

# Determination of Yield and Yield Components of Vetch and Cereal Mixture and Evaluation Using by GGE-Biplot Analysis

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**Abstract**— This study was carried out to determine forage and seed yield and its effecting component in different vetch and cereal mixtures, ecological condition of Tekirdag-Thrace region of Turkey. The study was conducted using a total 5 vetch and cereals varieties includes three different vetch species orakefe, Hungarian vetch species sariefe, narbon vetch candidate variety 570, two different cereals (barley variety scarpia, oat variety sebat) and their mixture combinations, each species were sown as sole, double and triple mixtures (8 combinations for each genotype) a randomized complete block design with 3 replications was laid out on 2013-2015 growing season. Eight mixture combinations for each genotype (common vetch, Hungarian vetch, Narbon vetch, barley and oat) were evaluated for yield and major plant structural characteristics. Genotype-Trait (GT) biplot analyses were used. Applying type of analyses to the multiple trait data revealed that GT biplot graphically displayed the interrelationships among traits and facilitated visual comparison of mixtures and selection for each genotype. Wide variation was observed for traits plant structure and components (plant height, branch number and pod number/plant, 1000 seed weight, individual genotype ratio). It was found that; seed yield, 1000 seed weight, plant height, branch and pod number /plant were the highest value in pure stand NV570. In addition (570+sebat) had maximum dry forage yield, sariefe as pure stand has the maximum plant height and seed yield value. On the other had sariefe+scarpia combination showed very high value for pod number/plant, branch number/plant and fresh and dry forage yield. Pure stand scarpia was the best performer in seed yield. CV+NV+scarpia had the highest 1000 seed weight, fresh and dry forage yield. Maximum plant height was determined from sebat+scarpia, CV+O+B and CV+HV+B combinations. Favorable seed yield and 1000 seed weight value was produced under pure stand sebat seeding. Intercrop NV570+O combination had the higher fresh and dry forage yield. CV+HV+O, CV+NV+O and CV+B+O combination had the highest plant height of sebat.

**Keywords**— GGE-Biplot, Forage Crops, Vetch, Cereal, Yield

## I. INTRODUCTION

In the Thrace region, there are many wild relatives of cultivated plants such as wheat, cheakpea, broad bean, vetch as well as many endemic species. <http://www.fao.org/ag/agp/agpc/doc/counprof/Turkey/Turkey.htm>. Thrace is topographically very heterogeneous. Most of this region is occupied by undulating plains drained by the river Ergene. Vetches (*Vicia* spp.) are legumes well adapted to winter growth in the Mediterranean environments throughout the world on a variety of soil types and are used in west Asia, North Africa, Avustralia, and Turkey for varied purposes such as dry matter, silage and green manure [1],[4]. A number of vetch species (*Vicia* spp.) have considerable potential as grain and forage legumes since they are well adapted to the different soil and climate conditions throughout the Mediterranean environments [3], [20], [25]. In the Mediterranean Basin, including Turkey, vetches are the most common annual forage crops cultivated for hay, pasturage or silage production either alone or mixed with cereals. Hungarian vetch (*Vicia pannonica* Crantz.) is a winter hardy species, which is widely used in regions with cool winter growing conditions [2],[3]. Intercropping has been practiced traditionally in tropical regions for centuries. Interest in the intercropping of cereal-legume has been growing in many temperate and tropical regions in recent years [15], [35]. This is due to the numerous benefits. Intercropping of cereals and legumes is widely used in low-input agriculture because the mixture of nitrogen (N)-fixing and non-N-fixing crop species provides complementarities in the utilization of resources [16]. In Mediterranean countries, intercropping of cereals with legumes has been a common cropping system in short-season-rain-fed environments, especially because of their increased productivity and sustainability [26]. However, vetch has a vine growing habit and if sown as monocrop, it lodges heavily [11]. Because of having weak and thin stem, the vetches spread. So harvest gets difficult

and because of losing leaves [24], its forage and quality decrease. Vetch should be sown mixture with cereals in order to overcome spreading. As a result, forage yield and quality start to decrease due to the decomposition of herbage. Due to this, it is sown with winter-sown small grains, such as oats (*Avena sativa* L.), wheat (*Triticum aestivum* L.), barley (*Hordeum vulgare* L.) and triticale (*x Triticosecale* Wittmack). In mixtures, companion cereals provide structural support for common vetch growth, improve light interception and facilitate mechanical harvest, whereas common vetch in mixtures improves the quality of forage [28], [11], [18]. So far, there has been no report on what cereal is the best companion crop for vetch [10], [30]. [28] reported that the most suitable cereal for mixtures with common vetch is wheat. Because high cereal rate in botanic composition of legume + cereal mixture cause low protein content which is one of the most important factors to determine the forage quality [7], [4]. Use of cereals and grasses in mixtures with legumes not only increases forage yield, but also provides physical support to, which facilitates mechanical harvesting and other benefits of these mixtures include greater use of light, greater uptake of water and nutrients, enhanced weed suppression, and increased soil conservation, [24], [6]. Disadvantages of mixtures may include extra work in preparing and planting seed and lack of mixed crop tolerance to herbicides. Higher yields have been documented for intercropping oats and vetches [13]. Generally green herbage and dry matter yield and percentage of mixture decreased by increasing seed rate of legume in the mixture [19], [28], [7]. Mixtures containing 25-50 % legume produced more quality forage and yield per unit area than those of pure sowings [27], [5], [9]. [31] and [8] found the highest forage yield at the ratio 75:25 (oats: vetch) differences in soil fertility might be a factor in this response. As any of the species increased in the mixture yield gradually decreased, minimum yield was found when each species were sown alone and the yield was 55% lower than the maximum. Dry weight was affected by the common vetch ratio as was reported before [8], [31], [11]. The highest protein and digestibility contents were found in common vetch sown alone, as the proportion of oats increased there was a gradual reduction in both protein and digestibility, the lowest contents for both were with oats sown alone, with no difference ( $P > 0.05$ ) between oats cultivars. [11] also pointed out that protein content of rainfed oat-common vetch mixture depends on vetch proportion. Recently, many new researches have been conducted where vetch species were grown with different cereals in an intercropping system, whereas generally associated with oat; wheat and barley have been carried in recent years. However few related with more than double mixture have been carried in recent

