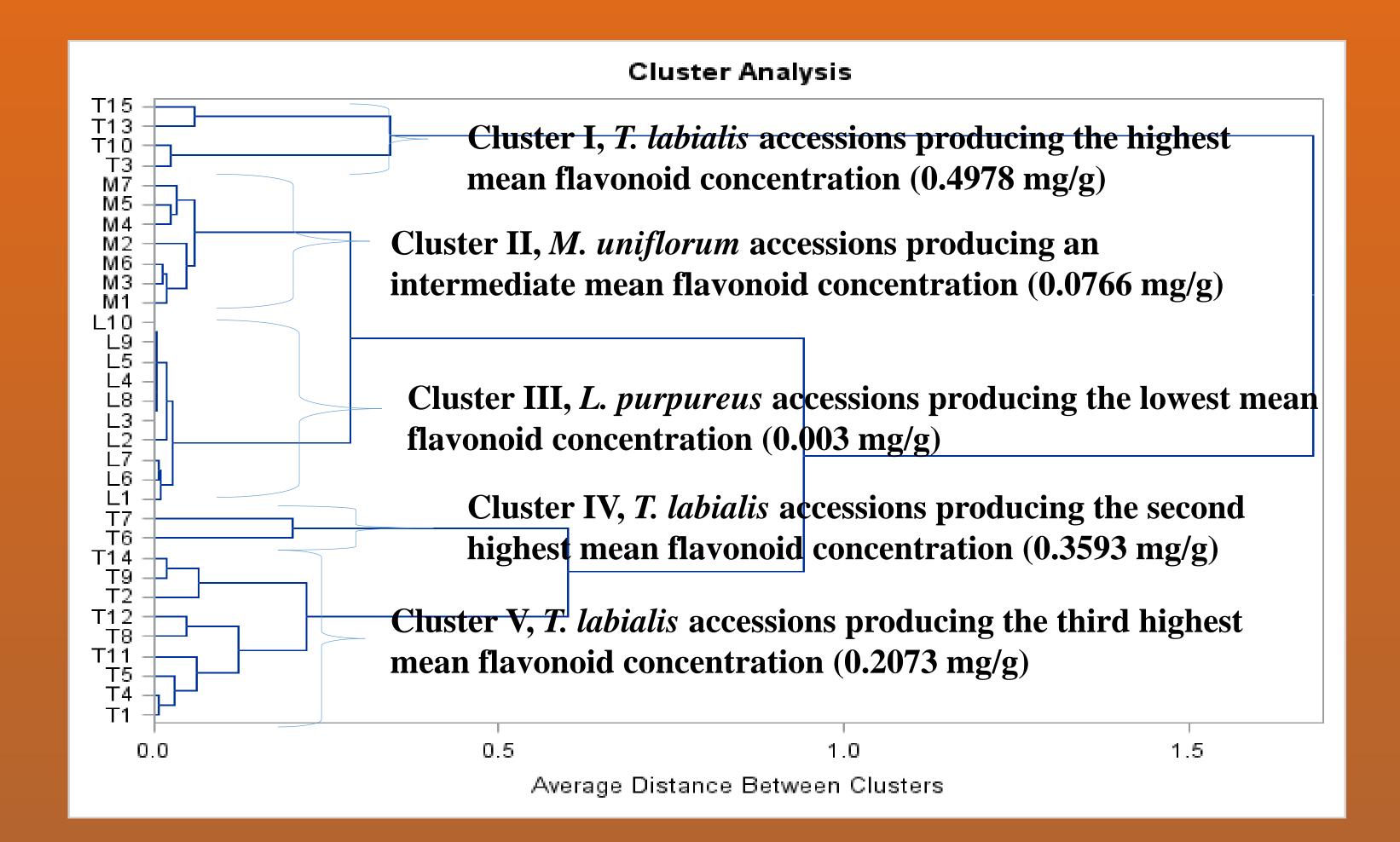
Flavonoid Concentration Diversity of 3 Different Species in the Phaseoleae Tribe

J.B. Morris and M.L. Wang, USDA, ARS, PGRCU, Griffin, GA, 30223, U.S.A. (Brad.Morris@ars.usda.gov)



