

Analysis of Starch Grains Produced in Select Taxa Encountered in Southwest Asia

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Abstract: Starch grain analysis is a rapidly growing field of archaeological research in Southwest Asia. However, much work still remains regarding which taxa produce starch grains that can be identified in the archaeological record. In this paper, I centralize what is known about starch production patterns within regional flora and analyze 64 previously unstudied taxa from 22 families. The results of this study demonstrate that descriptions of starch grains from Southwest Asian taxa are scattered between archaeological and plant and food science publications. Ten of the species examined in this study, most of whom are grasses, produced starch grains that can be identified at varying taxonomic levels.

Keywords: Paleoethnobotany, Starch Grains, Southwest Asia

Introduction

Recent starch grain analysis in Southwest Asia has provided insight into new areas of research such as beer brewing in ancient Egypt (Samuel 1996) and the diets of middle Holocene farmers (Henry and Piperno 2008), Upper Paleolithic hunters and gatherers (Piperno et al. 2004), and Neanderthals (Henry et al. 2011). Despite these promising strides in archaeological research, much remains to be done in regards to discovering which plants produce starches in Southwest Asia and whether or not these starch grains can be used to aid archaeological and paleoecological endeavors. In this paper I seek to understand the research potential of archaeological starch grain research in Southwest Asia by: 1) centralizing where starch grain information about Southwest Asian taxa can be found; 2) examining 64 previously unstudied taxa from 22 families to assess their production patterns; and 3) examining the diagnostic potential of starches found in these new taxa if present.

Organization of Comparative Southwest Asian Publications

The most comprehensive and detailed information about Southwest Asian taxa are embedded within archaeological site reports from this region. These publications provide the best source of data because they cover almost all of the domesticated Southwest Asian taxa except for *Triticum durum* Desf. (durum wheat), *Triticum compactum* L. (club wheat), *Hordeum distichon* L. (two-rowed barley), and *Vicia ervilia* (L.) Willd. (bitter vetch) and numerous wild grasses (Henry et al. 2011; Henry and Piperno 2008; Piperno et al. 2004). Unfortunately, not all of these publications provide detailed descriptions of the taxa they discuss despite their pioneering endeavors. Having thorough descriptions of starch producing taxa included in publications provides information about which taxa do and do not produce starches thereby helping researchers in identifying their own archaeological starch materials.

Southwest Asian taxa are also described in archaeological publications from other parts of the world either because these taxa were introduced to the region or because their natural distribution overlaps with that of Southwest Asia. For example, Yang and Perry (2013) analyze starch grains from the tribe Triticeae that grow in north China. These taxa include introduced Southwest Asian domesticates, such as *Triticum aestivum* L. (bread wheat), and wild taxa that are native to both China and Southwest Asia, such as *Aegilops tauschii* Coss. (Tauschs goatgrass). A list of publications detailing starch grains from Poaceae taxa that grow in Southwest Asia can be found in Table 1.



Table 1. Poaceae (Gramineae) taxa that have been published. Note, not all of these publications include descriptions of optical properties.