years. [32] developed a site regression methodology which contains genotype main effect plus genotype by environment (GGE) using biplot procedure as a graphical tool for analysis of multi environment trials dataset. A biplot is a plot that simultaneously displays the effects of genotypes and the environment [14]. The biplot according to GGE model is a biplot that displays the GGE of multi-environment trials dataset. It is constructed by plotting the first two principal components (PC1 and PC2) derived from singular value decomposition of the environment-centered data [36]. However, it can also be equally used for all types of two-way data that assume a two way structure. The genotypes can be generalized as rows and the multiple traits as columns. [34] used a genotype by trait (GT) biplot, which is an application of the GGE biplot technique to study the genotype by trait data. The objectives of this study were to evaluate the forage production capacities of vetch and cereals mixtures, to assess the effects of different mixture levels on the yield, yield components of the mixtures. And also determining the most convenient mixture combinations and interactions in this intercropping system vetch and cereal mixtures using by GT biplot technique, under the sea coast condition of Northern Turkey.

## II. MATERIALS AND METHODS

### 2.1 Field Experiments

Plant materials, common vetch (*Vicia sativa* L. "Orakefe") Hungarian vetch (*Vicia pannonica* Crantz. "Sariefe") Narbonne vetch (*Vicia narbonensis* candidate variety "570"), Oats (*Avena sativa* variety "Sebat" and Barley (*Hordeum vulgare* variety "Scarpia") were used as plant material of the research. Each species were sown as sole, double and triple mixtures. Experiments were conducted during the growing seasons 2013-2014 and 2014-2015 at the experimental area of field crops department of Namık Kemal University Tekirdağ, Turkey. Experimental area is located at 36° 15' N and 36° 30' E direction and has a typical mediterranean climate generally. Before seeding 50 kg/ha of N and 50 kg/ha of phosphorus were applied as a starter fertilizer (Using complex fertilizer 20-20-0). Seeding was made in October of both years at same rate. Field soil was loamy structure and had low organic matter which was poor for nitrogen and phosphorus. The plots were six rows of 5m long with 25 cm row spacing. Central two rows harvested for fresh forage, the other inner two rows harvested for grain yield. The sowing rate of Common vetch "Orakefe" (CV) 100kg/ha, Hungarian vetch "Sariefe" (HV) 80kg/ha, Narbonne vetch "570" (NV) Candidate variety 150kg/ha, Barley "Scarpia" (B) 180kg/ha and Oat "Sebat" (O) 180kg/ha were sown alone as well as double and triple mixtures. The mixtures were planned and seeding %60 legumes, % 40 cereals

designated. Plot combination of each species was planned different ratio. Plant materials were sown Orakefe” (CV), Candidate variety Narbon vetch “570” Cultivar “Sariefe”(HV), Cultivar “Sebat” (O), Cultivar “Scarpia” (B) as sole crop, double combinations (CV-NV, CV-HV, CV-O and CV-B) and triple combinations (HV-NV-CV; HV-B-CV and HV-O-CV) . Yield and yield characters were determined for each cultivar in all combination individually. The experimental design was randomized complete block design with three replications combined over years. All the data were subjected to analysis of variance (ANOVA) procedures using the MSTATC statistical software package. The weed Control by hand was carried out twice when density was high, in the preflowering and postflowering stages. The meteorological data of the experimental area during growing season from October to July was given at table 1. It has a longterm rainfall of 477,9 mm during growing season, while the average temperature of the first (13.8 °C) and second growing season (13,3 °C) was higher than the long term average value (12.5°C). The total precipitation was 524.2 mm and 598,1mm during the 2013-2014 and 2014-2015 growing seasons respectively.