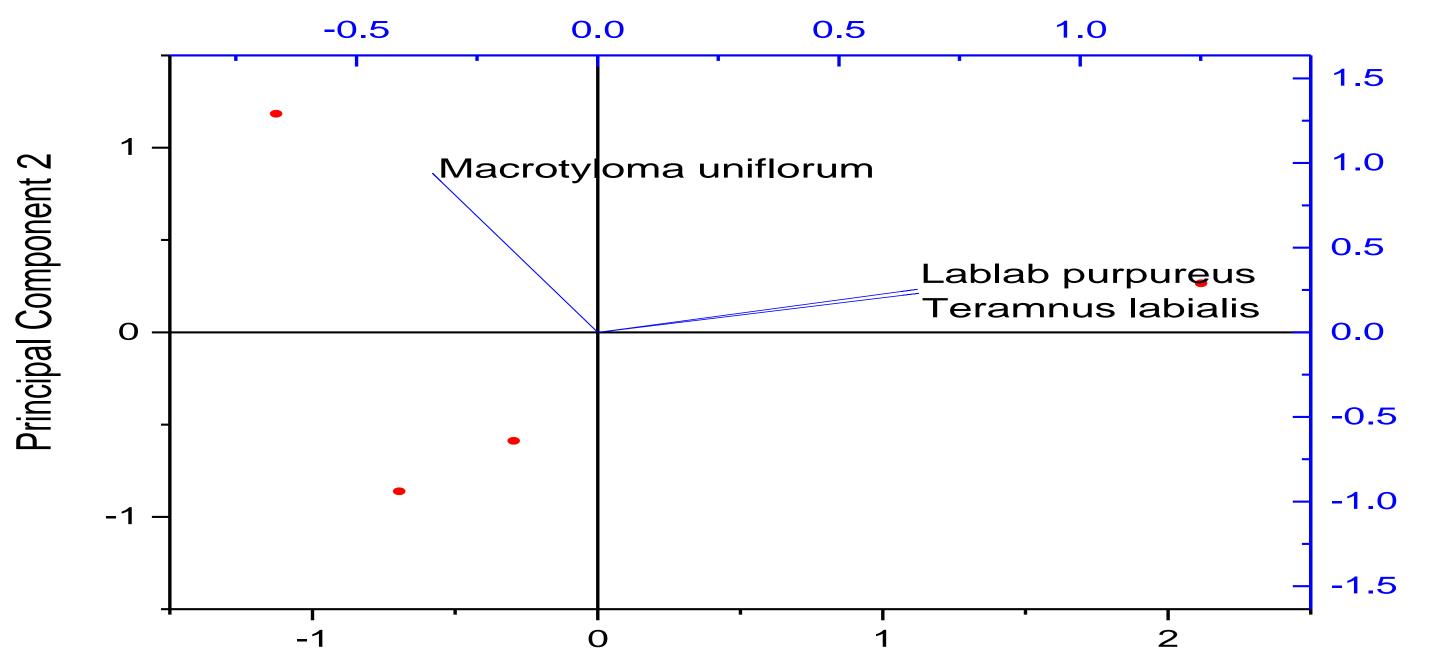
Abstract

The functional vegetable species including *Lablab purpureus* L. Sweet, Macrotyloma uniflorum (Lam.) Verdc., and Teramnus labialis (L.f.) Spreng are in the *Phaseoleae* tribe. Seeds from 10 L. purpureus and 7 M. uniflorum accessions originated from the field during 2009 and 2009 to 2010, respectively at Griffin, GA. However, seeds from 15 T. labialis accessions originated from greenhouse grown plants because of photoperiod sensitivity. Flavonoids have been shown to have anti-cancer potential. The flavonoid data from all three species were subjected to an ANOVA, mean separation, cluster, and principal component analysis. Significant flavonoid concentrations were identified from all species. Overall, T. labialis seeds produced significantly more quercetin (ranging from 0.616 – 2.12 mg/g) than the other 2 species. The cluster analysis showed 5 distinct clusters for flavonoid content in the 3 species. The *Teramnus labialis* accessions showed 3 clusters with the highest flavonoid concentrations (ranging from 0.2073 to 0.4978 mg/g). The *M. uniflorum* accessions produced an intermediate concentration of flavonoids (0.0766 mg/g), and the lowest flavonoid concentrations were observed in the L. purpureus accessions (0.0030) mg/g). The principal component analysis showed that the first and second components accounted for 70 and 28% of the total variation. *Teramnus labialis* and L. purpureus were high in principal component 1 with coefficients of 0.6659 and 0.6626, respectively. However, *M. uniflorum* was high in principal component 2 with a coefficient of 0.9393. The T. labialis accessions produced superior flavonoid concentrations relative to the other two species. Separate clusters for L. purpureus and T. labialis were observed in principal component **1.** A common cluster for *M. uniflorum* was observed in principal component **2**.



Materials and Methods

17 accessions of *Lablab purpureus* L. Sweet and *Macrotyloma uniflorum* (Lam.) Verdc. were grown in the field from 2009 to 2010. However, 15 *Teramnus labialis* (L.f.) Spreng accessions were grown in the greenhouse during 2009 because of photoperiod sensitivity. All seeds were harvested at maturity. Dry and mature seeds from each genera and accession were analyzed for flavonoid content using a similar reverse phase HPLC technique. See Morris et al., 2013a, 2013b, and 2014 for details. Published data from (Morris 2013a, 2013b, and 2014) was reanalyzed using mean separations from ANOVA as well as principal component and cluster analysis to identify which taxon produced superior flavonoid content. Dendrogram of the distance between clusters based on flavonoid concentration differences. Species are given on the y-axis (T, M, and L represent *Teramnus labialis*, *Macrotyloma uniflorum*, and *Lablab purpureus*, respectively). Values on the x-axis indicate average flavonoid distances between the species. 5 clusters for flavonoid concentrations can be distinguished.



Results and Discussion						
	<i>L. purpureus</i> (n=10)	<i>M. uniflorum</i> (n=7)	<i>T. labialis</i> (n=15)			
Flavonoid	Range (mg/g)					
Myricetin Quercetin Kaempferol Isorhamneti		0c – 0.036a 0g – 0.0272ab 0.2403c – 0.3155a Not detected	Not detected 0.616fg – 2.12fg 0d – 0.07d 0b – 0.086a			

Principal Component 1

	Eigenvalue	Percentage	of Variance	Cumulative
1	2.10491	70.16%		70.16%
2	0.85323	28.44%		98.60%
3	0.04186	1.40%		100.00%
	Coefficients	of PC1	Coefficients	of PC2
Teramnus labialis		0.66597		0.23062
Lablab purpureus		0.66261		0.25399
Macrotyloma uniflorum		-0.34269		0.93931

References

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