Genus and Species	Source
Aegilops bicomis (Forsk.) Jaub. & Spach.	Henry et al., 2011
Aegilops caudate auct.	Reichert, 1913
Aegilops geniculata Roth	Piperno et al., 2004
Aegilops peregrina (Hackel) Maire et Weiler	Piperno et al., 2004
Aegilops speltoides Tausch	Henry et al., 2011
Aegilops truincialis L.	Reichert, 1913
Agropyron cristatum (L.) Gaertn.	Reichert, 1913
Agropyron rigidum (Schrad.) P. Beauv.	Reichert, 1913
Agrostis spica-venti L.	Reichert, 1913
Aira caespitosa L.	Reichert, 1913
Alopecurus arundinaceus Poir	Piperno et al., 2004
Alopecurus geniculatus L.	Reichert, 1913
Alopecurus utriculatus Banks & Sol.	Piperno et al., 2004; Reichert, 1913
Alopecurus pratensis L.	Reichert, 1913
Avena barbata Pott ex Link	Piperno et al., 2004
Avena sterilis L.	Henry et al., 2011
Brachypodium distachyon (L.) P.Beauv.	Piperno et al., 2004
Bromus brachystachys Hornung	Reichert, 1913
Bromus pseudobrachystachys H. Scholz	Piperno et al., 2004
Bromus squarrosus L.	Reichert, 1913
Gastridium ventricosum (G. australe) (Gouan) Schinz & Thell.	Piperno et al., 2004; Reichert, 1913
Hordeum bulbosum L.	Piperno et al., 2004
Hordeum glaucum Steudel	Henry et al., 2011; Piperno et al., 2004
Hordeum hexastichon L.	Henry et al., 2011
Hordeum marinum Huds.	Piperno et al., 2004
Hordeum sativum var. (Champion) Jess.	Reichert, 1913
Hordeum spontaneum L.	Henry et al., 2011; Piperno et al., 2004
Hordeum vulgare L.	Henry et al., 2011; Reichert, 1913
Koeleria macrantha (Ledeb.) Schult.	Messner, 2011
Lolium multiflorum Lam.	Piperno et al., 2004
Lolium rigidum Gaudin	Piperno et al., 2004
Lolium temulentum var. speciosum L.	Reichert, 1913
Phalaris minor Retz.	Piperno et al., 2004
Phalaris paradoxa L.	Piperno et al., 2004
Piptatherum holciforme (M.Bieb.) Roem. & Schult.	Piperno et al., 2004
Poa pratensis L.	Messner, 2011
Poa nemoralis L.	Messner, 2011
Puccinellia distans (Jacq.) Parl.	Piperno et al., 2004
Puccinellia gigantea (Grossh.) Grossh.	Piperno et al., 2004
Secale cereale L.	Reichert, 1913
Secale cereale var. MammothWinter L.	Reichert, 1913
Secale cereale var. Spring L.	Reichert, 1913
Secale cereale ssp. ancestrale L.	Henry et al., 2011
Secale vavilovii Grossh.	Henry et al., 2011

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Genus and Species	Source
Triticum aegilopoides (T. monococcum subsp aegilopoides) (Link) Balansa ex	Henry et al., 2011
Körn.	
Triticum aestivum (T. aestivum ssp aestivum) L.	Henry et al., 2011; 2009
Triticum dicoccum (T. turgidum ssp. dicoccum) Schrank ex Schübl	Reichert, 1913
Triticum dicoccoides Schrank ex Schübl	Piperno et al., 2004
Triticum monococcum L.	Reichert, 1913
Triticum monococcum subsp. aegilopoides	Henry et al., 2011
Triticum sativum var.dicoccum (Schrank)	Reichert, 1913
Triticum sativum var.vulgare	Reichert, 1913
Triticum turgidum Desf.	Henry et al., 2011; Reichert, 1913
Triticum urartu Tumanian ex Gandilyan	Henry et al., 2011
Vulpia persica (Boiss. & Buhse) Krecz. & Bobrov	Piperno et al., 2004

Bibliography for Poaceae of Southwest Asia:

Henry, A. G., A. S. Brooks, D. R. Piperno. 2011. Microfossils in Calculus Demonstrate Consumption of Plants and Cooked Foods in Neanderthal Diets (Shanidar III, Iraq; Spy I and II, Belgium). Proceedings of the National Academy of Sciences 108:486-491.

Henry, A. G., H. F. Hudson, and D. R. Piperno. 2009. Changes in Starch Grain Morphologies from Cooking. Journal of Archaeological Science 36:915–922.

Messner, T. C. 2011. Acorns and Bitter Roots: Starch Grain Research in the Prehistoric Eastern Woodlands. University of Alabama Press, Tuscaloosa, AL.

Piperno, D. R., E. Weiss, I. Holst, and D. Nadel. 2004. Processing of Wild Cereal Grains in the Upper Palaeolithic Revealed by Starch Grain Analysis. Nature 430:670-673.

Messner (2011) analyzes starch grains in seeds and underground storage organs (USOs) produced by taxa that grow in the Delaware River valley, USA. Two of these taxa, *Typha latifolia* L. (cattail) and *Cyperus esculentus* L. (yellow nutsedge), are also found in Southwest Asia (Davis 1965; Migahid 1988). Finally, a few Southwest Asian taxa are discussed in experimental archaeological publications where researchers examine how food-processing activities affect starch grain morphology and how these changes can be detected archaeologically (Ge et al. 2010; Henry et al. 2009).