## 2.2 Statistical analysis

The Anderson and Darling normality test was used to normality test of dataset using MINITAB version 14 (2005) software. Cluster analysis is a tool for classifying objects into groups. The GT biplot analysis was performed on the eight cluster groups and eight for legume base and six for cereal base quantitative traits. The genotype by trait (GT) biplot method [34] was used to show the vetch+cereal mixture by trait two-way data in a biplot. These statistical methods have been described in detail by [32] and [34]. All biplots presented in this study were generated using the software GGEbiplot package [33].

### 2.2.1 Principal component analysis (PCA)

Principal component analysis (PCA) of the quantitative data was performed to investigate the importance of different characters in explaining vetch and cereals mixtures and the most commonly is useful tools for screening mixtures, and thus guide in the choice of combination for high and quality production. Biplot was used to assess the GGE biplot method [32] to assess the patterns of relations among morphological attributes, mixtures and their combinations and conducted in the dimension of first two principal components (PC1 and PC2), using a singular-value decomposition procedure.

### 2.2.2 Cluster analysis

In order to determine the similarities or differences among mixtures, a cluster analysis was performed as well. The

statistics of the cluster analysis based on the seven morpho-agronomic traits for eight different mixture combinations for legumes. In addition, six morpho-agronomic traits for eight different mixture combinations were examined for cereals.

## III. RESULTS AND DISCUSSION

The mixture combinations and evaluated of characters for each genotype were given individual as well. The following can be seen from fig. 1. The combinations of each genotype characters Orakefe (CV), 570(NV) +CV, Sariefe (HV), Sebat (O)+CV, Scarpia(B)+CV, HV+NV+CV, HV+B+CV, HV+O+CV were given an individual. The first one was given as common vetch variety and its combination can be seen from fig. 1. The GT biplot for Common vetch variety “Orakefe and its mixtures with cereals dataset, based on proposed method of [32] explained 66% of the total variation of the standardized data. This relatively moderate percentage reflects the complexity of the relationships among the measured traits. The first two principal components (PC1 and PC2) explained 44% and 22%, respectively. In the GT biplot, a vector is drawn from the biplot origin to each marker of the mixture to facilitate visualization of the relationships between and among the characters as well as mixture combinations

The vertex combinations in this investigation are (CV), (HV+NV+CV), (HV+O+CV), (B+CV), (O+CV) and (HV+CV). These combinations are the best or poorest combination in some or all the parameters since they had the longest distance from the origin of biplot. Therefore, it seems that Orakefe (CV) had the high value. Seed yield (SY), Pod number/plant (PNP) common vetch ratio (CVR) and plant height (PH) value. Sebat (O)+CV and Scarpia (B)+Sariefe (HV)+CV mixture of this sector had good amount branch number/plant (BNP) and thousand seed weight (TSW). The after combination (HV+O+CV) and related sector was not good performance for parameter HV+NV+CV, HV+B+CV and NV+CV mixtures. Which fall in its sector showed good performance for dry forage yield (DFY) and Fresh forage yield (FFY). Cluster analysis was done and a dendrogram generated from combinations. Distance matrices could distinguished clearly (similarity 66,67%) six cluster or another cluster dendrogram was done and a dendrogram generated from combination distance matrices could distinguished clearly (similarity %33) four clusters (Fig 2).

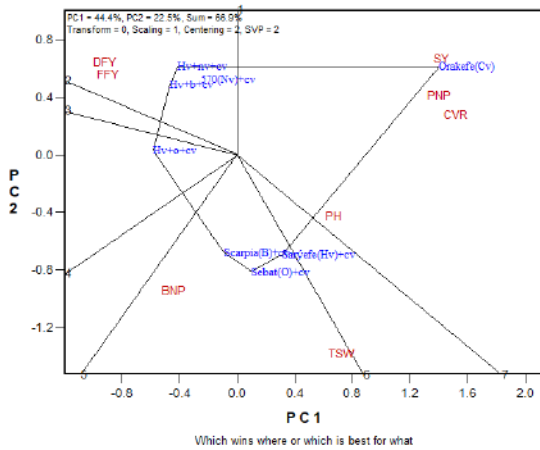


Figure 1. Polygon view, presents data of Common vetch variety “Orakefe” with 8 different mixture combinations in seven different traits.

Table 1. Correlation coefficient among seven Common vetch varieties Orakefe and its mixtures traits

Characters	BNP	PNP	SY	TSW	FFY	DFY	CVR
PH	0,081	0,056	-0,433**	0,630**	-0,239	0,093	0,373*
BNP	1	0,206	0,167	0,451**	0,162	0,117	-0,224
PNP		1	-0,191	0,137	0,241	0,273*	0,931**
SY			1	-0,097	-0,038	-0,452**	-0,401**
TSW				1	-0,123	0,053	-0,368*
FFY					1	0,698**	0,888**
DFY						1	0,245
CVR							1

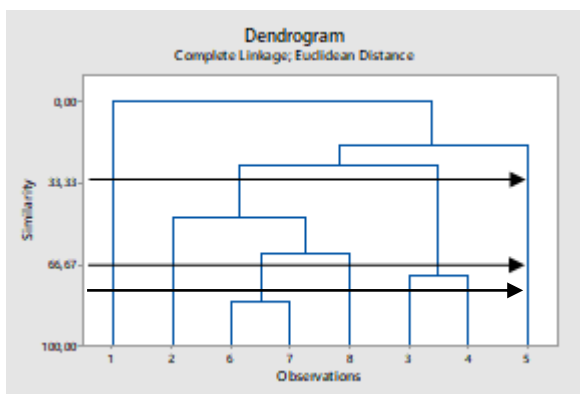


Figure 2. Clusters dendrogram of Common vetch variety Orakefe with 8 different mixture combinations in seven different traits

The number of clusters was verified by multivariate analysis of variance test. This relatively moderate percentage reflects the complexity of the relationships among the measured parameters. Thus reducing entry numbers of mixtures from 8 combinations to 6 clusters.

