Phenotypic Variation in Adaptive Traits Among Western Populations of Showy Milkweed (Asclepias speciose) Michaela Sonnen, Daison Weedop, Sacha Wells, Leslie Blackburn, Dusty Perkins College of Western Idaho College of Western Idaho

INTRODUCTION

•Losses of milkweeds in monarch butterfly (*Danaus plexippus*) breeding habitats and migratory corridors are hypothesized to be a chief contributor to widespread declines in monarch populations across North America (Pleasants and Oberhauser 2013, Flockhart et al. 2015). • Successful habitat restoration requires germplasm that is adapted to target restoration environments.

•Genecology evaluates variation in putative traits related to survival among populations in relation to source environments and local selection pressures (St. Claire et al. 2013). • Previous approaches suggest local seeds are best, but may not be available in suitable quantities, or may not be adapted to projected climate change scenarios (Kilkenny 2015). •We evaluated phenotypic variation in growth form traits among 36 populations of showy milkweed (Asclepias speciosa) to evaluate genetically appropriate adaptive traits among local climates to aid in identifying seed sources for monarch habitat restoration.



METHODS

Plant Propagation and Transplanting

• Procured seeds from 36 populations (Fig. 1) •BLM, USFWS germplasm accessions

•Citizen scientist collections

•Seeds were sowed in February in plug trays. Seedlings were transplanted to pots in April and transplanted in the common garden in late May.

 $\circ 0.5$ m between plants, 1 m between rows

• Prior to planting, the site was treated with herbicide, and covered in weed barrier and crushed gravel to reduce weed competition and heat loads.

•After planting, the garden received supplemental irrigation and was weeded on a weekly basis.

•Among the 360 experimental plants (ten per a population), we used 158 nonexperimental units as buffer rows and columns, and to spaces in the experimental layout. **Data Collection**

•Weekly collection of morphological measurements and plant growth data (Table 1). •Remotely sensed climate data retrieved from ClimateWNA v5.30 (Wang et al. 2012).

Analysis

• Principal components analysis:

•Conducted in program R version 3.5.1 using prcomp () function in package (stats)

•We used single factor ANOVAs to determine which measured traits significantly varied among populations.

• Tukey's HSD was used to determine where the variance was among populations at a significance level of $\alpha = 0.05$.

•Relationships between collections site characteristics and measured traits were analyzed with a simple linear regression models (Table 1).





Fig. 1- Locations of 36 populations of showy milkweed used in this study. Fig. 2- Principal components analysis separations among 36 populations of Showy Milkweed. Fig. 3- Biplot with principal components analysis separations with dimensional loading factors among 36 populations of Showy Milkweed. Table 1- Environmental variables and phenotypic characteristics used in analyses. Fig. 4a- Regression analysis of MWMT (°C) and plant height. Fig. 4b- Regression analysis of MAT (°C) and plant height. Table 2- Relative contributions of plant phenotypic characteristics (%) to top two principle components.



Environmental Variables	Description					
MAT (°C)	Mean average temperature					
MWMT (°C)	Mean warmest month temperature					
MSP (mm)	Mean annual summer (May to Sept.) precipitation					
MAP (mm)	Mean annual precipitation					
Elevation (m)	Site elevation					
Phenotypic Characteristics	Description					
Leaf Length (cm)	Length of the most mature leaf from petiole to terminal tip					
Leaf Width (cm)	Widest point of the most mature leaf					
Est. Leaf Area (cm ²)	Product of Leaf length x Leaf width					
Leaf L/W Ratio	Quotient of Leaf length / Leaf width					
Plant Height (cm)	Height measurement from the emergence from soil					
	to the top of the plant					
Canopy Area (cm ²)	Maximum distance of the canopy extent (cm) multiplied					
., . ,	by the maximum canopy extent of perpendicular axis.					
Volume (cm ³)	Plant Height multiplied by canopy area.					
Total Stem Length (cm)	Sum length of the three longest stems					
Number of stems	Total number of basal and axillary stems					

Deg. F._ /_ 6-12_C/m_Precip. Deg. F. / 3-6 C/m Precip. Deg. F. / 6-12 C/m Precip. 15_Deg._F./_3-6_C/m_Precip. 15-20_Deg._F._/_3-6_C/m_Precip. 15-20_Deg._ F._ /_ 6-12_C/m_Precip. 20-25_Deg._ F._ /_ 12-30_C/m_Precip 20-25_Deg._ F._ /_ 6-12_C/m_Precip. 25-30_Deg._ F._ /_ 12-30_C/m_Precip 25-30_Deg._ F._ /_ 6-12_C/m_Precip. 25-30_Deg._ F._ /_3-6_C/m_Precip. 30-35_Deg._ F._ /_ < 2_C/m_Precip.</p> 8 30-35_Deg._ F._ /_2-3_C/m_Precip. 35-40_Deg._ F._/_ 3-6_C/m_ Precip.



Fig. 3



•79% of variation amo
OPC1 associated
OPC2 associated
•Mixed evidence that p
have similar trai
 Some populat
■ 20-25
■ 15-20°
 Some populat
■ 25-30°
■ 25-30°

Table 2									
		No. of	Total Stem		Leaf	Leaf	LW	Leaf	Canopy
	Volume	Stems	Length	Height	Length	Width	Ratio	Area	Area
Dimension 1	15.8	1.3	10.8	14.2	12.8	13.3	1.9	14.8	15
Dimension 2	1.3	8.6	6.4	0.01	4.1	14.3	55.5	4.8	4.9

•ANOVAs revealed significant ($\alpha = 0.05$, df=35) effect of populations on all measured traits •Statistical differences driven mostly by 3-4 populations with significantly different growthforms. ■OR8, OR1, WY2, CA1

•MWMT(°C) and MAT (°C) show significant positive relationships with plant height (Figs. 4a, 4b).

western US.

characters (Fig. 2).

•Regressions across individuals nested within populations (not shown) indicate that the bulk of variation in traits exists among individuals. •Phenotypic plasticity may play an important role in showy milkweed responses to annual weather conditions (Sultan 2000).

•Environmental maternal effects may also serve as a source of variation in evaluating trait responses between generations(Galloway 2005, Donohue 2009). • Regression analyses indicated that plant height may be adaptive. • Higher MWMT(°C) and MAT (°C) correlated with increased height

•Cool, wet spring in 2018 may be responsible for large divisions in growth form. • Variation among principal components does not strongly correspond to

provisional seed zones.

• Closely clustered populations suggest some populations from common seed zones may have similar character traits (Fig. 3).

• Multi-year study (\leq three years)

•Examine population genetics.

• Explore variations in cardenolide toxins with respect to ecotypes and stress.

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RESULTS

ng populations is explained by two Principal Components (Fig. 2) with larger and greater surface areas, volumes and growth traits with lower LW ratio, reduced leaf width, and fewer stems. populations from common provisional seed zones (Bower et al. 2014)

its (Fig. 3)

tions from common seed zones cross both PCs

°C, 6-12 mm

°C, 6-12mm

tions cluster closely °C, 6-12mm

°C. 3-6mm

■ 20-25°C, 12-30mm



DISCUSSION

• Preliminary support that Showy milkweed exhibits genecological differences across

•Patterns do not clearly correspond to provisional seed zones, suggesting that species specific seed zones should be constructed for showy milkweed.

•Predominate divisions among populations due to differences in growth form and leaf

•Plant size often correlates to competitive ability (Westoby 1998).

FUTURE WORK

•Incorporate measures of pubescence, above ground biomass, precise leaf surface area, surface area, emergence and flowering phenology.

• Explore correlations with additional climate variables in ClimateWNA.

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