Food and plant science research on Southwest Asian taxa is extensive, focusing on understanding the chemical and physical attributes of main Southwest Asian domesticates: *Triticum* spp. (wheat), *Hordeum* spp. (barley), *Secale* spp. (rye), *Vicia faba* L. (faba bean), *Lens culinaris* Medikus (lentil), *Pisum sativum* L. (pea), and *Cicer arietinum* L. (chickpea). Other domesticates, such as *Vicia sativa* (common vetch) and *Vicia ervilia* (bitter vetch), have received little attention.

Reichert (1913) provides the most comprehensive analysis of starch grains produced by taxa and remains

one of the seminal publications used by many paleoethnobotanists. In this publication, he reviews the state of starch grain research at the beginning of the 20th century, discusses the chemical and physical properties of specific taxa, and provides an assessment on how these taxa can be identified based on their chemical and physical characteristics. Many of the taxa that he describes are found in Southwest Asia and can be referenced by comparing the list of species he covers with the species listed in one of the regional floras such as the *Flora of Turkey and East Aegean Islands* (Davis 1965).

Materials and Methods

Selecting species for analysis

Sixty-four species representing 22 families that currently grow in Syria were collected from Professor Joy McCorriston's extensive Southwest Asian herbarium collection at Ohio State University. The 64 species were subdivided into their constituent parts resulting in eighty-two samples (Tables 2 and 3). These samples included seeds, pericarps, synconia, legumes, and legume capsules. In this study, the generic term "seed" is used for simplicity. No leaves, stems, or small roots

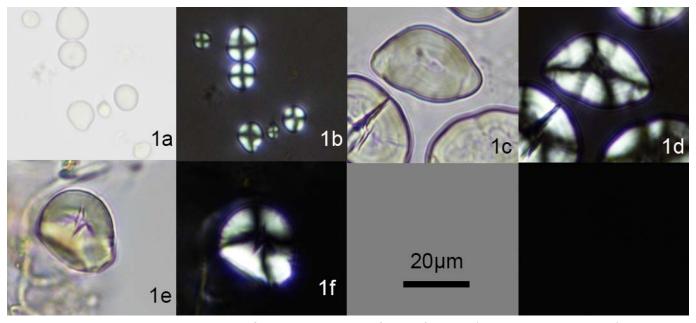


Figure 1. Transmitted and polarized views of starch at 400 × magnification from: a, b) *Cyperus esculentus*; c, d) *Vicia ervilia*; and e, f) *Moringa peregrina*.

were analyzed because they rarely produce large storage starch grains (Haslam 2004). Underground storage organs of important wetland taxa from the Cyperaceae family (Ryan 2011) were not available for analysis because they are difficult to store and are rarely found in herbarium collections.

Processing the samples

Samples were cleaned according to the protocol outlined by Pearsall (2000: 436–437), cut into small pieces using a sterile scalpel, or gently crushed using a sterile mortar and pestle. Very little pressure was applied when using the mortar and pestle to minimize potential damage to the starch grains. Two drops of a one to one glycerol/distilled water mix were placed on a 25 × 75 × 1mm microscope slide for each comparative sample. This medium was chosen, as opposed to a more permanent medium such as Permount or Entellen, in order to allow potential starch grains to be rotated when examined. The sample was gently covered with a microscope cover slip and the edges were sealed using finger nail polish and allowed to dry before being examined.

Recording methods

Samples were examined at 500× magnification using a Zeiss AxioStar Plus microscope. Each starch grain was given an identification number, described according to terms defined in the International Code

for Starch Grain Nomenclature (ICSN 2014) and measured using NIS Elements software. Photos of individual starch grains were taken at the Environmental Archaeology Lab at University of Texas. In order to minimize researcher bias, starch grains were chosen at random for description by using the random number generator function within Excel to provide x and y coordinates on the microscope stage. Fifty simple or half compound starch grains were described and photographed when present for each sample. Compound and aggregate starch grains were noted although excluded from the total count because clustering would often obscure their optical attributes making the individual starch grains difficult to describe and quantify. Starches less than five microns were typically omitted because their optical attributes were often hard to distinguish. Starch grains less than five microns in length were only counted in instances where they constituted the bulk of the starch grains produced.