Additional, 6 clusters are necessary to explain 66.67% similarity of the mixture combinations.

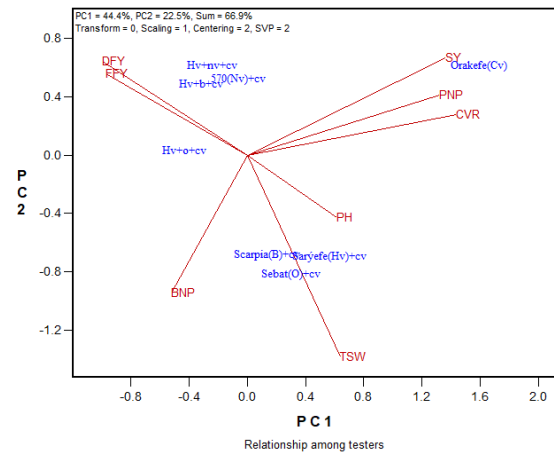


Figure 3. Vector view mixtures by trait biplot, showing the interrelationship among measured traits for Common vetch variety Orakefe with 8 mixtures combinations

In the Fig. 3, the correlation coefficient between any two traits is approximated by the cosine of the angle between the vectors. The correlation coefficients among the traits indicate that the biplot currently shows relationship among the traits that had relatively large loading on both PC1 and PC2. Therefore, the most prominent relations by this figure (8 mixtures and seven traits) are: a strong positive association among SY and PNP, among SY and CVR, among DFY and FFY had positive correlation as well. There was a near zero correlation between BNP and TSW with SY, PNP and CVR (Fig. 3) as indicated by the near perpendicular vectors ( $r = \cos 90 = 0$ ). Some of mentioned results can be verified using correlation coefficients of Table. 1.

The GT biplot for Narbon vetch candidate variety (NV570) and its mixtures with cereals based on method of [34] explained 66% total variation of standardized data. The principle component vectors (PC1 and PC2) explained 61.4 % and 18.1 %, sum of the vectors was 79.5%. In the GT biplot, this view helps identify mixtures with highest values for one or more traits. The scores of six traits seed yield (SY), plant height (PH), narbon vetch ratio (NVR), tausand seed weight (TSW), pod number /plant (PNP) and branch number /plant (BNP) fell in the NV570 sector. NV570 had highest and nearest highest values for this traits. Orakefe+NV570 combination was second to NV570 for these traits. Similarly combination of sebat+NV570 mixture was the highest in fresh and dry forage yield, other mixture combination. CV+B+NV570 had the highest fresh forage yield. Since the biplot did not explain all variation of these predictions may not exactly reflect the observed values.

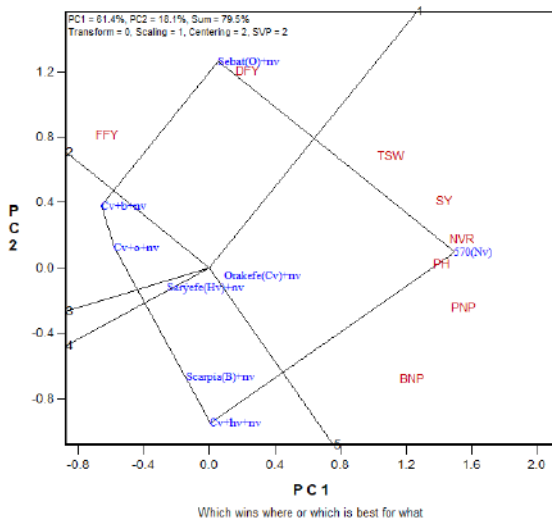


Figure 4. Polygon view, presents data of “Narbon vetch candidate variety” with 8 different mixture combinations in seven different traits

Fig. 5 displays the tree diagram. The figure provides a graphical view of the clusters. Cluster analysis was approved as a suitable method for data classifying and suggested by [22]. Based on the cluster analysis in Fig. 5, we can divide the 8 mixture combinations can divide into 6 clusters based on the studied agronomic characters. In other words, 6 clusters are necessary to explain 66.67% similarity of the mixture combinations.

