Genome-wide sequence-based genotyping supports a nonhybrid origin of *Castanea alabamensis*

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Short title: Nonhybrid origin of *Castanea alabamensis*

Abstract: The genus *Castanea* in North America contains multiple tree and shrub taxa of conservation concern. The two species within the group, American chestnut (*Castanea dentata*) and chinquapin (*C. pumila* sensu lato), display remarkable morphological diversity across their distributions in the eastern United States and southern Ontario. Previous investigators have hypothesized that hybridization between *C. dentata* and *C. pumila* has played an important role in generating morphological variation in wild populations. A putative hybrid taxon, *Castanea alabamensis*, was identified in northern Alabama in the early 20th century; however, the question of its hybridity has been unresolved. We tested the hypothesized hybrid origin of *C. alabamensis* using genome-wide sequence-based genotyping of *C. alabamensis*, all currently recognized North American *Castanea* taxa, and two Asian *Castanea* species at >100,000 single-nucleotide polymorphism (SNP) loci. With these data, we generated a high-resolution phylogeny, tested for admixture among taxa, and analyzed population genetic structure of the study taxa. Bayesian clustering and principal components analysis provided no evidence of

admixture between *C. dentata* and *C. pumila* in *C. alabamensis* genomes. Phylogenetic analysis of genome-wide SNP data indicated that *C. alabamensis* forms a distinct group within *C. pumila* sensu lato. Our results are consistent with the model of a nonhybrid origin for *C. alabamensis*. Our finding of *C. alabamensis* as a genetically and morphologically distinct group within the North American chinquapin complex provides further impetus for the study and conservation of the North American *Castanea* species.

Key words: American chestnut; *Castanea dentata*; *Castanea pumila*; chinquapin; genotyping-by-sequencing; hybridization; Fagaceae

Hybridization is an important and widespread phenomenon in plants (Ellstrand et al., 1996; Mallet, 2005; Soltis and Soltis, 2009; Whitney et al., 2010). Evolutionary outcomes of hybridization can include reinforcement or breakdown of reproductive barriers, increased intraspecific diversity, transfer of genetic adaptations between species, and the origin of new ecotypes or species (Rieseberg, 1997). Many authors have asserted that the descendants of natural interspecific hybridization warrant conservation because they represent a natural part of the evolutionary legacy of taxa (Whitham et al., 1991; Allendorf et al., 2001; Allendorf et al., 2013; Stronen and Paquet, 2013; Jackiw et al., 2015). Thus, the identification and characterization of natural hybrids, particularly in groups of conservation concern, is a priority for understanding and protecting plant biodiversity. The genus *Castanea* Mill. (Fagaceae) in North America exemplifies this concept.

There exists remarkable morphological diversity within the North American *Castanea*, a group that is thought to be comprised of just two species, American chestnut (*Castanea dentata* (Marsh.) Borkh.) and North American chinquapin (*C. pumila* (L.) Mill.) (Johnson, 1988). Within *C. pumila* alone, plant habit at maturity varies from rhizomatous subshrub in some populations of the Gulf Coastal Plain to canopy tree in other populations (Johnson, 1988; Nixon, 1997). Such variation in plant habit, as well as variation in flower, fruit, leaf, and twig morphology has led several authors to speculate that hybridization between *C. dentata* and *C. pumila* has contributed significantly to the variation observed in wild North American *Castanea* populations (Dode, 1908; Camus, 1929; Elias, 1971; Little, 1979; Nixon, 1997; Kubisiak and Roberds, 2006; Binkley, 2008; Shaw et al., 2012; Li and Dane, 2013). Reports of naturally occurring hybrids between *C. dentata* and *C. pumila* var. *pumila* have resulted in the description of two hybrid taxa. Dode (1908) published the first description of putative hybrid progeny of *C. dentata* and *C.*

pumila, referring to these plants from North Carolina, Virginia, and Maryland as *C. × neglecta* Dode. The second hybrid taxon resulted from Camus' (1929) revised treatment of the nonhybrid species *C. alabamensis* Ashe, a tree endemic to the Appalachian Mountains of northern Alabama. The putative hybrid taxa, *C. alabamensis* and *C. × neglecta*, both possess a combination of traits typically used to differentiate *C. dentata* from *C. pumila*. In *C. alabamensis*, specifically, plants produce one pistillate flower per cupule and one nut per bur, like *C. pumila*, yet mature plants grow as medium to large trees, have glabrous twigs, and have apparently glabrous abaxial leaf surfaces when viewed without magnification, like *C. dentata* with which they are sympatric (Ashe, 1925).

Multiple genetic studies over the past decade have investigated the putative hybridity of several North American *Castanea* populations (Binkley, 2008; Dane, 2009; Shaw et al., 2012; Li and Dane, 2013). Binkley (2008) and Shaw et al. (2012) used a small number of noncoding cpDNA regions to investigate populations of suspected hybrids in northwestern Georgia, which they described as having an “intermediate morphology” that corresponded to the hybrid taxon *C. × neglecta*. Their *C. × neglecta* accessions had a unique cpDNA haplotype that was not found in a diverse panel of the putative parental species, *C. dentata* and *C. pumila*, contrary what they predicted would have been observed in hybrid individuals. Authors of both studies concluded that further investigation would be needed to rigorously test the hybridity of their study populations in northwestern Georgia (Binkley, 2008; Shaw et al., 2012). The most recent study of reputed naturally-occurring hybrids between *C. dentata* and *C. pumila* used six noncoding cpDNA regions and 680 bp from two nuclear DNA regions to investigate the extent of hybridization between these two species (Li and Dane, 2013). An important finding of Li and Dane (2013) related to their discovery of two different morphological groups at a site in northern

Alabama, which they designated as distinct morphological types of *C. dentata*: Type I *C. dentata* (possessing morphological features typical of *C. dentata* throughout its range) and Type II *C. dentata* (possessing several leaf trichome features that the authors described as indicative of morphological intermediacy between *C. dentata* and *C. pumila*). In phylogenetic analyses of cpDNA and nuclear DNA sequences, Type I *C. dentata* grouped with other *C. dentata* plants sampled throughout the species' range, while Type II *C. dentata* grouped with *C. pumila* in analyses of both chloroplast and nuclear DNA datasets. Li and Dane (2013) concluded that the genetic and morphological patterns observed in Type II *C. dentata* might be the result of past hybridization between *C. dentata* and *C. pumila*.

A limitation of previous genetic studies of putative hybrids between *C. dentata* and *C. pumila* is that nearly all of the analyses relied principally on datasets comprised of information from a small number of maternally-inherited cpDNA loci. In the few cases where loci from the nuclear genome were analyzed, these data were not tested for signatures of hybridization. Although, Li and Dane (2013) used both cpDNA and nuclear DNA, results of their analysis of 680 bp from two nuclear loci showed no genetic evidence of *C. dentata* ancestry in their Type II *C. dentata* individuals. Their dataset also lacked the information necessary to more finely discern relationships between different putatively hybridized Type II *C. dentata* individuals and the *C. pumila* individuals found on the same branch of the phylogenetic tree. Moreover, analytical methods used to make inferences about past hybridization in this group have been limited to phylogenetic trees and cpDNA haplotype networks, using models that assume a strictly bifurcating pattern of lineage evolution (Binkley, 2008; Dane, 2009; Shaw et al., 2012; Li and Dane, 2013). Therefore, despite much speculation, convincing evidence regarding the occurrence

or extent of natural hybridization between the North American *Castanea* taxa has not been produced.

The North American *Castanea* are currently thought to consist of either two or three species. The most recent formal taxonomic revision of the group relied on morphology and concluded that North American *Castanea* is comprised of two species, American chestnut (*C. dentata*) and North American chinquapin (*Castanea pumila*)—the latter species containing the botanical varieties Allegheny chinquapin (*C. pumila* (L.) Mill. var. *pumila*) and Ozark chinquapin (*C. pumila* var. *ozarkensis* (Ashe) Tucker) (Johnson, 1988). Other taxonomists contend that the ecological and morphological differences between Allegheny chinquapin and Ozark chinquapin are substantial enough to justify recognition at the level of species, and they recognize three species of North American *Castanea*: *C. dentata*, *C. pumila*, and *C. ozarkensis* (Nixon, 1997; Weakley, 2015). American chestnut (*C. dentata*) produces three pistils per cupule, three nuts per bur, and apparently glabrous abaxial leaf surfaces that rarely contain stellate trichomes, and healthy individuals grow as trees at maturity (Nixon, 1997). Ozark chinquapin and Allegheny chinquapin typically produce one pistil per cupule, one nut per bur, and have sun leaves with abaxial surfaces that are covered in stellate trichomes (Johnson, 1988; Nixon, 1997). However, Ozark chinquapin grows as a tree at maturity and Allegheny chinquapin varies from rhizomatous shrub, to non-rhizomatous shrub, to small tree at maturity (Johnson, 1988). Dode (1908) describes the hybrid taxon *C. × neglecta* as having slightly pubescent twigs and leaves, reminiscent of the putative parent *C. pumila*, and larger burs and arborescent habit, like the putative parent *C. dentata*. Ashe (1925) originally described *C. alabamensis* as a distinct chinquapin species, but it was treated as a hybrid taxon throughout most of the 20th century (Camus, 1929; Elias, 1971; Little, 1979). In the most recent taxonomic revision of the group,

Johnson (1988) found no living plants that corresponded to the description *C. × neglecta*. Johnson (1988) treated *C. alabamensis* as a synonym of *C. pumila* var. *ozarkensis* on the basis of morphological similarities between the two taxa and, after field work near Ashe's *C. alabamensis* type locality in northern Alabama, he concluded that *C. pumila* var. *ozarkensis* had been extirpated from east of the Mississippi River by chestnut blight.

The advent of affordable genome-wide sequence-based genotyping methods has recently allowed new insights into questions of hybridization and admixture in plants (Escudero et al., 2014; Eaton et al., 2015; Baute et al., 2016; Owens et al., 2016; Leroy et al., 2017; McVay et al., 2017; Zhao et al., 2018; Turner et al., 2018). The availability of large numbers of markers distributed across the genome has allowed researchers to address questions that were previously intractable with less comprehensive datasets. Notably, these methods have allowed researchers to understand hybridization, introgression, and species boundaries on a fine scale in *Quercus*, a genus closely related to *Castanea* that is well known for a history of interspecific hybridization (Eaton et al., 2015; Leroy et al., 2017; McVay et al., 2017). Thus, genome-wide sequence-based genotyping represents a potentially useful approach for understanding hybridization and admixture in the North American *Castanea* species.

Here, we investigate the origins of purported hybrid populations of *Castanea* in the eastern United States. Our main objective was to test the hypothesized hybrid origin of samples corresponding to the morphological description of *C. alabamensis*. A secondary objective was to test for admixture in sympatric populations of *C. dentata* and *C. pumila*. We used sequence-based genotyping to gather genome-wide single nucleotide polymorphism (SNP) data from samples representing *C. alabamensis*, all currently recognized North American *Castanea* taxa (*C. dentata*, *C. pumila* var. *pumila*, *C. pumila* var. *ozarkensis*) and two East Asian outgroup

species (*C. crenata* and *C. mollissima*). Additionally, we examined the morphology of 865 herbarium accessions representing the type specimens of *C. alabamensis* and all extant *Castanea* species to place our samples in taxonomic context with recent studies of hybridization in the North American *Castanea* (Binkley, 2008; Shaw et al., 2012; Li and Dane, 2013) and earlier hypotheses based on morphology (Ashe, 1925; Camus, 1929; Elias, 1971; Little, 1979; Johnson, 1988).

MATERIALS AND METHODS

Plant materials, DNA extraction, and morphological comparisons—Young leaf tissues for DNA extraction were collected from 106 naturally-occurring North American *Castanea* and cultivated East Asian *Castanea* plants. We sampled plants corresponding to Ashe’s (1925) description of *C. alabamensis*, the putative parent taxa, *C. dentata* and *C. pumila* var. *pumila*, the remaining North American taxon, *C. pumila* var. *ozarkensis*, and two East Asian outgroup species, *C. mollissima* Blume (Chinese chestnut) and *C. crenata* Siebold & Zucc. (Japanese chestnut) (Table 1; Appendix S1). Samples representing two of the *C. × neglecta* populations studied by Binkley (2008) and Shaw et al. (2012) were included in sequencing (Table 1; Appendix S1). Total genomic DNA was extracted from 100 mg of fresh or frozen leaf tissue using the Qiagen DNeasy Plant Mini Kit (Qiagen, Valencia, California, USA) or a modified version of the CTAB protocol of Kubisiak et al. (2013). DNA quality and integrity were examined using a NanoDrop ND-8000 spectrophotometer (Thermo Fisher Scientific Inc., Waltham, Massachusetts, USA) and electrophoresis on 1% agarose gels. Genomic DNA was quantified using the Qubit 2.0 fluorometric quantification assay (Life Technologies, Carlsbad, California, USA).

Corresponding herbarium vouchers were obtained for wild-collected North American *Castanea* samples and assessed for a suite of morphological features that differentiate *C. dentata*, *C. pumila* var. *pumila*, *C. pumila* var. *ozarkensis*, and *C. alabamensis* (Table 1). Herbarium vouchers were also collected from the northern Alabama population containing the suspected hybrid “Type II *C. dentata*” studied by Li and Dane (2013) (Appendix S2). Finally, morphological comparisons were made between the genotyped plants and the entire *Castanea* collection at the University of North Carolina Chapel Hill Herbarium (NCU), which contains a total of 865 specimens representing all extant *Castanea* species and the type specimens of *C. alabamensis* (Appendix S3). Morphological features assessed included four leaf characters (sun leaf abaxial surface covered with stellate trichomes or not; simple trichomes present or absent on abaxial lamina; abaxial surface glaucous or not; leaf margin ciliate or eciliate), one twig character (twigs pubescent or not), and one flower/fruit character (pistil per cupule ratio or nut per bur ratio).