Results

Ten of 64 species produced starch grains. All of the starches were produced in the seeds with the exception of *Moringa peregrina* (Forssk.) Fiori (Yusor tree) that concentrated its starch in the pericarp (Table 2). The 54 species that did not produce starch grains were from wild taxa that were related to the domesticated grains and legumes or from other types of domesticat-

Table 2. Taxa that produced starch grains in abundance in this study.

Family	Genus/Species	Plant Part
Cyperaceae	Cyperus esculentus L.	Seed
Fabaceae (Leguminosae)	Vicia ervilia	Legume
Moringaceae	Moringa peregrina	Pericarp
Poaceae (Gramineae)	Aegilops crassa Boiss	Seed
	Aegilops triaristata Willd.	Seed
	Aegilops vavilovii (Zhuk.) Chennav.	Seed
	Hordeum distichon L.	Seed
	Pennisetum americanum (L.) Leeke	Seed
	Triticum compactum Host.	Seed
	Triticum durum Desf.	Seed

ed taxa (Table 3). These taxa produced seeds that were very small and contained almost no starches.

Cyperus esculentus

The starches formed within *Cyperus esculentus* (yellow nutsedge) seeds have a mean length of four microns, range in size from one to eight microns and are mostly ovoid in shape (Figure 1a, b). They differ markedly in size and shape from the starches produced in the tuber or root-stock, which have an average length of 12 to 14mm and are conical to oval in shape (Reichert 1913). The seed starches are diagnostic to Cyperaceae because of their size and rounded, oval, compressed lenticular, angular, or polyhedral shapes that are commonly associated with other Cypereraceae seeds discussed in Reichert (1913).

Vicia ervilia

Vicia ervilia (bitter vetch) starches have an average length of 16mm and range in size from five to 27mm (Reichert 1913) (Figure 1c, d). Vicia ervilia starches from seeds can be identified to the family level because they exhibit what Reichert (1913) refers to as "bean type" features (spherical to ovoid in shape, half to as broad as long, slightly compressed with a distinct longitudinal cleft) that are characteristic of the Fabaceae (Leguminosae) family. These starches are mostly ovoid to elliptical and reniform shape and have deep longitudinal clefts.

Moringa peregrina

Moringa peregrina starches are mostly angular rounded, range in size from four to 27mm, and have an average length of ten microns (Figure 1. e, f). It is hard to determine if these starches are diagnostic because there are no close relatives discussed in Reichert (1913) or any of the other publications mentioned in this study. More studies should be conducted on

Moringa and closely related taxa to determine the diagnostic status of these starches. It is important to note that starch grains were extracted from the pericarp of the *M. peregrina* sample, and not the seed. This species suggests that tissues surrounding the seed, and not just the seed itself, need to be studied when conducting comparative starch grains research.

Aegilops crassa, A. triaristata, Hordeum distichon, Triticum durum, and T. compactum

The seeds from the species Aegilops crassa (Persian goatgrass) (Figure 2a, b), A. triaristata (three awn-goat

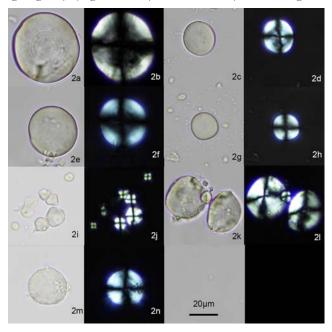


Figure 2. Transmitted and polarized views of starch at 400× magnification from: a, b) *Aegilops crassa*; c, d) *Aegilops triaristata*; e, f) *Aegilops vavilovii*; g, h) *Hordeum distichon*; i, j) *Pennisetum americanum*; k, l) *Triticum durum*; and m, n) *Triticum compactum*.



Table 3. Taxa that did not produce abundant starch grains.