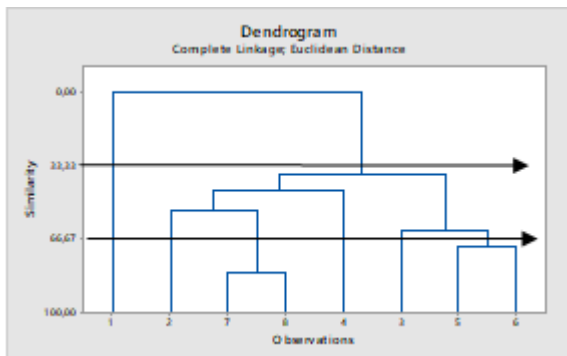


Fig. 5. The cluster dendrogram of Narbon vetch candidate variety with seven different traits of eight mixture combinations.

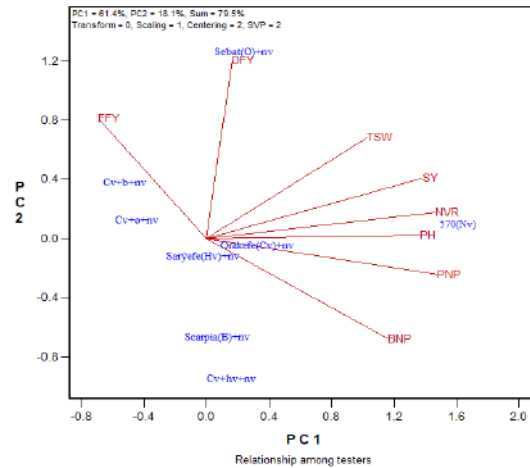


Figure 6. Vector view mixtures by trait biplot, showing the interrelationship among measured traits for Narbon vetch candidate variety with 8 mixtures combinations.

The correlation coefficients among the seven traits are presented in Table 2. The vector view of a GGE biplot provides a succinct summary of the interrelationships among the traits [34]. Figure 6 is the vector view of the GGE biplot, in which the mixture combination is connected with the biplot origin via traits. This view of the biplot aids understanding of the interrelationships among mixture combination. The cosine of the angle between the vectors of two traits approximates the correlation coefficient between them. Therefore, the most prominent relations were: (a) near-zero correlations between TSW and DFY, between PNP and TSW, and between BNP and DFY as indicated by the near-perpendicular vectors ( $r = \cos 90 = 0$ ); and (b) positive associations among PH, NVR, PNP and SY, and between PNP and BNP as indicated by acute angles.

Table 2. Correlation coefficient among seven traits of Narbon vetch candidate variety 570 and its mixtures combinations

Character s	PH	BNP	PNP	SY	TSW	FFY	DFY	NVR
PH	1	0,511**	0,651**	0,724**	0,168	-0,357**	-0,049	0,868**
BNP		1	0,561**	0,441**	0,018	-0,438**	-0,110	0,647**
PNP			1	0,804**	0,269	-0,385**	0,041	0,960**
SY				1	0,334*	-0,314*	0,088	-0,409**
TSW					1	0,074	-0,006	0,226
FFY						1	0,612**	0,948**
DFY							1	0,732**
NVR								1,000

The GT biplot for Hungarian vetch “Sariefe” and its mixtures with cereals based on method of [34] explained

66% total variation of standardized data. The principle component vectors (PC1 and PC2) explained 41.8 % and 25.7 %, sum of the vectors was 67.5%. The biplot representing a polygon view (Fig. 7), having some vertex mixtures while the rest are inside the polygon. These vertex mixtures are supposed to be the most responsive since they have the longest distance from the biplot origin. Responsive mixtures are either best or poorest at one or all mixtures [34]. Thus it seems that Scarpia (B) +HV and CV+ Scarpia (B) +HV mixture combinations had the highest values for DFY, BNP, PNP, FFY and TSW. The other vertex mixture combination which fall in its sector were good for plant height (PH), Hungarian vetch ratio (HVR) and seed yield (SY). The other vertex mixtures (CV+HV and CV+NV+HV) and related sectors were not good performance for the measured traits.

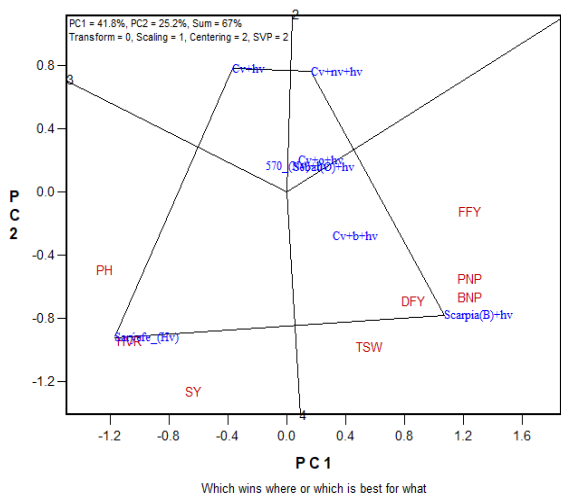


Figure 7. Polygon view, presents data of Hungarian vetch variety Sariefe with 8 different mixture combinations in seven different traits.