190 Table 1. Morphological and locality information for *Castanea* populations genotyped in this study. Data for numbers of pistils per
 191 cupule and nuts per bur reflect observations from spring and fall, respectively.

Taxon/County, State	Site name	No. of plants	GBS IDs	Morphological features					
				Abaxial with abundant stellate trichomes	Simple trichomes on laminar surfaces	Glaucous abaxial surface	Ciliate leaf margin	Pubescent twigs	Pistils per cupule/ nuts per bur
<i>C. pumila</i> var. <i>pumila</i>				Yes	Yes	No	Yes	Yes	1 (-3)
Oconee, SC	BF	4	29, 33, 37, 42	Yes	Yes	No	Yes	Yes	NA
Oconee, SC	CR	5	15, 19, 21, 25, 30	Yes	Yes	No	Yes	Yes	NA
Cherokee, NC	SE	8	53, 54, 55, 57, 58, 60, 61, 62	Yes	Yes	No	Yes	Yes	1
Charleston, SC	AW	9	1, 5, 9, 13, 17, 23, 28, 32, 36	Yes	Yes	No	Yes	Yes	NA
Sumter, SC	WB	2	41, 44	Yes	Yes	No	Yes	Yes	NA
Suwannee, FL	SS	3	80, 81, 82	Yes	Yes	No	No	No	1,2,3
Transylvania, NC	GO	2	20, 24	Yes	Yes	No	Yes	Yes	NA
Baker, GA	JR	1	26	Yes	Yes	No	Yes	Yes	1
Pickens, GA	TM	6	35, 40, 43, 46, 47, 48	Yes	Yes	No	Yes	Yes	1
<i>C. pumila</i> var. <i>ozarkensis</i>				Yes	Yes	No	Yes	No	1
Lawrence, MO	MO	1	63	Yes	Yes	No	Yes	No	1
Sharp, AR	SH	1	64	Yes	Yes	No	Yes	No	1
<i>C. alabamensis</i>				No	Yes	Yes	Yes	No	1
Cleburne, AL	HU	3	67, 86, 89	No	Yes	Yes	Yes	No	1
Clay, AL	AG	5	73, 75, 76, 85, 87	No	Yes	Yes	Yes	No	1
Calhoun, AL	CH	6	90, 92, 93, 94, 95, 96	No	Yes	Yes	Yes	No	1
Floyd, GA	JM [†]	1	22	No	Yes	Yes	Yes	No	1
Floyd, GA	PO [†]	1	38	No	Yes	Yes	Yes	No	1
<i>C. dentata</i>				No	No	No	No	No	3
Cleburne, AL	HU	3	68, 69, 70	No	No	No	No	No	NA
Clay, AL	AG	2	74, 77	No	No	No	No	No	NA
Calhoun, AL	CH	1	91	No	No	No	No	No	NA
Cleburne, AL	FR	3	71, 72, 84	No	No	No	No	No	NA
Morgan, AL	LA	3	78, 79, 88	No	No	No	No	No	NA
Pickens, GA	TM	2	27, 31	No	No	No	No	No	NA
Lincoln, TN	LI	1	83	No	No	No	No	No	3
Oconee, SC	BF	2	3, 45	No	No	No	No	No	NA
Oconee, SC	CR	2	7, 11	No	No	No	No	No	NA
Pickens, SC	CE	4	2, 6, 10, 14	No	No	No	No	No	NA
Transylvania, NC	GO	1	18	No	No	No	No	No	NA

Cherokee, NC	JB	3	50, 51, 52	No	No	No	No	No	NA
Cherokee, NC	SE	1	59	No	No	No	No	No	NA
Buncombe, NC	DO	1	102	No	No	No	No	No	3
Avery, NC	CB	1	103	No	No	No	No	No	3
Washington, VA	WN	1	97	No	No	No	No	No	3
Adair, KY	KY	1	16	NA	NA	NA	NA	NA	NA
Westmoreland, PA	WE	1	12	No	No	No	No	No	NA
Warren, PA	WR	1	8	No	No	No	No	No	NA
Huntingdon, PA	RO	1	4	No	No	No	No	No	NA
Knox, ME	KN	1	34	No	No	No	No	No	NA
Piscataquis, ME	AT	1	39	No	No	No	No	No	NA
Piscataquis, ME	SB	1	66	No	No	No	No	No	NA
Waldo, ME	WA	1	65	No	No	No	No	No	NA
<i>C. mollissima</i>				NA	NA	NA	NA	NA	3
‘Nanking’	NK	1	100	NA	NA	NA	NA	NA	3
‘Mahogany’	MY	1	98	NA	NA	NA	NA	NA	3
<i>C. crenata</i>				NA	NA	NA	NA	NA	3
‘Fort Defiance’	FD	1	104	NA	NA	NA	NA	NA	3
‘Morrow Mountain’	MM	1	107	NA	NA	NA	NA	NA	3
‘Rita 1’	RI	1	108	NA	NA	NA	NA	NA	NA
‘Rita 2’	RA	1	110	NA	NA	NA	NA	NA	NA

Note: NA = character was not observed in the individual or species; † = samples 22 and 38, from the JM and PO sites, were previously identified as *C. × neglecta* and sequenced in the studies of Binkley (2008) and Shaw et al. (2012).

GBS library construction, data processing, and SNP discovery—Genotyping-by-sequencing (GBS) libraries were prepared using a modified version of the methods of Elshire et al. (2011), as described by Zhebentyayeva et al. (2019). We followed all aspects of the GBS library preparation protocol of Zhebentyayeva et al. (2019), with one exception—we double-digested DNA samples with *Pst*I and *Mse*I. Briefly, we performed double digestion of DNA samples, ligated fragments to Illumina sequencing adapters and custom barcoded adapters (see Appendix S4 for barcode sequences), pooled samples, then purified the pooled samples using a QIAquick PCR purification kit (Qiagen, Valencia, California, USA). Pooled samples were amplified using 18 cycles of PCR and size-selection was performed using 0.4× and 0.8× volumes of Mag-Bind Total Pure NGS magnetic beads (Omega Bio-Tek, Georgia, USA) to remove fragments larger than 1.5 kb and smaller than 121 bp, respectively. The quality of GBS libraries was validated using a 2100 Bioanalyzer (Agilent Genomics, Santa Clara, California, USA). 96-plex GBS libraries were paired-end sequenced (2 × 125 bp) on an Illumina HiSeq 2500 (Illumina Inc., San Diego, California, USA) at the Hollings Cancer Center at the Medical University of South Carolina.

Data processing of Illumina reads was performed using Clemson University's Palmetto Cluster high-performance computing resource as described by Zhebentyayeva et al. (2019). Default settings of the Stacks 1.45 program 'process_radtags' (Catchen et al., 2011; Catchen et al., 2013; Rochette and Catchen, 2017) were used to demultiplex raw Illumina reads according to barcodes, discard reads with uncalled bases, discard reads with low quality scores, and filter reads for the presence of *Pst*I and *Mse*I restriction sites. Demultiplexed reads were uploaded to the National Center for Biotechnology Information (NCBI) Sequence Read Archive (SRA) under

222 BioProject ID: PRJNA541592. Individual plants with less than 100,000 total reads were removed
 223 from further processing. Samples from the *C. mollissima* and *C. crenata* outgroup cultivars were
 224 replicated twice in GBS libraries, and technical replicates with the highest number of retained
 225 reads were used for further processing. Demultiplexed reads were aligned to the *C. mollissima*
 226 reference genome v.1.1 (<https://www.hardwoodgenomics.org>) using the GSNAP software
 227 package (Wu and Nacu, 2010). Single nucleotide polymorphisms were called against the *C.*
 228 *mollissima* reference genome sequence and a catalog of tags and SNPs was generated using the
 229 ‘ref_map.pl’ command in Stacks, with default settings. SNP genotypes were generated in the
 230 ‘populations’ program in Stacks and filtered to remove indels and multi-nucleotide variants,
 231 leaving only bi-allelic SNPs and invariant sites. To avoid issues resulting from analysis of tightly
 232 linked markers, we used the *whitelist* feature in the ‘populations’ program to retain only one SNP
 233 per locus and create a list of 500,000 randomly selected SNPs for further processing. Individual
 234 genotypes were filtered to retain only SNPs supported by five or more reads and SNPs with a
 235 minimum allele frequency of 0.01. The SNPs were further filtered to produce three different
 236 datasets for analysis, two of which were exported in Variant Call Format (VCF) and the third as
 237 a .txt file: (1) a dataset comprised of 103,616 SNPs and 103 individuals representing North
 238 American and east Asian *Castanea* species was produced by filtering for SNPs present in >80%
 239 of individuals (dataset ‘NAC+EAC’); a dataset comprised of 190,656 SNPs and 96 individuals
 240 representing the North American *Castanea* taxa was produced by filtering for SNPs present in
 241 >75% of individuals (dataset ‘NAC’); and a smaller dataset comprised of 583 SNPs and 103
 242 individuals representing North American and east Asian *Castanea* species for STRUCTURE
 243 analysis was produced by filtering for SNPs present in >95% of individuals (dataset
 244 ‘NAC+EAC583’). All datasets exported by Stacks were deposited on GitHub:

https://github.com/MTPerkins/Nonhybrid_origin_of_Castanea_alabamensis (see Appendix S5 for correspondence of dataset names to files on GitHub).

Phylogeny and genetic differentiation—We inferred a maximum likelihood phylogenetic tree for the North American and East Asian *Castanea* taxa (dataset ‘NAC+EAC’) using the RAxML 8.2.11 (Stamatakis 2014) plugin for Geneious 11.1.5 (<http://www.geneious.com/>). We used the GTRCAT model of nucleotide evolution with 100 rapid bootstrap replicates and a subsequent search for the best scoring maximum likelihood tree.

To compare levels of genetic differentiation between the groups identified in the phylogenetic tree, we estimated Weir and Cockerham’s F_{ST} (1984) between pairs of taxa in the ‘NAC+EAC’ dataset using the ‘SNPRelate’ package (Zheng et al. 2012) in R (R Core Team 2018), with default settings.

STRUCTURE analyses—To determine the number of genetically differentiated clusters (K) and to detect admixture among all samples (i.e., North American and East Asian species), we performed Bayesian clustering analysis on the ‘NAC+EAC583’ dataset with STRUCTURE 2.3.4 software (Pritchard et al. 2000; Falush et al. 2003). We ran STRUCTURE using 35,000 burn-in repetitions, 35,000 Markov Chain Monte Carlo repetitions after burn-in, the correlated allele frequencies setting, and the admixture model for 10 iterations at $K = 1-9$ (i.e., the number of putative taxa, plus three). After finding that *C. alabamensis* grouped with *C. pumila* sensu lato in phylogenetic analysis and the first STRUCTURE analysis, we determined the number of genetic clusters and levels of admixture in this group by removing *C. dentata*, *C. crenata*, and *C. mollissima* from the ‘NAC+EAC583’ dataset and running STRUCTURE with the same settings as above, apart from using $K = 1-16$ (i.e., the number of chinquapin sample sites, plus one). The number of clusters within our *Castanea* samples from North America and eastern Asia and

within the chinquapins were determined by the ΔK method (Evanno et al. 2005) implemented within the STRUCTURE HARVESTER program (Earl and vonHoldt 2012). Clustering analysis output was summarized and visualized using the Cluster Markov Packager Across K algorithm (CLUMPAK), with default settings (Kopelman et al. 2015). Because the STRUCTURE analysis of the full dataset identified one putative *C. dentata* sample and one putative *C. pumila* var. *ozarkensis* sample as admixed with East Asian *Castanea* spp., we did not include these two samples in further analysis.

Principal components analysis—To understand the partitioning of genetic variation within the North American *Castanea* species and determine the relative placement of *C. alabamensis* samples, we performed principal components analysis (PCA) using the ‘NAC’ dataset. Principal components analysis was performed using the ‘SNPRelate’ package in R, with settings “remove.monosnp=TRUE”, “maf=NaN”, and “missing.rate=NaN”. Because missing genotypes as a result of variation in DNA quality can introduce apparent, but erroneous, population structure in PCA (Patterson et al., 2006), the dataset was further filtered for PCA by identifying individuals with <30% missing SNP loci using TASSEL v5.0 (Bradbury et al. 2007) and specifying for analysis only individuals below this threshold.

RESULTS

Morphological comparisons—Our collections from northern Alabama *Castanea* populations yielded samples of *C. dentata* and several samples that corresponded to the taxonomic description of *C. alabamensis* (Ashe, 1925). Plants identified as *Castanea alabamensis* differed from all other North American *Castanea* taxa in certain key aspects (Table 1; Fig. 1).

Fig. 1 Morphological features distinguishing *C. alabamensis* from *C. dentata* and *C. pumila* var. *ozarkensis*. (A) Eciliate leaf margin of *C. dentata*. (B) Ciliate leaf margin of *C. alabamensis*. (C) Abaxial leaf surface of *C. pumila* var. *ozarkensis* densely covered with stellate trichomes. (D) Abaxial leaf surface of *C. alabamensis* lacking stellate trichomes, but with occasional simple trichomes on laminar surface. Simple trichomes projecting from lateral vein are visible in the lower righthand corner of the image. Scale bars = 5 mm.

While the prevalence of chestnut blight infection prevented us from gathering data on floral/fruit morphology in many wild populations (i.e., plants were typically not reproductively mature), the nut per bur and pistil per bur ratios observed were consistent with the most recent taxonomic treatments of the group (Johnson, 1988; Nixon, 1997; Weakley, 2015). Chestnut blight infection also prevented us from assessing mature plant habit in most North American *Castanea* populations; however, we did observe a few reproductively mature *C. alabamensis* individuals where we could determine that plant habit of mature, healthy trees is typically that of a single-stemmed tree, rather than a shrub, consistent with Ashe's description of the species (1925).