Family	Genus/Species	Plant Part
Anacardiaceae	Pistacia atlantica Desf.	Seed
	Pistacia khinjuk Stocks	Seed
	Pistacia palaestina Boiss.	Seed
	Pistacia terebinthus L.	Seed
	Rhus coriaria L.	Seed
Apiaceae (Umbelliferae)	Bupleurum lancifolium Hornem.	Seeds
	Cuminum cyminum L.	Seeds
Arecaceae (Palmae)	Phoenix dactylifera L.	Seed
Asteraceae (Compositae)	Carthamus tinctorius L.	Seed
	Guizotia abyssinica (L.) Cass.	Seed
	Helianthus annus L.	Seed
	Notobasis syriaca (L.) Cass.	Seed
	Onopordum illyricum L.	Seed
	Onopordum palaestinum Eig.	Seed
	Silybum marianum (L.) Gaertn	Seed
Euphorbiaceae	Chorozophora tinctoria (L.) A. Juss.	Seed
Fabaceae (Leguminosae)	Acacia farnesiana (L.) Willd.	legume
	Acacia nilotica (L.) Delile	Seed
	Hymenocarpos circinnatus (L.) Savi	Legume
	Prosopis farcta Banks & Sol.) J. F. Macbr.	Legume capsule
	Trigonella foenum-graecum L.	Legume
	Trigonella monantha C. A. Mey.	Legume
	Trigonella stellata Forssk.	Legume
Geraniaceae	Erodium ciconium (L.) L'Hér. ex Aiton	Seed
	Erodium gruinum (L.) L'Hér. ex Aiton	Seed
Malvaceae	Malva parviflora L.	Seed
Moraceae	Ficus carica L.	Synconium, seed
Moringaceae	Moringa peregrina (Forssk.) Fiori	Seed
Oleaceae	Olea europaea L.	Pericarp, seed
Pedaliaceae	Sesamum indicum L.	Seed
Poaceae	Bromus scoparius Scop.	Seed
Polygonaceae	Polygonum patulum M. Bieb	Seed
	Polygonum venantianum Clementi	Seed
Ranunculaceae	Adonis dentata Delile	Seed
Rhamnaceae	Rhamnus palaestinus Boiss.	Pericarp, seed
	Zizyphus spina-christi (L.) Desf.	Exocarp, pericarp, seed
Rosaceae	Amygdalus arabica (Oliv.)	Pericarp, seed
	Amygdalus communis L.	Pericarp, seed
	Amygdalus orientalis Mill.	Exocarp, pericarp, seed

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Family	Genus/Species	Plant Part
	Crataegus aronia (L.) DC	Pericarp, seeds
	Prunus domestica L.	Seeds
	Prunus mahaleb L.	Seeds
	Prunus persica (L.) Stokes	Pericarp, seed
	Rosa canina L.	Pericarp/seed, seeds
	Rosa phoenicea Boiss.	Pericarp, seeds
	Sarcopoterium sinposum (L.) Spach.	Seeds
Rubiaceae	Asperula arvensis L.	Seeds
	Coffea arabica L.	Beans
	Galium tricornutum Dandy	Seeds
Solanaceae	Hyscamus muticus L.	Seed
	Physalis alkekengi L.	Seed
	Physalis angulata L.	Pericarp
	Solanum sepicula Dunal	Seed, fruit
Urticaceae	Urtica pilulifera L.	Seed
Zygophllaceae	Balanites aegyptiaca (L.) Delile	Exocarp, pericarp, seed

grass) (Figure 2c, d), Hordeum distiction (two-rowed barley) (Figure 2g, h), Triticum durum (durum wheat) (Figure 2k, l), and T. compactum (club-wheat) (Figure 2m, n) in this study all exhibit features that are diagnostic of the tribe Triticeae within the Poaceae (Gramineae) family. In general, starch grains from this tribe have simple, lenticular, oval, kidney (reniform) or dicoid in shapes with small reticulate surface depressions (Piperno et al. 2004; Yang and Perry 2013). The five Triticeae taxa that yielded abundant starch grains within this study exhibited all of these features

Seed starch grains from Aegilops, Hordeum, and Triticum taxa (AHT) and the Triticeae tribe are also much larger in general than the seed starch grains from non-Triticeae taxa. This feature can be used to identify individual starches at least to the tribe level when shape and size attributes are analyzed together. The mean length for the Poaceae starch grains observed in this study follow the pattern observed by Piperno et al. (2004) where AHT taxa can be distinguished from other grass taxa, such as the Pennisetum americanum, based on their overall large size (Table 4). The average length of the 18 AHT seed starch grain taxa in Table 4 with a sample size of 50 is 17.7mm with a standard deviation of 5.7mm. This length is well above the average length of the 15 non-Triticeae with an average of 5.1mm and a standard deviation 2.6mm. Recent work by Yang and Perry (2013) on 38 grass species from China supports this hypothesis and goes one step further, suggesting that all members of the tribe Triticeae produce larger starches relative to other Poaceae.