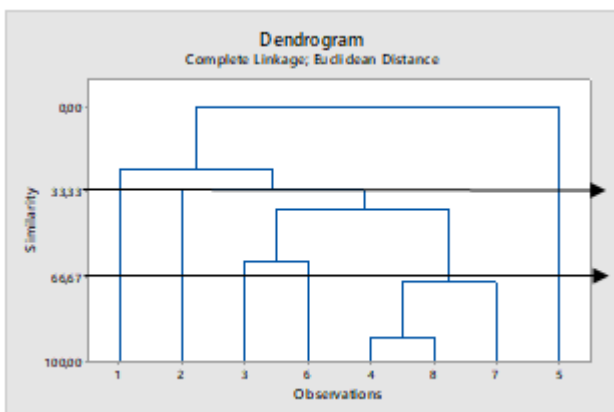


Figure 8. The cluster dendrogram of Hungarian vetch variety Sariefe with seven different traits of eight mixture combinations

Based on the cluster analysis in Fig. 8, we can divide the 8 mixture combinations can divide into 4 clusters based on the studied agronomic characters in similarity level 33.33 % of mixture combination. In other words, 4 clusters [1, 2, (3,6), (4, 7, 8), 5] are necessary to explain 33.33% similarity of the mixture combinations.

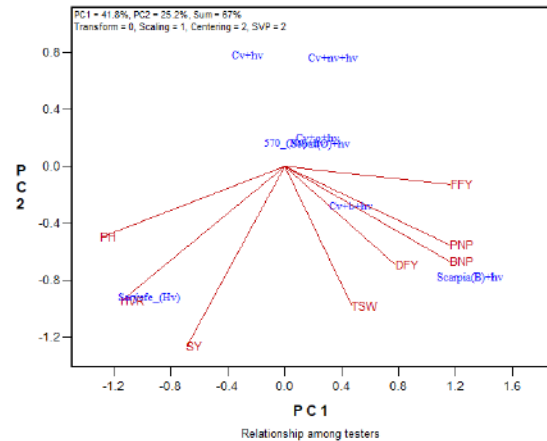


Figure 9. Vector view mixtures by trait biplot, showing the interrelationship among measured traits for Hungarian vetch variety Sariefe with 8 mixture combinations.

The mixtures-traits vectors and groups illustrate the specific interactions of each mixture with each trait (Fig 9). The biplot showed DFY that was highly correlated in terms of PNP (Fig.9). Positive correlations were found between FFY (Fresh forage yield), DFY (Dry forage yield), PNP (Pod number/plant) and BNP (Branch number/plant), as indicated by the acute angles (vector angles < 90°) of their respective vectors. There was a near zero correlation between BNP, PNP and FFY with PH (fig 7) as indicated by near perpendicular vectors ( $r = \cos 90 = 0$ ). The biplot of the relationship between traits negative correlation was observed between BNP and PNP with PH as indicated by the large obtus angles between their vectors. Having same result observed from Table 3.

Table 3. Correlation coefficient among seven traits of Hungarian vetch variety Sariefe and its mixtures combinations

Charact ers	P H	BN P	PNP	SY	TS W	FF Y	DFY	HVR
PH	1	-0,459**	-0,497**	0,339*	0,020	-0,035	0,144	0,848**
BNP		1	0,637**	0,167	0,029	0,348**	0,273	-0,350**
PNP			1	0,222	0,108	0,414**	0,419**	-0,278
SY				1	0,233	-0,030	0,128	-0,352**
TSW					1	0,085	0,081	0,084
FFY						1	0,939**	0,747**
DFY							1	-0,018
HVR								1

The GT biplot can be used to compare mixture combination on the basis of multiple traits and to identify mixture that are particularly good in certain aspects and therefore can be candidates for high production in mixture sowing. The GT biplot for oat variety “Scarpia” and its mixtures with cereals based on method of [34] explained 66% total variation of standardized data. The principle component vectors (PC1 and PC2) explained 39.9 % and 28.2 %, sum of the vectors was 68.8%. The perpendicular lines to the polygon sides facilitate comparison between neighboring mixtures. Specifically, comparison between CV+NV+ (Scarpia) B and (Scarpia) B neighboring vertex mixtures. According to [34] we have used the GT biplot to compare mixture combinations on the basis of multiple traits and to identify mixtures or groups of mixtures that are particularly good in certain aspects and therefore can be determined the best combination in vetch species+ cereal mixtures. The equality lines, which originate from the centre of biplot and are perpendicular to the sides of polygon, divide the graph into four sectors. The partitioning of GT interaction through GT biplot analysis shows that the first and second principal components (PC1 and PC2) together can explain 68% of the total variation. The yields of these mixtures were either the highest or lowest in one or more test environments. From the polygon view of biplot analysis, the genotypes fell into four sections and the test environments can be grouped in three sections (Section 1, 2 and 3). CV+NV+ B (Scarpia) combination were the highest thousand seed weight (TSW), fresh forage yield (FFY) and Dry forage yield (DFY). In addition Barley variety (Scarpia) was the highest seed yield (SY) and barley ratio value (Fig. 10). On the otherhand these vertex pure stant Scarpia and its mixtures, NV570+B, HV+B and CV+B were the highest value of Barley ratio and seed yield. (Fig 10). Highest plant height of Scarpia (B) was recorded on O+B;CV+O+B and CV+HV+B mixture combinations.