Morphological study of the entire *Castanea* collection at NCU, the voucher specimens from Binkley (2008) and Shaw et al. (2012), and recent collections from the Ruffner Mtn., AL, site studied by Li and Dane (2013) revealed that the *C. × neglecta* populations studied by Binkley (2008) and Shaw et al. (2012) and the Type II *C. dentata* studied by Li and Dane (2013) are morphologically identical to *C. alabamensis* (Ashe, 1925) (Appendix S2). The *C. × neglecta* samples of Binkley (2008) and Shaw et al. (2012) and the Type II *C. dentata* samples of Li and Dane (2013) are identical to Ashe's (1925) original collections of *C. alabamensis* in all aspects of leaf, twig, and flower/fruit morphology that we assessed.

Illumina sequencing and SNP discovery—A total of 489.9 million Illumina reads were obtained for 106 plants (including both technical replicates of the East Asian cultivars) (Appendix S4). Three plants, Haun *C. dentata*, SE56 *C. pumila* var. *pumila*, and TM49 *C. pumila* var. *pumila*, had <100,000 retained reads and were excluded from further processing. Only the higher quality technical replicate of each *C. mollissima* and *C. crenata* cultivar was retained. After removal of the three failed samples and four lower quality technical replicates of

Asian *Castanea* samples, the average clean reads per individual was 4.384 million. In the 103 samples that were retained for genotyping, average coverage depth was 44×. Three datasets were produced for analysis by Stacks 1.45: dataset ‘NAC+EAC’ contained 103 individuals of North American and eastern Asian *Castanea* species and 103,616 SNPs; dataset ‘NAC’ contained 96 individuals of North American *Castanea* species and 190,656 SNPs; and dataset ‘NAC+EAC583’ contained 103 individuals of North American and eastern Asian *Castanea* species and 583 SNPs.

Phylogeny and genetic differentiation—The maximum likelihood phylogeny inferred from the ‘NAC+EAC’ dataset indicated the existence of six distinct groups corresponding to the morphologically-defined taxa present in our dataset: *C. mollissima*, *C. crenata*, *C. dentata*, *C. pumila* var. *ozarkensis*, *C. pumila* var. *pumila*, and *C. alabamensis* (Fig. 2). The North American *Castanea* species formed a monophyletic group. Two main clades were present within the North American *Castanea* group: (1) *C. dentata* and (2) the North American chinquapins, inclusive of *C. alabamensis*, *C. pumila* var. *pumila*, and *C. pumila* var. *ozarkensis*. Bootstrap values for the nodes separating named taxa were 100% in all cases, except the node separating *C. alabamensis* and *C. pumila* var. *pumila*, which was 91% (Fig. 2; Appendix S6).

Fig. 2 Maximum likelihood phylogenetic tree inferred for North American and eastern Asian *Castanea* samples using genome-wide data from 103,616 SNP loci. Asterisks indicate interspecific hybrids identified by STRUCTURE analysis. One asterisk (*) corresponds to an interspecific hybrid derived from *C. pumila* var. *ozarkensis*, *C. dentata*, and an unidentified East Asian *Castanea* sp.; two asterisks (**) correspond to a first-backcross descendant of *C. dentata* and *C. mollissima*. Bootstrap support for the nodes separating the highlighted taxa was 100% in all cases except the node separating *C. alabamensis* and *C. pumila* var. *pumila*, where bootstrap support was 91%. Scale bar is proportional to 0.03 substitutions/site.

The *C. dentata* clade was comprised of two distinct groups: (1) a group containing only individuals from the northern portion of the species' range (samples from Maine, Pennsylvania, and Virginia) and individuals from the Blue Ridge of the southern portion of the species' range (samples from North Carolina, South Carolina, and Georgia) and (2) a group containing only individuals from outside the Blue Ridge in the southern portion of the species' range (samples from Piedmont of South Carolina, Interior Plateau of Tennessee, and Ridge and Valley and Southwestern Appalachians of Alabama). Bootstrap support for the two *C. dentata* groups was 100% (Appendix S6).

Within the *C. pumila* sensu lato clade, *C. alabamensis* and the different botanical varieties of chinquapin, *C. pumila* var. *pumila* and *C. pumila* var. *ozarkensis*, formed three distinct groups. The *C. pumila* var. *pumila* group was comprised of two subgroups: (1) a group containing only individuals from the southern Blue Ridge (samples from North Carolina, South Carolina, and Georgia) and (2) a group containing one individual from the southern Blue Ridge (a sample from South Carolina) and all individuals from the Coastal Plain (samples from South Carolina, Florida, and Georgia). Bootstrap support for the split separating *C. pumila* var. *pumila* samples into two groups was 100% (Appendix S6).

The phylogenetic placement of two samples did not match their putative species assignments. The first sample (dent hyb16_KY), thought to represent a *C. dentata* collection, was placed at an intermediate position between *C. dentata* and the East Asian *Castanea* species on the phylogenetic tree; review of our records showed that this sample was a first-backcross hybrid of *C. dentata* and *C. mollissima* ancestry. The second sample (ozar hyb63_MO), thought to represent *C. pumila* var. *ozarkensis*, was placed at an intermediate position between *C. pumila* sensu lato and *C. dentata* on the phylogenetic tree; STRUCTURE analysis results (detailed

below) showed that this individual derived ancestry from *C. pumila* var. *ozarkensis*, *C. dentata*, and an undetermined East Asian *Castanea* species.

Mean genetic differentiation, as estimated by Weir and Cockerham's (1984) F_{ST} , ranged from 0.71 (*C. mollissima* – *C. pumila* var. *ozarkensis*) to 0.11 (*C. pumila* var. *pumila* – *C. alabamensis*) (Appendix S7). In contrast to what would be expected if *C. alabamensis* were a hybrid taxon derived from *C. dentata* and *C. pumila* var. *pumila*, genetic differentiation was greater between *C. alabamensis* and *C. dentata* ($F_{ST} = 0.40$) than between *C. pumila* var. *pumila* and *C. dentata* ($F_{ST} = 0.38$).

STRUCTURE analyses—STRUCTURE analysis indicated the presence of three genetically differentiated clusters ($K = 3$) in dataset 'NAC+EAC583' (Fig. 3A) (see Appendix S7 for ΔK values) These three clusters corresponded to (1) *C. dentata*, (2) *C. pumila* var. *pumila*, *C. pumila* var. *ozarkensis*, and *C. alabamensis* combined, and (3) the eastern Asian *Castanea* samples. STRUCTURE did not identify a *C. dentata* contribution to *C. alabamensis* genomes (Fig. 3A). Although one *C. alabamensis* sample, AG87_AL, appeared to have a small fraction of *C. dentata* ancestry (*C. dentata* ancestry proportion estimate = 0.05), this sample had the highest percentage of missing SNPs (82% missing SNPs) of any samples in the 'NAC+EAC583' dataset and the lowest number of retained reads of any plants included in analyses; this result is likely an artifact of low sample quality.

Analysis of *C. dentata* samples also failed to detect a genetic contribution from *C. pumila* sensu lato in sympatric populations. However, two *C. pumila* var. *pumila* individuals, both from the allopatric site SS_FL, had evidence of *C. dentata* ancestry. Of these two admixed samples, SS80_FL, was estimated to derive as much as 0.15 of its genome from *C. dentata*.

STRUCTURE analysis of the North American chinquapins *Castanea pumila* sensu lato (including *C. alabamensis*) indicated the presence of two genetically differentiated clusters ($K = 2$)—the first corresponding to *C. pumila* var. *pumila* and the second corresponding to *C. alabamensis* and *C. pumila* var. *ozarkensis* combined (Fig. 3B; see Appendix S7 for ΔK values). Low to moderate levels of admixture between the different botanical varieties of chinquapin were found in the majority of populations analyzed. The single *C. pumila* var. *ozarkensis* individual (ozar_SH_AR) remaining after removal of the interspecific hybrid from Missouri showed evidence of substantial admixture with *C. pumila* var. *pumila*. Evidence of admixture was also present in *C. alabamensis* populations, with higher levels of *C. pumila* var. *pumila* ancestry in the more southerly *alabamensis* populations, CH_AL and AG_AL (Fig. 3B).

Principal component analysis—To provide an additional test of the hybridization hypothesis for *C. alabamensis* and to better understand partitioning of genetic variation within the North American clade, we performed PCA using the 80 North American *Castanea* samples in dataset ‘NAC’ with <30% missing SNP loci. Principal component analysis of the North American *Castanea* identified 32 significant principal components (PCs) that explained 56.52% of the total variation (Appendix S7). The first two principal components provided clear separation of the three currently recognized North American taxa and *C. alabamensis* from one another (Fig. 4). The first component explained 10.96% of the total variation and separated *C. dentata* samples from samples of *C. alabamensis*, *C. pumila* var. *pumila*, and *C. pumila* var.

ozarkensis (Fig. 4; Appendix S7). The second component explained 2.75% of the total variation and separated *C. pumila* var. *pumila*, *C. pumila* var. *ozarkensis*, and *C. alabamensis* from one another. *C. pumila* var. *pumila* samples from the Coastal Plain of Florida and South Carolina were placed along the leftmost portion of the axis created by PC2 in Fig. 4 (i.e., lower values for PC2). However, samples from other parts of the geographical distribution of *C. pumila* var. *pumila* were not completely excluded from the leftmost portion of PC2, as one sample from the Blue Ridge province of South Carolina, CR6_SC, clustered with Coastal Plain samples. *C. pumila* var. *pumila* samples from sites farther inland—specifically, the Blue Ridge of South Carolina, North Carolina, and Georgia—clustered near the midrange of PC2. In contrast to what would be expected for a hybrid taxon, PCA did not place any *C. alabamensis* samples intermediate to the *C. pumila* sensu lato and *C. dentata* clusters.

Fig. 4. Graph of the first two axes from a principal components analysis of 80 individuals representing the putative hybrid, *C. alabamensis*, and all other North American *Castanea* taxa. Dot colors correspond to the following taxa: blue = *C. pumila* var. *pumila*, green = *C. pumila* var. *ozarkensis*, black = *C. alabamensis*, and red = *C. dentata*.

DISCUSSION

C. alabamensis is a distinct variety of North American Castanea that includes plants previously identified as hybrids between Castanea species—While agreeing with previous studies that North American *Castanea* is composed of two species, *C. dentata* and *C. pumila* sensu lato, we have presented compelling evidence that plants identified by Ashe (1925) as *C. alabamensis* comprise a distinct variety within *C. pumila* sensu lato along with *C. pumila* var. *pumila* and *C. pumila* var. *ozarkensis*. Our study represents the first use of genome-wide data to understand evolution and hybridization in the North American *Castanea* species. With different population sampling methods, our analytical approach can be applied to other questions of importance for conservation and restoration of the American chestnut and chinquapins. Our findings provide much needed clarification of the evolutionary relationships, species boundaries, and history of admixture of the North American *Castanea* taxa and will allow workers in the field of chestnut and chinquapin conservation to more efficiently conserve and restore the biodiversity of these imperiled taxa.

In contrast to multiple recent studies of closely-related sympatric species in the Fagaceae (Cavender-Bares and Pahlich, 2009; Leroy et al. 2017; Kim et al. 2018), we find that *C. dentata* and *C. pumila* are both genetically and morphologically discrete where they co-occur, with no evidence of local hybridization and introgression. Jaynes (1964) reported that fertile hybrids can be made among all *Castanea* species. The reproductive barrier between the species is most likely flowering time; *C. pumila* blooms 1-2 weeks earlier than *C. dentata* (P.H. Sisco, J.H. Craddock, unpublished data).

Admixture among taxa—Although we did not document admixture with *C. dentata* in our *C. alabamensis* samples, many other cases of intra- and interspecific admixture were

identified. Perhaps the most interesting case of interspecific admixture that we documented involved samples from a morphologically intriguing population of *C. pumila* var. *pumila* in Florida. Despite the stark morphological differences between these *C. pumila* var. *pumila* individuals and *C. dentata*, two of three individuals had evidence of introgression from *C. dentata* and one of the plants had an ancestry proportion estimate of 0.15 from *C. dentata* in STRUCTURE. These plants, which were perhaps the most morphologically unique of the study, are the rhizomatous subshrub form (only ~0.5-2 m tall at reproductive maturity) of chinquapin that have been treated as *C. alnifolia* (common name: trailing chinquapins) in older taxonomic works (Nuttall, 1818; Sargent, 1919; Ashe, 1922). These Florida *C. pumila* populations are currently separated from the southernmost known *C. dentata* populations by hundreds of kilometers. The finding of *C. dentata* ancestry in these *C. pumila* plants is consistent with the hypothesis that the range of *C. dentata* once extended much farther south than is currently observed (Davis, 1983). While other population genetics studies have inferred a post-Pleistocene expansion of *C. dentata* populations from south to north along the Appalachians (e.g., Gailing and Nelson, 2017), our finding of *C. dentata* ancestry in *C. pumila* populations from Florida is the most direct genetic evidence produced in support of a Pleistocene refugium for *C. dentata* in the southeastern Coastal Plain.

Finally, many cases of intraspecific admixture within the chinquapin clade were also documented in our data. Within *C. pumila* sensu lato, admixture between the two genetic groups identified by STRUCTURE—(1) *C. pumila* var. *pumila* and (2) *C. alabamensis* and *C. pumila* var. *ozarkensis*—was observed in most populations, indicating a recent history of shared ancestry between the different botanical varieties of chinquapin. It should be noted, however, that none of the chinquapin populations analyzed in this study were collected from areas of sympatry for

distinct chinquapin varieties (i.e., where *C. pumila* var. *pumila* and *C. pumila* var. *ozarkensis* occur at the same site or within effective pollination distance). Thus, levels of admixture in sympatric populations containing multiple chinquapin varieties cannot be assessed from our data. Given the low to moderate levels of admixture between chinquapin varieties documented in allopatric populations in the present study, we expect sympatric populations to display even greater signatures of shared ancestry.

