to find that d_g with the highest degree of confidence than d_{S_1} , d_{S_2} and d_{S_4} (Shahzadi et al. 2017), d_V (Ye 2014b). For the results of Table 5, the distance d_g not only can give correct classification results but also obtain a much higher degree of confidence. Therefore, the distance d_g can be seen as a reasonable application to pattern recognition.

4.2.3 Applications for medical diagnosis

Example 4 (Samuel and Narmadhagnanam 2018; Shahzadi et al. 2017) Given three patients $P = \{\text{Ali, Hamza, Imran}\}$, five symptoms $S = \{\text{Temperature, Insulin, Blood}$

pressure, Blood plates, Cough\} and three diseases $D = \{\text{Diabetes, Dengue, Tuberculosis}\}$. The relation $P \rightarrow S$ and $D \rightarrow S$ are shown in Tables 6 and 7, respectively. What kind of disease could each patient suffer from?

The medical diagnostic results by the minimum distance d_g are shown in Table 8. Obviously, the first line shows Ali suffering from Dengue, the second line shows Hamza suffering from Diabetes, the third line shows Imran suffering from Tuberculosis. By comparing, our results are the same as Samuel and Narmadhagnanam (2018) and Shahzadi et al. (2017) in Table 9. Further, the diagnostic results based on the proposed distance are more believable than these diagnostic results in Samuel and Narmadhagnanam (2018) and Shahzadi et al. (2017).

Example 5 (Garg Nancy 2017; Ye and Fu 2016) Given four patients $P = \{p_1, p_2, p_3, p_4\}$, five symptoms $S = \{\text{Temperature, Headache, Stomach pain, Cough, Chest pain}\}$, five possible diseases $D = \{\text{Viral fever, Stenocardia, Typhoid, Gastritis, Malaria}\}$. The relation $P \rightarrow S$ and $D \rightarrow S$ are shown in Table 10 and Table 11, respectively. What kind of disease could each patient suffer from?

The medical diagnostic results by the minimum distance d_g are given in Table 12. Obviously, the first line shows p_1 , p_3 suffering from Viral fever, the second line shows p_2 suffering from Malaria, the fourth line shows p_4 suffering from Stenocardia. In order to illustrate our results, we give a com-

Table 5 The results of distance measure application in pattern recognition

Distances	$d(P_1, S)$	$d(P_2, S)$	$d(P_3, S)$	Classification results	DoC
d_H (Broumi and Smarandache 2013a)	0.167	0.233	0.167	Cannot be determined	No
d_{NH} (Majumdar 2015)	0.150	0.200	0.150	Cannot be determined	No
d_{NE} (Majumdar 2015)	0.147	0.191	0.147	Cannot be determined	No
d_{S_1} (Shahzadi et al. 2017)	0.207	0.267	0.219	P_1	0.072
d_{S_2} (Shahzadi et al. 2017)	0.205	0.267	0.214	P_1	0.071
d_{S_3} (Shahzadi et al. 2017)	0.100	0.133	0.100	Cannot be determined	No
d_{S_4} (Shahzadi et al. 2017)	0.114	0.154	0.122	P_1	0.048
d_{CS_1} (Ye 2015a)	0.037	0.069	0.037	Cannot be determined	No
d_{CS_2} (Ye 2015a)	0.012	0.022	0.012	Cannot be determined	No
d_{T_1} (Ye and Fu 2016)	0.132	0.186	0.132	Cannot be determined	No
d_{T_2} (Ye and Fu 2016)	0.079	0.105	0.079	Cannot be determined	No
d_V (Ye 2014b)	0.030	0.020	0.035	P_1	0.015
d_g	0.090	0.175	0.142	P_1	0.137