Cluster analysis was given in fig 11. According to the result of Scarpia combination can divided in to 3 clusters based on studied. Five mixture combinations were extracted from the 8 studied combinations. In fact, with this method, 8 mixture combinations were reduced to three in 33.33 % similarity level of the mixture combination (Fig. 11). Another explanation, Three clusters [(1, 2, 3, 7, 8), (4, 5), 6] are necessary to explain %33 similarity level of the mixture combinations.

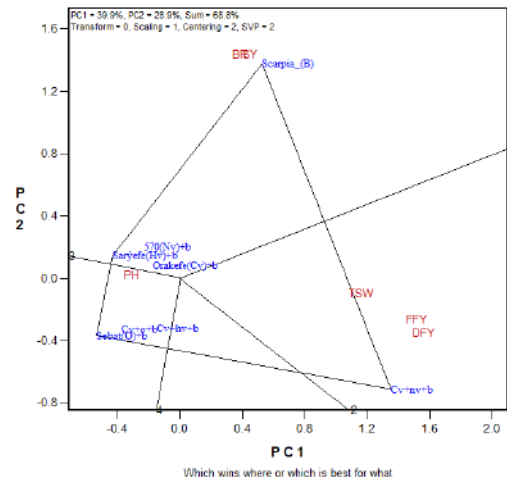


Figure 10. Polygon view, presents data of “Oat variety Scarpia” with 8 different mixture combinations in seven different traits

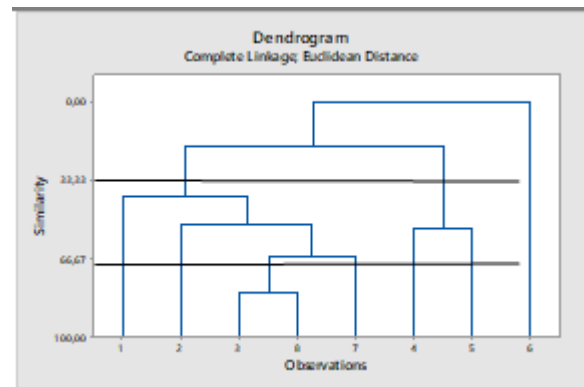


Figure 11. The dendrogram of Oat variety Scarpia with seven different traits of eight mixture combinations using hierarchial cluster analysis (Ward’s method and squared Euclidean distance

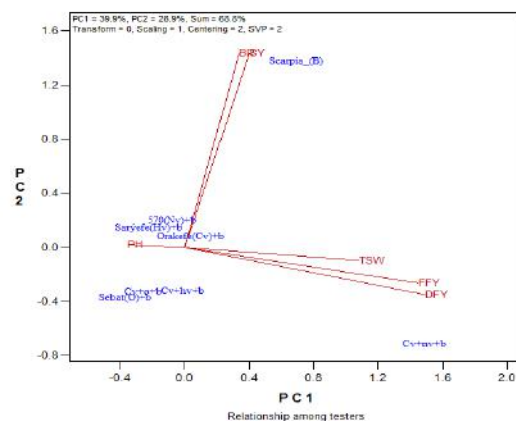


Figure 12, Vector view mixtures by trait biplot, showing the interrelationship among measured traits for “Oat variety Scarpia” with 8 mixtures combinations.

The correlation between these traits supports this theory that plant height and Scarpia ratio can improve Scarpia seed yield. The correlation coefficient between Scarpia seed yield and TSW is positive. TSW is one of the important components for increasing yield production, so, this is normal and expected. There is a positive and significant correlation between TSW and FFY; DFY. It is seen a close and positive association between FFY and DFY traits in many studies. The GT biplot for Barley variety “Sebat” and its mixtures with cereals based on method of [34] explained 66% total variation of standardized data. The principle component vectors (PC1 and PC2) explained 39.9 % and 28.2 %, sum of the vectors was 68.8%. The biplot representing a polygon view (Fig. 1). Having some vertex hybrids while the rest are inside the polygon. These vertex hybrids are supposed to be the most responsive since they have the longest distance from the biplot origin. Responsive hybrids are either best or the poorest at one or all locations [34]. Thus, hybrid Sebat was the highest seed yield and thousand seed weight and sebat ratio (OR). Similarly, in the next sector where the vertex 570(NV)+O excelled for Fresh and Dry forage yield. In addition sector where the vertex combination CV+(HV)+O followed by CV+(NV)+O and CV+B+O excelled for plant height. None of the traits fell in the sectors with Sariefe (HV)+O and Orakefe (CV)+O as the vertices genotypes, indicating that these mixture combinations were not best in any of the trait, Sebat (O) and 570(NV)+O combinations seemed to be highest- yielding for seed yield, thousand seed weight, Fresh and Dry forage yield in two mentioned combinations.