CONCLUSIONS

We have used genome-wide SNP data and morphology to show that *C. alabamensis* is a distinct variety of North American chinquapin (*C. pumila* sensu lato) that includes plants previously identified morphologically as hybrids between *Castanea* species. A combination of genome-wide genotyping and morphological analysis was required to better understand the origin of the putative hybrids and the nature of species boundaries in North American *Castanea*. Presumed naturally-occurring admixtures between different *Castanea* species and varieties were found, most notably between *C. pumila* var. *pumila* in northern Florida and *C. dentata*. Our results demonstrate the capability of genomic approaches to resolve previously intractable questions of *Castanea* evolution and highlight the need for further exploration of *Castanea* diversity.

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DATA ACCESSIBILITY

Sequence data are available at the NCBI SRA (BioProject ID: PRJNA541592). All datasets used in analyses are available on GitHub: https://github.com/MTPerkins/Nonhybrid_origin_of_Castanea_alabamensis.

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626 LITERATURE CITED

627 Allendorf, F.W., R.F. Leary, P. Spruell, and J.K. Wenburg. 2001. The problems with hybrids:
628 setting conservation guidelines. *Trends in Ecology and Evolution* 16(11): 613–622.

629 Allendorf, F.W., G. Luikart, and S.N. Aitken. 2013. Conservation and the genetics of
630 populations, 2nd ed. Wiley-Blackwell, West Sussex, UK.

631 Ashe, W.W. 1922. Notes on trees and shrubs of the southeastern United States. *Bulletin of the*
632 *Torrey Botanical Club* 49(9): 265–268.

633 Ashe, W.W. 1925. Notes on woody plants. *The Charleston Museum Quarterly* 1(2): 28–32.

634 Baute, G.J., G.L. Owens, D.G. Bock, and L.H. Rieseberg. 2016. Genome-wide genotyping-by-
635 sequencing data provide a high-resolution view of wild *Helianthus* diversity, genetic structure,
636 and interspecific gene flow. *American Journal of Botany* 103(12): 2170–2177.

637 Biomatters development team. 2018. Geneious 11.1.5. Biomatters Ltd., website:
638 <http://www.geneious.com/> [accessed 01 October 2018].

639 Binkley, M.A. 2008. The phylogeography of North American chestnuts and chinquapins
640 (*Castanea* Mill., Fagaceae). M.S. thesis, The University of Tennessee at Chattanooga,
641 Chattanooga, Tennessee, USA.

642 Bradbury, P.J., Z. Zhang, D.E. Kroon, T.M. Casstevens, Y. Ramdoss, and E.S. Buckler, E.S.
643 2007. TASSEL: Software for association mapping of complex traits in diverse samples.
644 *Bioinformatics* 23: 2633–2635.

645 Camus, A. 1929. Les chataigniers. Monographie des genres *Castanea* et *Castanopsis*.

646 Encyclopédie Économique de Sylviculture, vol. 3. Paul Lechevalier, Paris, France.

647 Catchen, J., A. Amores, P. Hohenlohe, W. Cresko, and J. Postlethwait. 2011. Stacks: building
648 and genotyping loci de novo from short-read sequences. *G3: Genes, Genomes, Genetics* 1:171–
649 182.

650 Catchen, J., P. Hohenlohe, S. Bassham, A. Amores, and W. Cresko. 2013. Stacks: an analysis
651 tools set for population genomics. *Molecular Ecology* 22(11): 3124–3140.

652 Cavender-Bares, J., and A. Pahlich. 2009. Molecular, morphological, and ecological niche
653 differentiation of sympatric sister oak species, *Quercus virginiana* and *Q. geminata* (Fagaceae).
654 *American Journal of Botany* 96(9): 1690–1702.

655 Dane, F. 2009. Comparative phylogeography of *Castanea* species. *Acta Horticulturae* 844:211–
656 222.

657 Davis, M.B. 1983. Quaternary history of deciduous forests of eastern North America and Europe.
658 *Annals of the Missouri Botanical Garden* 70(3): 550–563.

659 Dode, L.A. 1908. Notes dendrologiques. VI. Sur les chataigniers. *Bulletin de la Société*
660 *Dendrologique de France* 8: 140–166.

661 Earl, D.A., and B.M. vonHoldt. 2012. STRUCTURE HARVESTER: a website and program for
662 visualizing STRUCTURE output and implementing the Evanno method. *Conservation Genetics*
663 *Resources* 4(2): 359–361.

664 Eaton, D.A.R., A.L. Hipp, A. González-Rodríguez, and J. Cavender-Bares. 2015. Historical
665 introgression among the American live oaks and the comparative nature of tests for
666 introgression. *Evolution* 69(10): 2587–2601.

667 Elias, T.S. 1971. The genera of Fagaceae in the southeastern United States. *Journal of the Arnold*
668 *Arboretum* 52: 159–195.

669 Ellstrand, N.C., R. Whitkus, and L.H. Rieseberg. 1996. Distribution of spontaneous plant
670 hybrids. *Proceedings of the National Academy of Sciences, USA* 93: 5090–5093.

671 Elshire, R.J., J.C. Glaubitz, Q. Sun, J.A. Poland, K. Kawamoto, E.S. Buckler, and S.E. Mitchell,
672 S.E. 2011. A robust, simple genotyping-by-sequencing (GBS) approach for high diversity
673 species. *PLoS ONE* 6(5): e19379.

674 Escudero, M., D.A.R. Eaton, M. Hahn, and A.L. Hipp. 2014. Genotyping-by-sequencing as a
675 tool to infer phylogeny and ancestral hybridization: a case study in *Carex* (Cyperaceae).
676 *Molecular Phylogenetics and Evolution* 79: 359–367.

677 Evanno, G., S. Regnaut, and J. Goudet. 2005. Detecting the number of clusters of individuals
678 using the software STRUCTURE: a simulation study. *Molecular Ecology* 14: 2611–2620.

679 Falush, D., M. Stephens, and J.K. Pritchard. 2003. Inference of population structure: extensions
680 to linked loci and correlated allele frequencies. *Genetics* 164: 1567–1587.

681 Gailing, O., and C.D. Nelson. 2017. Genetic variation patterns of American chestnut populations
682 at EST-SSRs. *Botany* 95(8): 799–807.

683 Jackiw, R.N., G. Mandil, and H.A. Hager. 2015. A framework to guide the conservation of
684 species hybrids based on ethical and ecological considerations. *Conservation Biology* 29(4):
685 1040–1051.

686 Jaynes, R.A. 1964. Interspecific crosses in the genus *Castanea*. *Silvae Genetica* 13: 146–154.

687 Johnson, G.P. 1988. Revision of *Castanea* sect. *Balanocastanon* (Fagaceae). *Journal of the*
688 *Arnold Arboretum* 69: 25–49.

689 Kim, B.Y., X. Wei, S. Fitz-Gibbon, K.E. Lohmueller, J. Ortego, P.F. Gugger, and V.L. Sork.
690 2018. RADseq data reveal ancient, but not pervasive, introgression between Californian tree and
691 scrub oak species (*Quercus* sect. *Quercus*: Fagaceae). *Molecular Ecology* 27: 4556–4571.

692 Kopelman, N.M., J. Mayzel, M. Jakobsson, N.A. Rosenberg, and I. Mayrose. 2015. CLUMPAK:
693 a program for identifying clustering modes and packaging population structure inferences across
694 *K. Molecular Ecology Resources* 15(5): 1179–1191.

695 Kubisiak, T.L., C.D. Nelson, M.E. Staton, T. Zhebentyayeva, C. Smith, B.A. Olukolu, G.-C.
696 Fang, F.V. Hebard, S. Anagnostakis, N. Wheeler, P.H. Sisco, A.G. Abbott, and R.R. Sederoff.
697 2013. A transcriptome-based genetic map of Chinese chestnut (*Castanea mollissima*) and
698 identification of regions of segmental homology with peach (*Prunus persica*). *Tree Genetics &*
699 *Genomes* 9: 557–571.

700 Kubisiak, T.L., and J.H. Roberds. 2006. Genetic structure of American chestnut populations
701 based on neutral DNA markers. In K.C. Steiner and J.E. Carlson [eds.], *Restoration of American*
702 *Chestnut to Forest Lands – Proceedings of a Conference and Workshop, The North Carolina*
703 *Arboretum, N.C., 2004, 109-122. National Park Service, Washington, DC, USA.*

704 Leroy, T., C. Roux, L. Villate, C. Bodénès, J. Romiguier, J.A.P. Paiva, C. Dossat, J.-M. Aury, C.
705 Plomion, and A. Kremer. 2017. Extensive recent secondary contacts between four European
706 white oak species. *New Phytologist* 214: 865–878.

707 Li, X., and F. Dane. 2013. Comparative chloroplast and nuclear DNA analysis of *Castanea*
708 species in the southern region of the USA. *Tree Genetics and Genomes* 9(1): 107–116.

709 Little, E.L. 1979. Checklist of United States trees (native and naturalized), U.S.D.A. Agriculture
710 Handbook no. 541. U.S. Government Printing Office, Washington, DC, USA.

711 Mallet, J. 2005. Hybridization as an invasion of the genome. *Trends in Ecology and Evolution*
712 20(5): 229–237.

713 McVay, J.D., A.L. Hipp, and P.S. Manos. 2017. A genetic legacy of introgression confounds
714 phylogeny and biogeography in oaks. *Proceedings of the Royal Society B* 284: 20170300.

715 Nixon, K.C. 1997. Fagaceae Dumortier, Beech Family. *In* Flora of North America Editorial
716 Committee (eds.). Flora of North America north of Mexico, vol. 3. Oxford University Press,
717 New York, New York, USA.

718 Nuttall, T. 1818. The genera of North American plants, Vol. 2. D. Heart, Philadelphia,
719 Pennsylvania, USA.

720 Owens, G.L., G.J. Baute, and L.H. Rieseberg. 2016. Revisiting a classic case of introgression:
721 hybridization and gene flow in Californian sunflowers. *Molecular Ecology* 25: 2630–2643.

722 Patterson, N., A.L. Price, and D. Reich. 2006. Population structure and eigenanalysis. *PLoS*
723 *Genetics* 2(12): e190.

724 Pritchard, J.K., M. Stephens, and P. Donnelly. 2000. Inference of population structure using
725 multilocus genotype data. *Genetics* 155: 945–959.

726 R Core Team. 2018. R: A language and environment for statistical computing. R Foundation for
727 Statistical Computing, website: <https://www.R-project.org/> [accessed 01 February 2018].

728 Rieseberg, L.H. 1997. Hybrid origins of plant species. *Annual Review of Ecology and*
729 *Systematics* 28: 359–389.

730 Rochette, N.C., and J.M. Catchen. 2017. Deriving genotypes from RAD-seq short-read data
731 using Stacks. *Nature Protocols* 12: 2640–2659.

732 Sargent, C.S. 1919. North American Trees, IV. *Botanical Gazette* 67(3): 208–242.

733 Shaw, J., J.H. Craddock, and M.A. Binkley. 2012. Phylogeny and phylogeography of North
734 American *Castanea* Mill. (Fagaceae) using cpDNA suggests gene sharing in the Southern
735 Appalachians (*Castanea* Mill., Fagaceae). *Castanea* 77(2): 186–211.

736 Small, J.K. 1933. Manual of the Southeastern Flora. The University of North Carolina Press,
737 Chapel Hill, North Carolina, USA.

738 Soltis, P.S., and D.E. Soltis. 2009. The role of hybridization in plant speciation. *Annual Review*
739 *of Plant Biology* 60: 561–588.

740 Stamatakis, A. 2014. RAxML version 8: a tool for phylogenetic analysis and post-analysis of
741 large phylogenies. *Bioinformatics* 9: 1312–1313.

742 Steiner, K.C., J.W. Westbrook, F.V. Hebard, L.L. Georgi, W.A. Powell, and S.F. Fitzsimmons.
743 2017. Rescue of American chestnut with extraspecific genes following its destruction by a
744 naturalized pathogen. *New Forests* 48(2): 317–336.

745 Stronen, A.V., and P.C. Paquet. 2013. Perspectives on the conservation of wild hybrids.
746 *Biological Conservation* 167: 390–395.

747 Tucker, G.E. 1975. *Castanea pumila* var. *ozarkensis* (Ashe) Tucker, comb. nov. *Arkansas*
748 *Academy of Science Proceedings* 29: 67–69.

749 Turner, K.G., D.I. Huang, Q.C.B. Cronk, and L.H. Rieseberg. 2018. Homogenization of
750 populations in the wildflower, Texas Bluebonnet (*Lupinus texensis*). *Journal of Heredity* 109(2):
751 152–161.

752 Weakley, A.S. 2015. Flora of the Southern and Mid-Atlantic States. The University of North
753 Carolina Herbarium, Chapel Hill, North Carolina, USA.

754 Weir, B.S., and C.C. Cockerham. 1984. Estimating *F*-statistics for the analysis of population
755 structure. *Evolution* 38(6): 1358–1370.

756 Whitham, T.G., P.A. Morrow, and B.M. Potts. 1991. Conservation of hybrid plants. *Science* 254:
757 779–780.

758 Whitney, K.D., J.R. Ahern, L.G. Campbell, L.P. Albert, and M.S. King. 2010. Patterns of
759 hybridization in plants. *Perspectives in Plant Ecology, Evolution and Systematics* 12(3): 175–
760 182.

Wu, T.D., and S. Nacu. 2010. Fast and SNP-tolerant detection of complex variants and splicing in short reads. *Bioinformatics* 26(7): 873–881.

Zhao, P., H.-J. Zhou, D. Potter, Y.-H. Hu, X.-J. Feng, M. Dang, L. Feng, S. Zulfiqar, W.-Z. Liu, G.-F. Zhao, and K. Woeste. 2018. Population genetics, phylogenomics and hybrid speciation of *Juglans* in China determined from whole chloroplast genomes, transcriptomes, and genotyping-by-sequencing (GBS). *Molecular Phylogenetics and Evolution* 126: 250–265.

Zhebentyayeva, T., V. Shankar, R. Scorza, A. Callahan, M. Ravelonandro, S. Castro, T. DeJong, C.A. Saski, and C. Dardick. 2019. Genetic characterization of world-wide *Prunus domestica* (plum) germplasm using sequence-based genotyping. *Horticulture Research* 6:12.