The one non-Triticeae grass in this study that yielded abundant seed starch, *Pennesitum americanum* yielded semi-compound to compound, flat, angular, or irregular shaped starch grains (Figure 2. i, j). This compares well with other studies of non-Triticeae grasses such as *Bromus* sp. and *Pipatherum* sp. where similar features were observed (Piperno et al. 2004).

Discussion and Conclusions

Chemical and physical properties of starch grains from over 100 species from Southwest Asia have been published in archaeological reports and food and plant science literature. An additional 64 species were examined here, ten of which produced abundant starch grains in their seeds and pericarps that are diagnostic at the tribe, family, and potentially genus and species level. This project adds to the growing body of knowledge regarding archaeological starch grain analysis in Southwest Asia by centralizing the published comparative literature for this region and describing the starches produced in domesticated and wild taxa.

The starches from Cyperus esculentus seeds are



Table 4. Mean (±s.d.) length (mm) and range for Poaceae starch grains divided by subfamily and tribe.

Subfamily	Tribe	Genus/species	Mean	Range	n	Source
Panicodae	Paniceae	Pennisetum americanum (L.) Leeke	5.7 (1.4)	3-10	50	This study
Pooideae	Aveneae	Alopecurus arundinaceus Poir.	4 (0.9)	2–8	50	Piperno et al 2004
		Alopecurus utriculatus Banks & Sol.	5 (1.5)	2–8	50	Piperno et al 2004
		Avena barbata Pott ex Link	12 (2.9)	6–18	50	Piperno et al 2004
		Gastridium ventricosum (Gouan) Schinz & Thell.	4 (1.0)	2–6	50	Piperno et al 2004
		Phalaris minor Retz.	<2.0	-	50	Piperno et al 2004
		Phalaris paradoxa L.	<4.0	-	50	Piperno et al 2004
	Brachypodieae	Brachypodium distachyon (L.) P.Beauv.	9 (2.2)	4–16	50	Piperno et al 2004
	Bromeae	Bromus pseudobrachystachys H. Scholz	5 (1.4)	4–8	50	Piperno et al 2004
	Poeae	Lolium multiflorum Lam.	<6.0	-	50	Piperno et al 2004
		Lolium rigidum Gaudin	<6.0	-	50	Piperno et al 2004
		Puccinellia distans (Jacq.) Parl.	<4.0	-	50	Piperno et al 2004
		Puccinellia gigantea (Grossh.) Grossh.	<4.0	-	50	Piperno et al 2004
		Vulpia persica (Boiss. & Buhse) Krecz. & Bobrov	<2.0	-	50	Piperno et al 2004
	Stipeae	Piptatherum holciforme (M.Bieb.) Roem. & Schult.	3 (1.0)	2–4	50	Piperno et al 2004
	Triticeae	Aegilops crassa Boiss	16 (7.6)	5–31	50	This study
		Aegilops geniculata Roth	21 (6.4)	10–36	50	Piperno et al 2004
		Aegilops peregrina Hack.	25 (8.0)	12–52	50	Piperno et al 2004
		Aegilops speltoides Tausch	22 (4.5)	10-32	50	Henry et al 2011
		Aegilops triaristata Willd.	10 (3.4)	5–20	50	This study
		Aegilops vavilovii (Zhuk.) Chennav.	13 (6.2)	5–35	50	This study
		Hordeum bulbosum L.	17 (3.7)	10–24	50	Piperno et al 2004
		Hordeum bulbosum (with lamellae only)	21 (1.6)	18–24	50	Piperno et al 2004
		Hordeum distichon L.	11 (2.7)	5–18	50	This study
		Hordeum glaucum Steudel	18 (3.5)	10-30	39	Henry et al 2011
		Hordeum glaucum Steudel	18 (3.9)	8–24	50	Piperno et al 2004
		Hordeum glaucum (with lamellae only)	22 (1.4)	18–26	50	Piperno et al 2004