Table 4. Correlation coefficient among seven traits of Scarpia and its mixtures combinations

Characters	PH	SY	TSW	FFY	DFY	BR
PH	1	0,319*	0,267	0,102	0,017	-0,247
SY		1	0,326*	-0,099	0,011	0,139
TSW			1	0,288*	0,321*	0,029
FFY				1	0,886**	0,760**
DFY					1	-0,158
BR						1

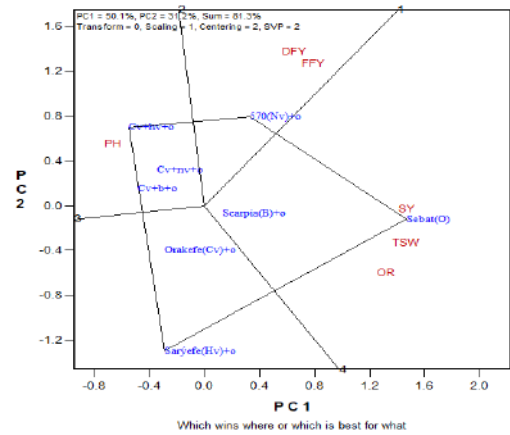


Figure 13. Polygon view, presents data of “Barley variety Sebat” with 8 different mixture combinations in seven different traits.

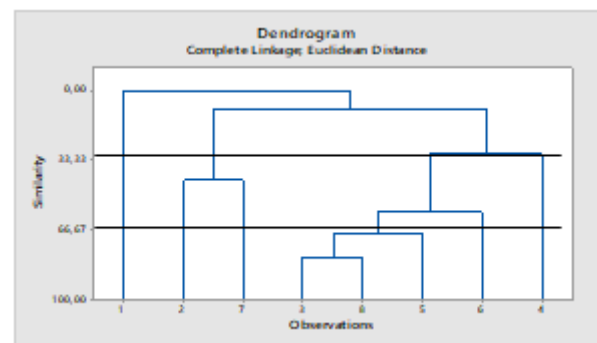


Fig. 14. The cluster dendrogram of Barley variety Sebat with seven different traits of eight mixture combinations.

Analysis of the results showed that the cultivar was clustered in to four main clusters in 33.33 % similarity level (Fig. 14). In the second cluster were included 1 (NV)+O and 7 (CV+HV+O) and third clusters were included 3 (CV+O), 8 (CV+B+O); 5 (B+O); 6 (CV+NV+O) which were similar each other. Four mixture combinations were extracted from the 8 studied combinations. In fact, with this method, 8 mixture combinations were reduced to four in 33.33 % similarity level (Fig. 14).

In GGE-biplot, the cosine of the angle between any two location vectors stands for correlation intensity. Less than 90° indicates a positive correlation, more than 90° a negative correlation, and close to 90° no correlation [37], [29]. According to Fig. 15, there were positive relationships between SY and TSW, FFY and DFY, PH and FFY, PH and DFY. TSW and OR. On the other hand, the negative correlations were observed between SY and PH, TSW and PH, OR and PH (Fig. 15).



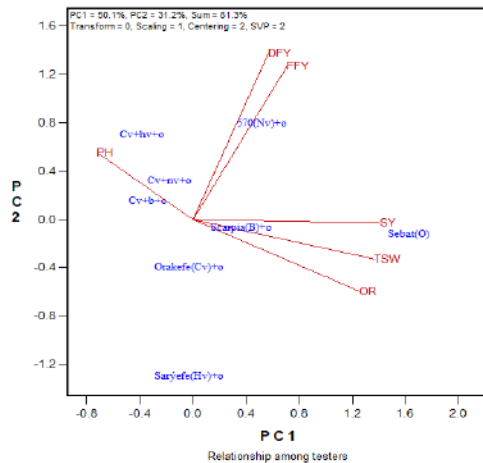


Fig. 15, Vector view mixtures by trait biplot, showing the interrelationship among measured traits for “Barley variety Sebat” with 8 mixtures combinations.

Table 5. Correlation coefficient among seven traits of Barley variety Sebat and its mixtures combinations

Characters	PH	SY	TSW	FFY	DFY	OR
PH	1	-0,334*	-0,097	0,085	0,165	-0,319*
SY		1	0,680**	0,357*	0,315*	0,737**
TSW			1	0,288*	0,321*	0,907**
FFY				1	0,886**	0,088
DFY					1	-0,095
OR						1

#### IV. CONCLUSION

The GT biplot analysis of 8 different combinations showed variable ranges of purestand and mixtures of vetches and cereals. According to polygon view of traits and combination; seed yield, pod number /plant and plant height were found to be heighest value from pure stand Orakefe. Seed yield, tausand seed weight, planth height, Branch and pod number /plant were the heighest value in pure stand NV570. In addition NV570+sebat had maximum dry forage yield. Sariefe as purestand has the maximum planth height and seed yield value. On the other hand Sariefe+Scarpia combination showed very high value for pod number/plant, branch number/plant, and fresh and dry forage yield. Purestand Scarpia was the best performer in seed yield. CV+NV+Scarpia had the highest tausand seed weight, fresh and dry forage yield. Maximum planth height was determined from Sebat+ scarpia, CV+O+B and CV+HV+B combinations. Favurable seed yield and tausand seed weight value was produced under purestand sebat seeding. Intercrop NV570+O combination had the higher fresh and dry forage yield. CV+HV+O, CV+NV+O

and CV+B+O combination had the highest plant height of sebat.

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