Zheng, X., D. Levine, J. Shen, S. Gogarten, C. Laurie, and B. Weir. 2012. A high-performance computing toolset for relatedness and principal component analysis of SNP data. *Bioinformatics* 28: 3326–3328.

APPENDICES

Appendix S1. Sample voucher and NCBI SRA information for individual *Castanea* plants genotyped. GBS IDs correspond to sample names in NCBI SRA BioProject ID “PRJNA541592”.

County, State or cultivar/selection name	Sample site code	GBS ID	Taxon	Herbarium voucher	DNA tube label
Charleston, SC	AW	1	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-1	AW1
Charleston, SC	AW	5	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-2	AW2
Charleston, SC	AW	9	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-3	AW3
Charleston, SC	AW	13	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-4	AW4
Charleston, SC	AW	17	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-5	AW5
Charleston, SC	AW	23	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-7	AW7
Charleston, SC	AW	28	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-8	AW8
Charleston, SC	AW	32	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-9	AW9
Charleston, SC	AW	36	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-10	AW10
Sumter, SC	WB	41	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-11	WB1
Sumter, SC	WB	44	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-12	WB2
Transylvania, NC	GO	18	<i>C. dentata</i>	Perkins 15-13	G1
Transylvania, NC	GO	20	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-14	G2
Transylvania, NC	GO	24	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-15	G3
Oconee, SC	CR	7	<i>C. dentata</i>	Perkins 15-16	CR1
Oconee, SC	CR	11	<i>C. dentata</i>	Perkins 15-17	CR2
Oconee, SC	CR	15	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-18	CR3
Oconee, SC	CR	19	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-19	CR4
Oconee, SC	CR	21	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-20	CR5
Oconee, SC	CR	25	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-21	CR6
Oconee, SC	CR	30	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-22	CR7
Oconee, SC	BF	29	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-23	BF1
Oconee, SC	BF	33	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-24	BF2
Oconee, SC	BF	37	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-25	BF3
Oconee, SC	BF	42	<i>C. pumila</i> var. <i>pumila</i>	Perkins 15-26	BF4
Oconee, SC	BF	45	<i>C. dentata</i>	Perkins 15-27	BF5
Oconee, SC	BF	3	<i>C. dentata</i>	Perkins 15-28	BF6
Pickens, SC	CE	2	<i>C. dentata</i>	Perkins 15-29	CEF1
Pickens, SC	CE	6	<i>C. dentata</i>	Perkins 15-30	CEF2
Pickens, SC	CE	10	<i>C. dentata</i>	Perkins 15-31	CEF3
Pickens, SC	CE	14	<i>C. dentata</i>	Perkins 15-32	CEF4
Suwannee, FL	SS	80	<i>C. pumila</i> var. <i>pumila</i>	Perkins 17-1	SS2
Suwannee, FL	SS	81	<i>C. pumila</i> var. <i>pumila</i>	Perkins 17-2	SS3
Suwannee, FL	SS	82	<i>C. pumila</i> var. <i>pumila</i>	Perkins 17-3	SS4

Lincoln, TN	LI	83	<i>C. dentata</i>	Perkins 17-4	TNLIN1
Calhoun, AL	CH	90	<i>C. alabamensis</i>	Perkins 17-5	CH21
Calhoun, AL	CH	91	<i>C. dentata</i>	Perkins 17-6	CH22
Calhoun, AL	CH	92	<i>C. alabamensis</i>	Perkins 17-7	CH23
Calhoun, AL	CH	93	<i>C. alabamensis</i>	Perkins 17-8	CH24
Calhoun, AL	CH	94	<i>C. alabamensis</i>	Perkins 17-9	CH25
Calhoun, AL	CH	95	<i>C. alabamensis</i>	Perkins 17-10	CH26
Calhoun, AL	CH	96	<i>C. alabamensis</i>	Perkins 17-11	CH27
Cleburne, AL	HU	67	<i>C. alabamensis</i>	Perkins 17-12	H1
Cleburne, AL	HU	68	<i>C. dentata</i>	Perkins 17-13	H2
Cleburne, AL	HU	69	<i>C. dentata</i>	Perkins 17-14	H3
Cleburne, AL	HU	70	<i>C. dentata</i>	Perkins 17-15	H4
Cleburne, AL	HU	86	<i>C. alabamensis</i>	Perkins 17-16	H9
Cleburne, AL	HU	89	<i>C. alabamensis</i>	Perkins 17-17	H6
Cleburne, AL	FR	71	<i>C. dentata</i>	Perkins 17-18	F1
Cleburne, AL	FR	72	<i>C. dentata</i>	Perkins 17-19	F5
Cleburne, AL	FR	84	<i>C. dentata</i>	Perkins 17-20	F3
Clay, AL	AG	73	<i>C. alabamensis</i>	Perkins 17-21	AG1
Clay, AL	AG	74	<i>C. dentata</i>	Perkins 17-22	AG2
Clay, AL	AG	75	<i>C. alabamensis</i>	Perkins 17-23	AG3
Clay, AL	AG	76	<i>C. alabamensis</i>	Perkins 17-24	AG6
Clay, AL	AG	77	<i>C. dentata</i>	Perkins 17-25	AG7
Clay, AL	AG	85	<i>C. alabamensis</i>	Perkins 17-26	AG4
Clay, AL	AG	87	<i>C. alabamensis</i>	Perkins 17-27	AG5
Morgan, AL	LA	78	<i>C. dentata</i>	Perkins 17-28	L14
Morgan, AL	LA	79	<i>C. dentata</i>	Perkins 17-29	L19
Morgan, AL	LA	88	<i>C. dentata</i>	Perkins 17-30	L10
Floyd, GA	JM	22	<i>C. alabamensis</i>	Binkley 412	412
Floyd, GA	PO	38	<i>C. alabamensis</i>	Binkley 420	420
Cherokee, NC	JB	50	<i>C. dentata</i>	Binkley 507	507
Cherokee, NC	JB	51	<i>C. dentata</i>	Binkley 513	513
Cherokee, NC	JB	52	<i>C. dentata</i>	Binkley 514	514
Cherokee, NC	SE	53	<i>C. pumila</i> var. <i>pumila</i>	Binkley 124	124
Cherokee, NC	SE	54	<i>C. pumila</i> var. <i>pumila</i>	Binkley 525	525
Cherokee, NC	SE	55	<i>C. pumila</i> var. <i>pumila</i>	Binkley 126	126
Cherokee, NC	SE	56	<i>C. pumila</i> var. <i>pumila</i>	Binkley 527	527
Cherokee, NC	SE	57	<i>C. pumila</i> var. <i>pumila</i>	Binkley 528	528
Cherokee, NC	SE	58	<i>C. pumila</i> var. <i>pumila</i>	Binkley 529	529
Cherokee, NC	SE	59	<i>C. dentata</i>	Binkley 531	531
Cherokee, NC	SE	60	<i>C. pumila</i> var. <i>pumila</i>	Binkley 532	532
Cherokee, NC	SE	61	<i>C. pumila</i> var. <i>pumila</i>	Binkley 535	535
Cherokee, NC	SE	62	<i>C. pumila</i> var. <i>pumila</i>	Binkley 149	149
Baker, GA	JR	26	<i>C. pumila</i> var. <i>pumila</i>	Binkley 735	735
Pickens, GA	TM	27	<i>C. dentata</i>	Binkley 645	645
Pickens, GA	TM	31	<i>C. dentata</i>	Binkley 646	646

Pickens, GA	TM	35	<i>C. pumila</i> var. <i>pumila</i>	Binkley 647	647
Pickens, GA	TM	40	<i>C. pumila</i> var. <i>pumila</i>	Binkley 648	648
Pickens, GA	TM	43	<i>C. pumila</i> var. <i>pumila</i>	Binkley 649	649
Pickens, GA	TM	46	<i>C. pumila</i> var. <i>pumila</i>	Binkley 650	650
Pickens, GA	TM	47	<i>C. pumila</i> var. <i>pumila</i>	Binkley 651	651
Pickens, GA	TM	48	<i>C. pumila</i> var. <i>pumila</i>	Binkley 652	652
Pickens, GA	TM	49	<i>C. pumila</i> var. <i>pumila</i>	Binkley 653	653
Knox, ME	KN	34	<i>C. dentata</i>	Binkley 671	671
Piscataquis, ME	AT	39	<i>C. dentata</i>	Binkley 672	672
Waldo, ME	WA	65	<i>C. dentata</i>	Binkley 673	673
Piscataquis, ME	SB	66	<i>C. dentata</i>	Binkley 674	674
Huntingdon, PA	RO	4	<i>C. dentata</i>	Binkley 669	669
Warren, PA	WR	8	<i>C. dentata</i>	Binkley 710	710
Westmoreland, PA	WE	12	<i>C. dentata</i>	Binkley 675	675
Sharp, AR	SH	64	<i>C. pumila</i> var. <i>ozarkensis</i>	Binkley 705	705
Lawrence, MO	MO	63	<i>ozarkensis</i> complex hybrid	Binkley 694	694
Adair, KY	KY	16	<i>dentata-mollissima</i> hybrid	NA	KYADA1
Buncombe, NC	DO	102	<i>C. dentata</i>	Sisco & Perkins 15-1	NCDOT
Avery, NC	CB	103	<i>C. dentata</i>	NA	Cranberry
Washington, VA	WN	97	<i>C. dentata</i>	NA	AD98
cultivar 'Nanking'	NK	100	<i>C. mollissima</i>	NA	Nanking
cultivar 'Mahogany'	MY	98	<i>C. mollissima</i>	NA	Mahogany
selection 'Fort Defiance'	FD	104	<i>C. crenata</i>	NA	Ft. Defiance
selection 'Morrow Mountain'	MM	107	<i>C. crenata</i>	NA	Morrow Mtn.
selection 'Rita 1'	RI	108	<i>C. crenata</i>	NA	Rita 1
selection 'Rita 2'	RA	110	<i>C. crenata</i>	NA	Rita 2

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Appendix S2. Annotations of plants studied by Binkley (2008), Shaw et al. (2012), and Li and Dane (2013). Two plants from the study of Binkley (2008) and Shaw et al. (2012) were genotyped here. Herbarium vouchers from all plants listed below were assessed for morphological traits listed in Table 1 and annotated.

County, State	Sample site	Sample ID in previous study	Taxonomic determination of previous study	Reference	Taxonomic determination of present study	Herbarium voucher	Sample ID in present study
Floyd, GA	the Pocket	420	<i>C. x neglecta</i>	Binkley (2008); Shaw et al. (2012)	<i>C. alabamensis</i>	Binkley 420	PO38_AL
Floyd, GA	Johns Mountain	412	<i>C. x neglecta</i>	Binkley (2008); Shaw et al. (2012)	<i>C. alabamensis</i>	Binkley 412	JM22_AL
Jefferson, AL	Ruffner Mountain	AL-M65	Type II <i>C. dentata</i>	Li and Dane (2013)	<i>C. alabamensis</i>	Perkins 15-101	Not genotyped
Jefferson, AL	Ruffner Mountain	AL-M35	Type II <i>C. dentata</i>	Li and Dane (2013)	<i>C. alabamensis</i>	Perkins 15-102	Not genotyped
Jefferson, AL	Ruffner Mountain	AL-M38	Type I <i>C. dentata</i>	Li and Dane (2013)	<i>C. dentata</i>	Deason MS38	Not genotyped
Jefferson, AL	Ruffner Mountain	AL-M68	Type II <i>C. dentata</i>	Li and Dane (2013)	<i>C. alabamensis</i>	Deason MS68	Not genotyped
Jefferson, AL	Ruffner Mountain	AL-4CN	Type I <i>C. dentata</i>	Li and Dane (2013)	<i>C. dentata</i>	Deason 4CN	Not genotyped
Jefferson, AL	Ruffner Mountain	AL-7CN	Type I <i>C. dentata</i>	Li and Dane (2013)	<i>C. dentata</i>	Deason 7CN	Not genotyped

Appendix S3. Herbarium accessions housed at the University of North Carolina Chapel Hill Herbarium (NCU) that were assessed for morphological characters and annotated. Vouchers collected by W.W. Ashe and used for his description of *C. alabamensis* are denoted with “collected by W.W. Ashe” in the identification remarks column. SERNEC catalog numbers are unique to each accession in the Southeast Regional Network of Expertise and Collections (SERNEC) project. Images of each accession, along with locality and collector information, can be accessed at the SERNEC website: <http://sernecportal.org/portal/index.php>.