Table 6 Characteristic valued between patients and symptoms

	Temperature	Insulin	Blood pressure	Blood plates	Cough
Ali	$\langle 0.8, 0.1, 0.1 \rangle$	$\langle 0.2, 0.2, 0.6 \rangle$	$\langle 0.4, 0.2, 0.4 \rangle$	$\langle 0.8, 0.1, 0.1 \rangle$	$\langle 0.3, 0.3, 0.4 \rangle$
Hamza	$\langle 0.6, 0.2, 0.2 \rangle$	$\langle 0.9, 0.0, 0.1 \rangle$	$\langle 0.1, 0.1, 0.8 \rangle$	$\langle 0.2, 0.1, 0.7 \rangle$	$\langle 0.5, 0.1, 0.4 \rangle$
Imran	$\langle 0.4, 0.2, 0.4 \rangle$	$\langle 0.2, 0.1, 0.7 \rangle$	$\langle 0.1, 0.2, 0.7 \rangle$	$\langle 0.3, 0.1, 0.6 \rangle$	$\langle 0.8, 0.0, 0.2 \rangle$

Table 7 Characteristic valued between diseases and symptoms

	Temperature	Insulin	Blood pressure	Blood plates	Cough
Diabetes	$\langle 0.2, 0.0, 0.8 \rangle$	$\langle 0.9, 0.0, 0.1 \rangle$	$\langle 0.1, 0.1, 0.8 \rangle$	$\langle 0.1, 0.1, 0.8 \rangle$	$\langle 0.1, 0.1, 0.8 \rangle$
Dengue	$\langle 0.9, 0.0, 0.1 \rangle$	$\langle 0.0, 0.2, 0.8 \rangle$	$\langle 0.8, 0.1, 0.1 \rangle$	$\langle 0.9, 0.0, 0.1 \rangle$	$\langle 0.1, 0.1, 0.8 \rangle$
Tuberculosis	$\langle 0.6, 0.2, 0.2 \rangle$	$\langle 0.0, 0.1, 0.9 \rangle$	$\langle 0.4, 0.2, 0.4 \rangle$	$\langle 0.0, 0.2, 0.8 \rangle$	$\langle 0.9, 0.0, 0.1 \rangle$

Table 8 Diagnosis results based on the proposed distance

	Diabetes	Dengue	Tuberculosis	Diagnosis result	DoC
Ali	0.296	0.179	0.246	Dengue	0.184
Hamza	0.185	0.281	0.238	Diabetes	0.149
Imran	0.244	0.258	0.174	Tuberculosis	0.154

Table 9 Comparative results

	Ali (DoC)	Hamza (DoC)	Imran (DoC)
The result in Samuel and Narmadhagnanam (2018)	Dengue (0.174)	Diabetes (0.145)	Tuberculosis (0.137)
The result in Shahzadi et al. (2017)	Dengue (0.167)	Diabetes (0.133)	Tuberculosis (0.146)
Our result	Dengue (0.184)	Diabetes (0.149)	Tuberculosis (0.154)

Table 10 Characteristic valued between patients and symptoms

	Temperature	Headache	Stomach pain	Cough	Chest pain
p_1	$\langle 0.8, 0.2, 0.1 \rangle$	$\langle 0.6, 0.3, 0.1 \rangle$	$\langle 0.2, 0.1, 0.8 \rangle$	$\langle 0.6, 0.5, 0.1 \rangle$	$\langle 0.1, 0.4, 0.6 \rangle$
p_2	$\langle 0.6, 0.6, 0.1 \rangle$	$\langle 0.1, 0.2, 0.6 \rangle$	$\langle 0.3, 0.2, 0.8 \rangle$	$\langle 0.6, 0.2, 0.3 \rangle$	$\langle 0.2, 0.3, 0.7 \rangle$
p_3	$\langle 0.3, 0.1, 0.2 \rangle$	$\langle 0.3, 0.2, 0.2 \rangle$	$\langle 0.7, 0.6, 0.7 \rangle$	$\langle 0.3, 0.2, 0.2 \rangle$	$\langle 0.4, 0.4, 0.3 \rangle$
p_4	$\langle 0.2, 0.1, 0.7 \rangle$	$\langle 0.2, 0.3, 0.7 \rangle$	$\langle 0.2, 0.2, 0.7 \rangle$	$\langle 0.2, 0.1, 0.8 \rangle$	$\langle 0.8, 0.2, 0.1 \rangle$