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Subfamily	Tribe	Genus/species	Mean	Range	n	Source
Pooideae	Triticeae	Hordeum hexastichon L.	20 (3.5)	10-30	52	Henry et al 2011
		Hordeum marinum Huds.	10 (1.8)	6–14	50	Piperno et al 2004
		Hordeum spontaneum L.	18 (3.8)	12-30	27	Henry et al 2011
		Hordeum spontaneum L.	20 (4.7)	10–26	50	Piperno et al 2004
		Hordeum spontaneum (with lamellae only)	28 (2.9)	18–26	50	Piperno et al 2004
		Secale vavilovii Grossh.	25 (4.2)	15-36	50	Henry et al 2011
		Triticum aestivum L.	24 (4.4)	15-35	52	Henry et al 2011
		Triticum compactum Host.	12 (4.9)	5–22	50	This study
		Triticum dicoccoides Schrank ex Schübl.	17 (6.1)	8–30	50	Piperno et al 2004
		Triticum durum Desf.	11 (4.0)	5-23	50	This study
		Triticum monococcum subsp. aegilo- poides (Link.) Thell.	15 (1.7)	10–20	46	Henry et al 2011

distinct from the starches produced in its tubers but are similar to the seeds of other related taxa within the Cyperaceae family making them diagnostic to this family. Vicia ervilia starches exhibit "bean type" features and can be identified to the genus and species level due to their small size and Fabaceae (Leguminosae)-like properties. Although the diagnostic ability of Moringa peregrina starches remains unclear, their production in the pericarp, and not the seed challenges assumptions originally made in this project, and in the general literature, about perceived starch production in particular plant parts and illustrates the importance of testing every part of a plant when possible. Finally, the Poaceae taxa in this study can be distinguished from each other at the tribe level by size and overall shape.

Centralization of information about taxa that produce starch grains will help specialists narrow down identification of unknown starch grains encountered in the archaeological record. The discovery of starch grains within important domesticated taxa such as *Hordeum distichon*, *Triticum durum*, and wild taxa such as *Cyperus esculentus* provides a clearer understanding of what can be identified within Southwest Asia and within these families and genera.

There are many avenues of comparative starch grain research that can be pursued to better aid archaeologists in their reconstruction of plant use in Southwest Asia. With a few exceptions, very little research has been conducted on starch grains pro-

duced by underground storage organs such as bulbs, corms, rhizomes, and tubers (Henry et al. 2009, 2011; Messner 2011; Piperno et al. 2004; Reichert 1913; Yang and Perry 2013). Macrobotanical and phytolith evidence suggests that wetland taxa played an important role as a source of food in Southwest Asia during the Epipaleolithic (Wollstonecroft et al. 2008), Pre-Pottery Neolithic (Balbo et al. 2012), Pottery Neolithic (Rosen 2005), and Ubaid (Kennett and Kennett 2006) periods. Aside from the research by Hather (1991, 1993), very little work has been conducted to establishd criteria for identifying underground storage organs at archaeological sites. Recovering and identifying starch grains associated with USO's would open a whole new avenue of research into wild resource exploitation, complement existing datasets, and allow for archaeologists to explore new topics through the analysis of starches contained in artifact residues and dental calculus.

The research on Triticeae taxa from China (Yang and Perry, 2013) and taxa from the Delaware River Valley, USA (Messner 2008, 2011) are excellent examples of how a regional synthesis can lead to the construction of standardized dichotomous keys for a region. In both of these papers, the researchers develop an easy to use dichotomous key that allows for quick identification of archaeological starch grains. Further research into starch grain production patterns of other taxa found in Southwest Asia and the identification of Southwest Asian taxa discussed in Reichert (1913) would eventually lead to the develop-



ment of a dichotomous key and the establishment of regional diagnostic starch grain types that archaeologists could use in this important area of the world.

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Biosketch

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