SERNEC catalog number	Taxonomic determination by Perkins et al.	Identification remarks
NCU00153249	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153250	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153251	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153252	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153253	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153254	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153255	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153248	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153247	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153241	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153242	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153243	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153244	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153246	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153238	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153239	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153240	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153231	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153232	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153233	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153235	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153237	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00027970	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153211	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153212	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153213	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00113544	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153215	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153216	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153217	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153204	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153199	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153130	<i>Castanea dentata</i>	
NCU00153128	<i>Castanea dentata</i>	
NCU00153127	<i>Castanea dentata</i>	

NCU00153125	Castanea dentata	
NCU00153124	Castanea dentata	
NCU00153123	Castanea dentata	
NCU00153057	Castanea dentata	
NCU00153131	Castanea dentata	Occasional simple trichome at leaf margin
NCU00153058	Castanea dentata	
NCU00153132	Castanea dentata	
NCU00153048	Castanea dentata	
NCU00027754	Castanea dentata	
NCU00153049	Castanea dentata	
NCU00145692	Castanea dentata	Rare stellate trichome
NCU00153136	Castanea dentata	
NCU00153134	Castanea dentata	Rare stellate trichome
NCU00153133	Castanea dentata	
NCU00153060	Castanea dentata	
NCU00153897	Castanea dentata	
NCU00153822	Castanea pumila var. pumila	
NCU00153823	Castanea pumila var. pumila	
NCU00153814	Castanea pumila var. pumila	
NCU00153809	Castanea pumila var. pumila	Specimen has section of stolon
NCU00153811	Castanea pumila var. pumila	But see Harbison's note re: "C. alnifolia past blooming while ashei is yet in full bloom..."
NCU00153798	Castanea pumila var. pumila	
NCU00153947	Castanea pumila var. pumila	
NCU00153800	Castanea pumila var. pumila	
NCU00153799	Castanea pumila var. pumila	
NCU00153794	Castanea pumila var. pumila	
NCU00153796	Castanea pumila var. pumila	
NCU00153804	Castanea pumila var. pumila	
NCU00153806	Castanea pumila var. pumila	
NCU00153813	Castanea pumila var. pumila	
NCU00153812	Castanea pumila var. pumila	
NCU00153802	Castanea pumila var. pumila	
NCU00153807	Castanea pumila var. pumila	
NCU00153144	Castanea dentata	
NCU00153143	Castanea dentata	
NCU00153140	Castanea dentata	
NCU00153139	Castanea dentata	
NCU00153138	Castanea dentata	
NCU00153160	Castanea dentata	
NCU00153209	Castanea pumila var. pumila	
NCU00153208	Castanea pumila var. pumila	
NCU00153207	Castanea pumila var. pumila	

NCU00153206	Castanea pumila var. pumila
NCU00153205	Castanea pumila var. pumila
NCU00153220	Castanea pumila var. pumila
NCU00055657	Castanea pumila var. pumila
NCU00153229	Castanea pumila var. pumila
NCU00153228	Castanea pumila var. pumila
NCU00153227	Castanea pumila var. pumila
NCU00153225	Castanea pumila var. pumila
NCU00153223	Castanea pumila var. pumila
NCU00153222	Castanea pumila var. pumila
NCU00153221	Castanea pumila var. pumila
NCU00153192	Castanea pumila var. pumila
NCU00008712	Castanea pumila var. pumila
NCU00153190	Castanea pumila var. pumila
NCU00093627	Castanea pumila var. pumila
NCU00153189	Castanea pumila var. pumila
NCU00153188	Castanea pumila var. pumila
NCU00153187	Castanea pumila var. pumila
NCU00153193	Castanea pumila var. pumila
NCU00153198	Castanea pumila var. pumila
NCU00153197	Castanea pumila var. pumila
NCU00153196	Castanea pumila var. pumila
NCU00153194	Castanea pumila var. pumila
NCU00008710	Castanea pumila var. pumila
NCU00153203	Castanea pumila var. pumila
NCU00153202	Castanea pumila var. pumila
NCU00153201	Castanea pumila var. pumila
NCU00153792	Castanea pumila var. pumila
NCU00153420	Castanea pumila var. pumila
NCU00153781	Castanea pumila var. pumila
NCU00153782	Castanea pumila var. pumila
NCU00153784	Castanea pumila var. pumila
NCU00153790	Castanea pumila var. pumila
NCU00153788	Castanea pumila var. pumila
NCU00153789	Castanea pumila var. pumila
NCU00153787	Castanea pumila var. pumila
NCU00153793	Castanea pumila var. pumila
NCU00153867	Castanea dentata
NCU00153866	Castanea dentata
NCU00153864	Castanea dentata
NCU00153862	Castanea dentata
NCU00153861	Castanea dentata
NCU00153860	Castanea dentata
NCU00153859	Castanea dentata
NCU00153403	Castanea pumila var. pumila

NCU00153379	Castanea pumila var. pumila	
NCU00153946	Castanea pumila var. pumila	Tips of leaves not acuminate, cupules four per spike.
NCU00153378	Castanea pumila var. pumila	
NCU00153377	Castanea pumila var. pumila	
NCU00153387	Castanea pumila var. pumila	
NCU00153376	Castanea pumila var. pumila	
NCU00153392	Castanea pumila var. pumila	
NCU00153391	Castanea pumila var. pumila	
NCU00153389	Castanea pumila var. pumila	
NCU00153401	Castanea pumila var. pumila	
NCU00153400	Castanea pumila var. pumila	
NCU00153399	Castanea pumila var. pumila	
NCU00153398	Castanea pumila var. pumila	
NCU00153396	Castanea pumila var. pumila	
NCU00153395	Castanea pumila var. pumila	
NCU00153393	Castanea pumila var. pumila	
NCU00153406	Castanea pumila var. pumila	
NCU00153175	Castanea pumila var. pumila	Abaxial leaf surfaces densely covered with stellate trichomes. Deep sinuses along leaf margin. Our population genomic work demonstrates that plants with similar morphology from Blue Ridge region of NC are not hybrids.
NCU00153173	Castanea pumila var. pumila	Leaf shape is lanceolate, like commonly seen in <i>C. dentata</i> , and abaxial surface is densely covered with stellate trichomes, like commonly seen in <i>C. pumila</i> . Yet our population genomic work demonstrates that plants with similar morphology are not hybrids.
NCU00153176	Castanea x neglecta	Note mixture of stellate trichomes and long, simple trichomes on abaxial surface; long, adpressed, simple trichomes on leaf veins.
NCU00129230	Castanea mollissima	
NCU00153896	Castanea dentata	
NCU00153895	Castanea mollissima	
NCU00153894	Castanea dentata	
NCU00153893	Castanea dentata	
NCU00153892	Castanea dentata	
NCU00050100	Castanea alabamensis	Collected by W.W. Ashe
NCU00053434	Castanea alabamensis	Collected by W.W. Ashe

NCU00050202	Castanea alabamensis	
NCU00053426	Castanea alabamensis	Collected by W.W. Ashe
NCU00053427	Castanea alabamensis	Collected by W.W. Ashe
NCU00053423	Castanea pumila var. pumila	Note 10 cupules per spike, slender twigs, and stellate interveinal trichomes on abaxial surface.
NCU00053421	Castanea pumila var. pumila	Note stellate trichomes on interveinal lamina of abaxial surface.
NCU00053420	Castanea alabamensis	Collected by W.W. Ashe
NCU00053410	Castanea pumila var. ozarkensis	
NCU00053415	Castanea pumila var. ozarkensis	
NCU00053404	Castanea pumila var. ozarkensis	
NCU00082746	Castanea pumila var. ozarkensis	
NCU00053424	Castanea pumila var. pumila	
NCU00053409	Castanea pumila var. ozarkensis	
NCU00053416	Castanea pumila var. pumila	See stellate trichomes on abaxial lamina.
NCU00053425	Castanea pumila var. pumila	See stellate trichomes on abaxial lamina, slender twigs, and four cupules on spike.
NCU00053417	Castanea pumila var. pumila	See stellate trichomes on abaxial lamina.
NCU00053411	Castanea pumila var. ozarkensis	
NCU00053408	Castanea pumila var. ozarkensis	
NCU00053414	Castanea pumila var. ozarkensis	
NCU00053403	Castanea pumila var. ozarkensis	Has stellate trichomes on lamina surface. Has simple adpressed trichomes on veins. On young, unexpanded leaves, it appears that abaxial lamina has simple trichomes, but these are actually on veins.
NCU00053407	Castanea pumila var. ozarkensis	
NCU00053412	Castanea pumila var. ozarkensis	
NCU00053406	Castanea pumila var. ozarkensis	
NCU00053405	Castanea pumila var. ozarkensis	
NCU00053431	Castanea alabamensis	Collected by W.W. Ashe. Note complete absence of stellate trichomes on abaxial lamina. Note ciliate leaf margin. Note glabrous, glaucous abaxial lamina.
NCU00053429	Castanea alabamensis	Collected by Mrs. H.J. McDowell for W.W. Ashe. Note ciliate leaf margin, absence of trichomes on abaxial lamina, presence of simple trichomes on veins only.

NCU00053428	<i>Castanea alabamensis</i>	Collected by W.W. Ashe. Note presence of leaf trichomes only on blade margin and veins. Complete absence of stellate trichomes on leaves.
NCU00053433	<i>Castanea alabamensis</i>	Collected by W.W. Ashe
NCU00050092	<i>Castanea alabamensis</i>	Collected by Mrs. H.J. McDowell for W.W. Ashe
NCU00050093	<i>Castanea alabamensis</i>	Collected by Mrs. H.J. McDowell for W.W. Ashe
NCU00050094	<i>Castanea alabamensis</i>	Collected by W.W. Ashe. Ciliate leaf margin. No stellate trichomes on abaxial lamina. Simple trichomes on veins of abaxial leaf surface.
NCU00050095	<i>Castanea alabamensis</i>	Collected by W.W. Ashe
NCU00050096	<i>Castanea alabamensis</i>	Collected by W.W. Ashe
NCU00053432	<i>Castanea alabamensis</i>	Collected by W.W. Ashe
NCU00050097	<i>Castanea pumila</i> var. <i>pumila</i>	Note stellate trichomes on abaxial lamina of leaves from both plants represented on this sheet.
NCU00050098	<i>Castanea pumila</i> var. <i>pumila</i>	Note stellate trichomes on abaxial lamina.
NCU00153328	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00085046	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153327	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153326	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153325	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153323	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153322	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153321	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153891	<i>Castanea dentata</i>	
NCU00153890	<i>Castanea dentata</i>	
NCU00153936	<i>Castanea mollissima</i>	
NCU00153935	<i>Castanea mollissima</i>	
NCU00153934	<i>Castanea mollissima</i>	
NCU00008707	<i>Castanea mollissima</i>	
NCU00153937	<i>Castanea mollissima</i>	
NCU00153940	<i>Castanea mollissima</i>	
NCU00153939	<i>Castanea mollissima</i>	
NCU00008708	<i>Castanea mollissima</i>	
NCU00153838	<i>Castanea dentata</i>	
NCU00153837	<i>Castanea dentata</i>	
NCU00153836	<i>Castanea dentata</i>	
NCU00153786	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153785	<i>Castanea pumila</i> var. <i>pumila</i>	
NCU00153858	<i>Castanea dentata</i>	

NCU00153857	Castanea dentata	
NCU00153856	Castanea dentata	
NCU00153881	Castanea dentata	
NCU00153880	Castanea dentata	
NCU00153879	Castanea dentata	
NCU00153878	Castanea dentata	
NCU00153876	Castanea dentata	
NCU00153874	Castanea dentata	
NCU00153872	Castanea pumila var. pumila	Note the abundant stellate trichomes on abaxial leaf surface, erect simple trichomes on abaxial midrib, and ciliate leaf margin.
NCU00153874	Castanea dentata	
NCU00153871	Castanea dentata	
NCU00153870	Castanea dentata	
NCU00153869	Castanea dentata	
NCU00153868	Castanea dentata	
NCU00153908	Castanea dentata	
NCU00153907	Castanea dentata	
NCU00153906	Castanea dentata	
NCU00153905	Castanea dentata	
NCU00153904	Castanea dentata	
NCU00140424	Castanea alabamensis	Note ciliate leaf margin and abundant, erect, simple trichomes on leaf abaxial veins.
NCU00153903	Castanea alabamensis	Note ciliate leaf margin and abundant, erect, simple trichomes on leaf abaxial veins.
NCU00153901	Castanea alabamensis	Note ciliate leaf margin and abundant, erect, simple trichomes on leaf abaxial veins.
NCU00153900	Castanea dentata	Note eciliate leaf margin, abundant glandular trichomes on abaxial leaf surface, and occasional appressed trichome on abaxial leaf veins.
NCU00153899	Castanea dentata	Although simple trichomes are infrequently present on leaf margin, there are far fewer trichomes here than is found C. alabamensis. Also note long, typically appressed trichomes on abaxial leaf veins and lack of erect, simple trichomes on abaxial leaf surface.
NCU00153898	Castanea dentata	

NCU00153909	Castanea dentata	Note frequent glandular trichomes on abaxial leaf surface, eciliate leaf margin, and long, often appressed, simple trichomes on abaxial leaf midrib.
NCU00153912	Castanea dentata	
NCU00153911	Castanea dentata	
NCU00153910	Castanea dentata	
NCU00153259	Castanea pumila var. pumila	
NCU00153258	Castanea pumila var. pumila	
NCU00153257	Castanea pumila var. pumila	
NCU00153263	Castanea pumila var. pumila	
NCU00153260	Castanea pumila var. pumila	
NCU00153270	Castanea pumila var. pumila	
NCU00153268	Castanea pumila var. pumila	
NCU00153266	Castanea pumila var. pumila	
NCU00153264	Castanea pumila var. pumila	
NCU00153414	Castanea pumila var. pumila	
NCU00153278	Castanea pumila var. pumila	
NCU00153277	Castanea pumila var. pumila	
NCU00116436	Castanea pumila var. pumila	
NCU00116437	Castanea pumila var. pumila	
NCU00153271	Castanea pumila var. pumila	
NCU00153281	Castanea pumila var. pumila	
NCU00153282	Castanea pumila var. pumila	
NCU00153280	Castanea pumila var. pumila	
NCU00153287	Castanea pumila var. pumila	
NCU00153285	Castanea pumila var. pumila	
NCU00153284	Castanea pumila var. pumila	
NCU00075919	Castanea pumila var. pumila	
NCU00153291	Castanea pumila var. pumila	
NCU00153290	Castanea pumila var. pumila	
NCU00153288	Castanea pumila var. pumila	
NCU00153149	Castanea dentata	
NCU00153151	Castanea dentata	
NCU00153152	Castanea dentata	
NCU00153153	Castanea dentata	
NCU00153154	Castanea dentata	
NCU00135776	Castanea pumila var. pumila	Note abundant stellate trichomes on abaxial leaf surface.
NCU00153157	Castanea dentata	
NCU00153158	Castanea dentata	
NCU00153159	Castanea dentata	
NCU00153185	Castanea pumila var. pumila	
NCU00153184	Castanea pumila var. pumila	