Table 11 Characteristic valued between diseases and symptoms

	Temperature	Headache	Stomach pain	Cough	Chest pain
Viral fever	$\langle 0.4, 0.6, 0.0 \rangle$	$\langle 0.3, 0.2, 0.5 \rangle$	$\langle 0.1, 0.3, 0.7 \rangle$	$\langle 0.4, 0.3, 0.3 \rangle$	$\langle 0.1, 0.2, 0.7 \rangle$
Malaria	$\langle 0.7, 0.3, 0.0 \rangle$	$\langle 0.2, 0.2, 0.6 \rangle$	$\langle 0.0, 0.1, 0.9 \rangle$	$\langle 0.7, 0.3, 0.0 \rangle$	$\langle 0.1, 0.1, 0.8 \rangle$
Typhoid	$\langle 0.3, 0.4, 0.3 \rangle$	$\langle 0.6, 0.3, 0.1 \rangle$	$\langle 0.2, 0.1, 0.7 \rangle$	$\langle 0.2, 0.2, 0.6 \rangle$	$\langle 0.1, 0.0, 0.9 \rangle$
Gastritis	$\langle 0.1, 0.2, 0.7 \rangle$	$\langle 0.2, 0.4, 0.4 \rangle$	$\langle 0.8, 0.2, 0.0 \rangle$	$\langle 0.2, 0.1, 0.7 \rangle$	$\langle 0.2, 0.1, 0.7 \rangle$
Stenocardia	$\langle 0.1, 0.1, 0.8 \rangle$	$\langle 0.0, 0.2, 0.8 \rangle$	$\langle 0.2, 0.0, 0.8 \rangle$	$\langle 0.2, 0.0, 0.8 \rangle$	$\langle 0.8, 0.1, 0.1 \rangle$

Table 12 Diagnosis results based on the proposed distance

	Viral fever	Malaria	Typhoid	Gastritis	Stenocardia	Diagnosis results	<i>DoC</i>
p_1	0.177	0.191	0.206	0.262	0.264	Viral fever	0.215
p_2	0.150	0.135	0.174	0.239	0.238	Malaria	0.261
p_3	0.212	0.256	0.241	0.232	0.245	Viral fever	0.126
p_4	0.256	0.263	0.246	0.226	0.205	Stenocardia	0.171

Table 13 Comparative results

	$p_1(DoC)$	$p_2(DoC)$	$p_3(DoC)$	$p_4(DoC)$
The result in Garg Nancy (2017) ($t = 3$)	Viral fever (0.187)	Malaria (0.224)	Viral fever (0.107)	Stenocardia (0.153)
The result in Ye and Fu (2016)	Viral fever (0.169)	Malaria (0.194)	Viral fever (0.116)	Stenocardia (0.147)
Our result	Viral fever (0.215)	Malaria (0.267)	Viral fever (0.126)	Stenocardia (0.171)

parison with the exist study in Table 13. By comparing with Table 13, we can easily see that p_1 , p_3 suffering from Viral fever, p_2 suffers from Malaria and p_4 suffering from Stenocardia the same as Garg Nancy (2017) and Ye and Fu (2016). Further, the diagnostic results based on the proposed distance are more believable than these diagnostic results in Garg Nancy (2017) and Ye and Fu (2016).

5 Conclusions

Although many distance between *SVNSs* have been presented to handle uncertainty and inconsistency in various fields, most of them exist counter-intuitive cases. In this paper, we present a new distance between single valued neutrosophic sets, which overcomes the counter-intuitive cases of the existing distances. Furthermore, we applied the proposed distance in pattern recognition and medical diagnosis. Obviously, experimental results show, the proposed distance not only can be accurately classified test sample to pattern but also obtain a much higher degree of confidence. In the future, we will consider the distance based on different binary functions and matrix norms. And we will apply the distance measure to other fields, such as the multi-attribute decision making, clustering analysis.

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Data availability Enquiries about data availability should be directed to the authors.

Declarations

Conflict of interest The authors declare that they have no conflict of interest.

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