NCU00153179	Castanea pumila var. pumila
NCU00153181	Castanea pumila var. pumila
NCU00153183	Castanea pumila var. pumila
NCU00008711	Castanea pumila var. pumila
NCU00153121	Castanea dentata
NCU00008703	Castanea dentata
NCU00153122	Castanea dentata
NCU00153047	Castanea crenata
NCU00008705	Castanea dentata
NCU00153051	Castanea dentata
NCU00153052	Castanea dentata
NCU00153054	Castanea dentata
NCU00153053	Castanea dentata
NCU00153056	Castanea dentata
NCU00153055	Castanea dentata
NCU00153336	Castanea pumila var. pumila
NCU00153337	Castanea pumila var. pumila
NCU00153339	Castanea pumila var. pumila
NCU00153338	Castanea pumila var. pumila
NCU00153340	Castanea pumila var. pumila
NCU00014050	Castanea pumila var. pumila
NCU00153341	Castanea pumila var. pumila
NCU00153342	Castanea pumila var. pumila
NCU00153344	Castanea pumila var. pumila
NCU00153345	Castanea pumila var. pumila
NCU00153346	Castanea pumila var. pumila
NCU00153348	Castanea pumila var. pumila
NCU00153335	Castanea pumila var. pumila
NCU00153334	Castanea pumila var. pumila
NCU00153333	Castanea pumila var. pumila
NCU00153332	Castanea pumila var. pumila
NCU00153331	Castanea pumila var. pumila
NCU00115121	Castanea pumila var. pumila
NCU00153329	Castanea pumila var. pumila
NCU00153178	Castanea pumila var. pumila
NCU00153304	Castanea pumila var. pumila
NCU00153294	Castanea pumila var. pumila
NCU00153295	Castanea pumila var. pumila
NCU00153297	Castanea pumila var. pumila
NCU00028020	Castanea pumila var. pumila
NCU00153298	Castanea pumila var. pumila
NCU00008709	Castanea pumila var. pumila
NCU00153299	Castanea pumila var. pumila
NCU00153300	Castanea pumila var. pumila
NCU00153293	Castanea pumila var. pumila

NCU00153320	Castanea pumila var. pumila	
NCU00153305	Castanea pumila var. pumila	
NCU00153306	Castanea pumila var. pumila	
NCU00153307	Castanea pumila var. pumila	
NCU00153310	Castanea pumila var. pumila	
NCU00153309	Castanea pumila var. pumila	
NCU00153312	Castanea pumila var. pumila	
NCU00153313	See note	Specimens from two different species are glued to this sheet. A = C. dentata; B = C. pumila var. pumila.
NCU00153313	See note	Specimens from two different species are glued to this sheet. A = C. dentata; B = C. pumila var. pumila.
NCU00153314	Castanea pumila var. pumila	
NCU00153315	Castanea pumila var. pumila	
NCU00153317	Castanea pumila var. pumila	
NCU00153319	Castanea pumila var. pumila	
NCU00153318	Castanea pumila var. pumila	
NCU00153350	Castanea mollissima	
NCU00110700	Castanea pumila var. pumila	
NCU00153349	Castanea pumila var. pumila	
NCU00153402	Castanea pumila var. pumila	
NCU00153404	Castanea pumila var. pumila	
NCU00153407	Castanea pumila var. pumila	
NCU00153418	Castanea pumila var. pumila	
NCU00153416	Castanea pumila var. pumila	
NCU00153413	Castanea pumila var. pumila	
NCU00153412	Castanea pumila var. pumila	
NCU00153411	Castanea pumila var. pumila	
NCU00153410	Castanea pumila var. pumila	
NCU00153409	Castanea pumila var. pumila	
NCU00153408	Castanea pumila var. pumila	
NCU00153419	Castanea pumila var. pumila	
NCU00153368	Castanea pumila var. pumila	
NCU00153374	Castanea pumila var. pumila	
NCU00153375	Castanea pumila var. pumila	
NCU00153386	Castanea pumila var. pumila	
NCU00153385	Castanea pumila var. pumila	
NCU00153382	Castanea pumila var. pumila	
NCU00153381	Castanea pumila var. pumila	
NCU00153913	Castanea dentata	
NCU00153817	Castanea pumila var. pumila	
NCU00153816	Castanea pumila var. pumila	

NCU00153815	Castanea pumila var. pumila
NCU00153824	Castanea pumila var. pumila
NCU00153826	Castanea mollissima
NCU00153825	Castanea mollissima
NCU00153840	Castanea dentata
NCU00153839	Castanea dentata
NCU00154913	Castanea pumila var. pumila
NCU00154906	Castanea pumila var. pumila
NCU00154907	Castanea pumila var. pumila
NCU00154908	Castanea pumila var. pumila
NCU00154909	Castanea pumila var. pumila
NCU00154911	Castanea pumila var. pumila
NCU00154912	Castanea pumila var. pumila
NCU00154905	Castanea pumila var. pumila
NCU00086049	Castanea pumila var. pumila
NCU00135544	Castanea pumila var. pumila
NCU00154896	Castanea pumila var. pumila
NCU00154897	Castanea pumila var. pumila
NCU00154898	Castanea pumila var. pumila
NCU00154899	Castanea pumila var. pumila
NCU00154901	Castanea pumila var. pumila
NCU00154902	Castanea pumila var. pumila
NCU00154903	Castanea pumila var. pumila
NCU00154904	Castanea pumila var. pumila
NCU00154894	Castanea pumila var. pumila
NCU00154876	Castanea dentata
NCU00154887	Castanea pumila var. pumila
NCU00154888	Castanea pumila var. pumila
NCU00154890	Castanea pumila var. pumila
NCU00154892	Castanea pumila var. pumila
NCU00154875	Castanea pumila var. pumila
NCU00154893	Castanea pumila var. pumila
NCU00154863	Castanea pumila var. pumila
NCU00154864	Castanea pumila var. pumila
NCU00154865	Castanea pumila var. pumila
NCU00154866	Castanea pumila var. pumila
NCU00154867	Castanea pumila var. pumila
NCU00154868	Castanea pumila var. pumila
NCU00154869	Castanea pumila var. pumila
NCU00154873	Castanea pumila var. pumila
NCU00154871	Castanea pumila var. pumila
NCU00154874	Castanea pumila var. pumila
NCU00153916	Castanea dentata
NCU00153914	Castanea dentata
NCU00153925	Castanea dentata

NCU00085296	Castanea dentata	Note absence of stellate trichomes on abaxial leaf surfaces. Note presence of long, appressed, simple trichomes on midribs. Note absence of trichomes on leaf margins.
NCU00085297	Castanea dentata	
NCU00085295	Castanea dentata	
NCU00085294	Castanea dentata	
NCU00085293	Castanea dentata	
NCU00085292	Castanea dentata	
NCU00085291	Castanea dentata	
NCU00085298	Castanea dentata	
NCU00153933	Castanea dentata	
NCU00153932	Castanea dentata	
NCU00153931	Castanea dentata	
NCU00153930	Castanea mollissima	See abundant stellate trichomes and tan twig color. There are also larger, fewer lenticels than would be seen in C. dentata.
NCU00153927	Castanea dentata	
NCU00153926	Castanea dentata	
NCU00153888	Castanea dentata	
NCU00153887	Castanea dentata	
NCU00153885	Castanea dentata	
NCU00153886	Castanea dentata	
NCU00153883	Castanea dentata	
NCU00153884	Castanea dentata	
NCU00153882	Castanea dentata	
NCU00153889	Castanea dentata	
NCU00153945	Castanea mollissima	
NCU00153944	Castanea mollissima	
NCU00153943	Castanea mollissima	
NCU00153942	Castanea mollissima	
NCU00153941	Castanea mollissima	
NCU00008706	Castanea mollissima	
NCU00153832	Castanea dentata	
NCU00153833	Castanea dentata	
NCU00153834	Castanea dentata	
NCU00153835	Castanea dentata	
NCU00153938	Castanea mollissima	
NCU00153830	Castanea dentata	
NCU00153829	Castanea dentata	
NCU00153841	Castanea dentata	
NCU00153855	Castanea dentata	
NCU00153854	Castanea dentata	

NCU00153853	Castanea dentata	
NCU00153852	Castanea dentata	
NCU00153851	Castanea mollissima	Note the dense covering of stellate trichomes on abaxial leaf surface; short, erect, simple trichomes on adaxial midrib; stellate trichomes on adaxial leaf surface adjacent to midrib; and stipules broader than stem with a flare at the base.
NCU00153850	Castanea dentata	
NCU00153848	Castanea dentata	
NCU00153846	Castanea dentata	
NCU00153844	Castanea dentata	
NCU00153842	Castanea dentata	
NCU00153843	Castanea dentata	
NCU00153819	Castanea pumila var. pumila	
NCU00153818	Castanea pumila var. pumila	
NCU00153924	Castanea dentata	
NCU00153923	Castanea dentata	
NCU00153920	Castanea dentata	
NCU00153922	Castanea dentata	
NCU00053292	Castanea pumila var. pumila	Twigs on this specimen are more slender than twigs we have observed in C. pumila var. ozarkensis.
NCU00153918	Castanea dentata	
NCU00053293	Castanea pumila var. ozarkensis	
NCU00153917	Castanea dentata	
NCU00053294	Castanea pumila	Leaves smaller than we have observed in C. pumila var. ozarkensis. Current season's twigs appear slender. Pistillate flowers absent, however.
NCU00053295	Castanea pumila	
NCU00053297	Castanea pumila	
NCU00053298	Castanea pumila	
NCU00053299	Castanea pumila	
NCU00053401	Castanea pumila	
NCU00053402	Castanea pumila	
NCU00053291	Castanea pumila	
NCU00153951	Castanea pumila var. pumila	
NCU00153950	Castanea pumila var. pumila	
NCU00153948	Castanea pumila var. pumila	
NCU00153952	Castanea pumila var. pumila	
NCU00153965	Castanea pumila var. pumila	
NCU00153963	Castanea pumila var. pumila	

NCU00153962	Castanea pumila var. pumila	
NCU00153961	Castanea pumila var. pumila	
NCU00153960	Castanea pumila var. pumila	
NCU00153959	Castanea pumila var. pumila	This specimen is similar to C. alabamensis in its absence of stellate trichomes and presence of ciliate leaf margins, but is different by having simple trichomes on the twigs.
NCU00153958	Castanea pumila var. pumila	
NCU00153957	Castanea pumila var. pumila	
NCU00153955	Castanea pumila var. pumila	
NCU00153954	Castanea pumila var. pumila	
NCU00153966	Castanea pumila var. pumila	
NCU00153953	Castanea pumila var. pumila	
NCU00153975	Castanea pumila var. pumila	
NCU00153974	Castanea pumila var. pumila	This specimen is similar to C. alabamensis in its absence of stellate trichomes and presence of ciliate leaf margins, but is different by having simple trichomes on the twigs. Leaf shape suggests that voucher may have been collected from shade leaves.
NCU00153973	Castanea pumila var. pumila	
NCU00153972	Castanea pumila var. pumila	This sheet may contain parts of two separate plants. See leaf color.
NCU00153971	Castanea pumila var. pumila	
NCU00153969	Castanea pumila var. pumila	
NCU00153968	Castanea pumila var. pumila	
NCU00153352	Castanea mollissima	
NCU00153351	Castanea henryi	
NCU00153355	Castanea seguinii	
NCU00153354	Castanea seguinii	
NCU00153353	Castanea mollissima	
NCU00013728	Castanea mollissima	
NCU00153366	Castanea dentata	
NCU00153364	Castanea dentata	
NCU00153363	Castanea dentata	
NCU00153362	Castanea dentata	
NCU00153361	Castanea dentata	
NCU00153360	Castanea dentata	
NCU00153358	Castanea dentata	
NCU00153356	Castanea dentata	
NCU00153367	Castanea dentata	
NCU00153373	Castanea pumila var. pumila	

NCU00153371	Castanea pumila var. pumila	
NCU00153370	Castanea pumila var. pumila	
NCU00153369	Castanea pumila var. pumila	
NCU00157730	Castanea mollissima	
NCU00154862	Castanea pumila var. pumila	
NCU00154014	Castanea pumila var. pumila	
NCU00154015	Castanea pumila var. pumila	
NCU00154013	Castanea pumila var. pumila	
NCU00154016	Castanea pumila var. pumila	
NCU00154016	Castanea pumila var. pumila	
NCU00154017	Castanea pumila var. pumila	
NCU00154019	Castanea pumila var. pumila	
NCU00154020	Castanea pumila var. pumila	
NCU00154861	Castanea pumila var. pumila	
NCU00154011	Castanea pumila var. pumila	
NCU00154001	Castanea pumila var. pumila	
NCU00154002	Castanea pumila var. pumila	
NCU00154004	Castanea pumila var. pumila	
NCU00154005	Castanea pumila var. pumila	
NCU00154006	Castanea pumila var. pumila	
NCU00154008	Castanea pumila var. pumila	
NCU00154008	Castanea pumila var. pumila	
NCU00154009	Castanea pumila var. pumila	
NCU00154010	Castanea pumila var. pumila	
NCU00154000	Castanea pumila var. pumila	
NCU00153991	Castanea pumila var. pumila	
NCU00153993	Castanea pumila var. pumila	
NCU00153995	Castanea pumila var. pumila	
NCU00153996	Castanea pumila var. pumila	
NCU00153997	Castanea pumila var. pumila	
NCU00153999	Castanea pumila var. pumila	
NCU00153990	Castanea pumila var. pumila	
NCU00153986	Castanea pumila var. pumila	
NCU00153987	Castanea pumila var. pumila	
NCU00153989	Castanea alabamensis	We observed only one stellate trichome in the axil of midrib and lateral vein on abaxial surface of leaf. Plant was sampled from Sumter Co., AL, a potential contact zone for different chinquapin varieties and may be a hybrid of C. alabamensis and C. p. var. pumila
NCU00153985	Castanea pumila var. pumila	
NCU00153977	Castanea pumila var. pumila	
NCU00153978	Castanea pumila var. pumila	

NCU00153980	Castanea pumila var. pumila	
NCU00153979	Castanea pumila var. pumila	
NCU00153982	Castanea pumila var. pumila	
NCU00153984	Castanea pumila var. pumila	
NCU00053435	Castanea pumila var. ozarkensis	
NCU00053446	Castanea pumila var. ozarkensis	
NCU00031434	Castanea pumila var. ozarkensis	
NCU00053444	Castanea pumila	
NCU00053441	Castanea pumila var. ozarkensis	
NCU00053442	Castanea pumila var. ozarkensis	
NCU00053439	Castanea pumila var. ozarkensis	
NCU00053440	Castanea pumila var. ozarkensis	
NCU00053447	Castanea pumila var. ozarkensis	
NCU00053452	Castanea pumila var. ozarkensis	
NCU00053451	Castanea pumila	
NCU00053450	Castanea pumila var. ozarkensis	
NCU00053449	Castanea pumila var. ozarkensis	
NCU00053453	Castanea pumila var. ozarkensis	
NCU00053460	Castanea pumila var. ozarkensis	
NCU00053459	Castanea pumila var. ozarkensis	
NCU00053457	Castanea pumila var. ozarkensis	
NCU00053418	Castanea pumila var. ozarkensis	
NCU00053456	Castanea pumila var. ozarkensis	
NCU00053461	Castanea pumila var. ozarkensis	
NCU00053463	Castanea pumila var. ozarkensis	
NCU00053462	Castanea pumila var. ozarkensis	
NCU00053436	Castanea pumila var. pumila	Note the small bur (less than 2.5 cm diameter, including spines) and short leaf lengths (10 cm or less).
NCU00053437	Castanea pumila	Mature bur diameter (2 cm), leaf lengths (11.5 cm in length, max.), and note on plant habit all suggest that plant could be either var. pumila or var. ozarkensis.
NCU00153150	Castanea dentata	
NCU00153148	Castanea dentata	
NCU00153169	Castanea dentata	
NCU00153166	Castanea dentata	
NCU00153165	Castanea dentata	
NCU00008701	Castanea dentata	
NCU00153164	Castanea dentata	
NCU00153162	Castanea dentata	
NCU00153161	Castanea dentata	
NCU00153170	Castanea dentata	
NCU00008702	Castanea dentata	

NCU00153177	Castanea pumila var. pumila	
NCU00031381	Castanea pumila var. pumila	
NCU00153059	Castanea crenata	Note the cordate leaf bases, mix of stellate and simple abaxial trichomes, slender twig, and oblong leaf shape.
NCU00116435	Castanea dentata	
NCU00153147	Castanea dentata	
NCU00153146	Castanea dentata	
NCU00153145	Castanea dentata	
NCU00113377	Castanea mollissima	Note cordate leaf bases, stipules wider than stem, crenate leaf margins.
NCU00154955	Castanea pumila var. pumila	
NCU00154953	Castanea pumila var. pumila	
NCU00154952	Castanea pumila var. pumila	
NCU00154949	Castanea pumila var. pumila	
NCU00154950	Castanea pumila var. pumila	
NCU00154947	Castanea pumila var. pumila	
NCU00154956	Castanea pumila var. pumila	
NCU00154966	Castanea pumila var. pumila	
NCU00154964	Castanea pumila var. pumila	
NCU00154962	Castanea pumila var. pumila	
NCU00154961	Castanea pumila var. pumila	
NCU00154960	Castanea pumila var. pumila	
NCU00154959	Castanea pumila var. pumila	
NCU00154957	Castanea pumila var. pumila	
NCU00154967	Castanea pumila var. pumila	
NCU00154973	Castanea pumila var. pumila	
NCU00154972	Castanea pumila var. pumila	
NCU00154971	Castanea pumila var. pumila	
NCU00154970	Castanea pumila var. pumila	
NCU00154969	Castanea pumila var. pumila	
NCU00154968	Castanea pumila var. pumila	
NCU00154974	Castanea pumila var. pumila	
NCU00155307	Castanea pumila var. pumila	
NCU00155306	Castanea pumila var. pumila	
NCU00155299	Castanea pumila var. pumila	
NCU00155298	Castanea pumila var. pumila	
NCU00155300	Castanea pumila var. pumila	
NCU00155301	Castanea pumila var. pumila	
NCU00155302	Castanea pumila var. pumila	
NCU00155217	Castanea pumila var. pumila	
NCU00155218	Castanea pumila var. pumila	
NCU00155213	Castanea pumila var. pumila	

NCU00155214	Castanea pumila var. pumila
NCU00155215	Castanea pumila var. pumila
NCU00155219	Castanea pumila var. pumila
NCU00155209	Castanea pumila var. pumila
NCU00155211	Castanea pumila var. pumila
NCU00155210	Castanea pumila var. pumila
NCU00155288	Castanea pumila var. pumila
NCU00155285	Castanea pumila var. pumila
NCU00155357	Castanea dentata
NCU00155283	Castanea pumila var. pumila
NCU00155355	Castanea dentata
NCU00155354	Castanea dentata
NCU00155349	Castanea pumila var. ozarkensis
NCU00155348	Castanea pumila var. ozarkensis
NCU00155352	Castanea pumila var. ozarkensis
NCU00155346	Castanea dentata
NCU00155104	Castanea pumila var. pumila
NCU00155103	Castanea pumila var. pumila
NCU00155102	Castanea pumila var. pumila
NCU00155101	Castanea pumila var. pumila
NCU00154979	Castanea pumila var. pumila
NCU00154980	Castanea pumila var. pumila
NCU00154977	Castanea pumila var. pumila
NCU00154976	Castanea pumila var. pumila
NCU00155105	Castanea pumila var. pumila
NCU00154975	Castanea pumila var. pumila
NCU00155114	Castanea pumila var. pumila
NCU00155113	Castanea pumila var. pumila
NCU00155112	Castanea pumila var. pumila
NCU00155111	Castanea pumila var. pumila
NCU00155109	Castanea pumila var. pumila
NCU00155108	Castanea pumila var. pumila
NCU00155107	Castanea pumila var. pumila
NCU00155106	Castanea pumila var. pumila
NCU00155115	Castanea pumila var. pumila
NCU00155120	Castanea pumila var. pumila
NCU00155119	Castanea pumila var. pumila
NCU00155118	Castanea pumila var. pumila
NCU00155117	Castanea pumila var. pumila
NCU00155116	Castanea pumila var. pumila
NCU00157719	Castanea pumila var. pumila
NCU00157720	Castanea pumila var. pumila
NCU00157721	Castanea pumila var. pumila
NCU00157723	Castanea pumila var. pumila
NCU00157724	Castanea pumila var. pumila

NCU00157725	Castanea pumila var. pumila
NCU00157726	Castanea pumila var. pumila
NCU00155289	Castanea pumila var. pumila
NCU00155297	Castanea pumila var. pumila
NCU00155296	Castanea pumila var. pumila
NCU00155295	Castanea pumila var. pumila
NCU00155294	Castanea pumila var. pumila
NCU00155293	Castanea pumila var. pumila
NCU00155292	Castanea pumila var. pumila
NCU00155291	Castanea pumila var. pumila
NCU00155220	Castanea pumila var. pumila
NCU00155281	Castanea pumila var. pumila
NCU00155284	Castanea pumila var. pumila
NCU00155318	Castanea pumila var. pumila
NCU00155317	Castanea pumila var. pumila
NCU00155316	Castanea pumila var. pumila
NCU00092135	Castanea pumila var. pumila
NCU00155320	Castanea pumila var. pumila
NCU00155310	Castanea pumila var. pumila
NCU00155309	Castanea pumila var. pumila
NCU00155308	Castanea pumila var. pumila
NCU00155315	Castanea pumila var. pumila
NCU00089777	Castanea pumila var. pumila
NCU00155311	Castanea pumila var. pumila
NCU00155314	Castanea pumila var. pumila
NCU00155304	Castanea pumila var. pumila
NCU00155303	Castanea pumila var. pumila
NCU00135543	Castanea pumila var. pumila
NCU00155319	Castanea pumila var. pumila
NCU00155321	Castanea dentata
NCU00135547	Castanea crenata
NCU00155322	Castanea dentata
NCU00155170	Castanea pumila var. pumila
NCU00155197	Castanea pumila var. pumila
NCU00155196	Castanea pumila var. pumila
NCU00155195	Castanea pumila var. pumila
NCU00155194	Castanea pumila var. pumila
NCU00155199	Castanea pumila var. pumila
NCU00155178	Castanea pumila var. pumila
NCU00155177	Castanea pumila var. pumila
NCU00155181	Castanea pumila var. pumila
NCU00155358	Castanea pumila var. pumila
NCU00155361	Castanea pumila var. pumila
NCU00155365	Castanea pumila var. pumila
NCU00155364	Castanea pumila var. pumila

NCU00155363	Castanea pumila var. pumila
NCU00155370	Castanea pumila var. pumila
NCU00155359	Castanea pumila var. pumila
NCU00155381	Castanea crenata
NCU00155380	Castanea crenata
NCU00155379	Castanea crenata
NCU00155378	Castanea crenata
NCU00155377	Castanea crenata
NCU00155376	Castanea crenata
NCU00155323	Castanea dentata
NCU00155330	Castanea dentata
NCU00155325	Castanea dentata
NCU00155327	Castanea dentata
NCU00155324	Castanea dentata
NCU00155334	Castanea dentata
NCU00155335	Castanea dentata
NCU00155336	Castanea dentata
NCU00155329	Castanea dentata
NCU00155326	Castanea dentata
NCU00155347	Castanea dentata
NCU00155332	Castanea dentata
NCU00155331	Castanea dentata
NCU00155338	Castanea dentata
NCU00155333	Castanea dentata
NCU00155339	Castanea dentata
NCU00155342	Castanea dentata
NCU00155341	Castanea dentata
NCU00155340	Castanea dentata
NCU00155345	Castanea dentata
NCU00155344	Castanea dentata
NCU00155343	Castanea dentata
NCU00155353	Castanea pumila var. ozarkensis
NCU00155350	Castanea pumila var. ozarkensis
NCU00154917	Castanea pumila var. pumila
NCU00154916	Castanea pumila var. pumila
NCU00154915	Castanea pumila var. pumila
NCU00154914	Castanea pumila var. pumila
NCU00135545	Castanea pumila var. pumila
NCU00154934	Castanea pumila var. pumila
NCU00154933	Castanea pumila var. pumila
NCU00154931	Castanea pumila var. pumila
NCU00154930	Castanea pumila var. pumila
NCU00154929	Castanea pumila var. pumila
NCU00154928	Castanea pumila var. pumila
NCU00154927	Castanea pumila var. pumila

NCU00154926	Castanea pumila var. pumila	
NCU00154925	Castanea pumila var. pumila	
NCU00135546	Castanea pumila var. pumila	
NCU00154924	Castanea pumila var. pumila	
NCU00154935	Castanea pumila var. pumila	
NCU00154945	Castanea pumila var. pumila	
NCU00154943	Castanea pumila var. pumila	
NCU00154942	Castanea pumila var. pumila	
NCU00154941	Castanea pumila var. pumila	
NCU00154940	Castanea pumila var. pumila	
NCU00154939	Castanea pumila var. pumila	
NCU00154939	Castanea pumila var. pumila	
NCU00154938	Castanea pumila var. pumila	
NCU00154936	Castanea pumila var. pumila	
NCU00154946	Castanea pumila var. pumila	
NCU00154919	Castanea pumila var. pumila	
NCU00154920	Castanea pumila var. pumila	
NCU00154922	Castanea pumila var. pumila	
NCU00154923	Castanea pumila var. pumila	
NCU00155162	Castanea pumila var. pumila	
NCU00155161	Castanea pumila var. pumila	
NCU00155160	Castanea pumila var. pumila	
NCU00155159	Castanea pumila var. pumila	
NCU00155158	Castanea pumila var. pumila	
NCU00155157	Castanea pumila var. pumila	
NCU00155156	Castanea pumila var. pumila	
NCU00155134	Castanea pumila var. pumila	
NCU00155121	Castanea pumila var. pumila	
NCU00155127	Castanea pumila var. pumila	
NCU00155123	Castanea pumila var. pumila	
NCU00155124	Castanea pumila var. pumila	
NCU00155125	Castanea pumila var. pumila	
NCU00155126	Castanea pumila var. pumila	
NCU00155154	Castanea pumila var. pumila	
NCU00155163	Castanea pumila var. pumila	
NCU00155144	Castanea pumila var. pumila	
NCU00155143	Castanea pumila var. pumila	
NCU00155153	Castanea pumila var. pumila	
NCU00155146	Castanea pumila var. pumila	
NCU00155374	Castanea sativa	Note the long petioles, stout twigs, small lenticels, elongate leaves, and large buds.
NCU00155372	Castanea mollissima	
NCU00155373	Castanea sativa	
NCU00155368	Castanea pumila var. pumila	

NCU00155367	Castanea pumila var. pumila	
NCU00155366	Castanea pumila var. pumila	
NCU00155371	Castanea pumila var. pumila	
NCU00155392	Castanea seguinii	Burs at multiple nodes along the stem are evidence of the continuous flowering typical of Castanea seguinii.
NCU00155396	Castanea sativa	
NCU00155382	Castanea crenata	
NCU00155394	Castanea sativa	We think this is from Manali, Himachal Pradesh, India
NCU00155393	Castanea sativa	
NCU00155384	Castanea henryi	
NCU00155383	Castanea seguinii	Burs at multiple nodes along the stem are evidence of the continuous flowering typical of Castanea seguinii.
NCU00155397	Castanea sativa	
NCU00155391	Castanea sativa	
NCU00155388	Castanea sativa	
NCU00155390	Castanea sativa	
NCU00155387	Castanea sativa	
NCU00155386	Castanea sativa	
NCU00155385	Castanea henryi	
NCU00155180	Castanea pumila var. pumila	
NCU00155182	Castanea pumila var. pumila	
NCU00155185	Castanea pumila var. pumila	
NCU00155190	Castanea pumila var. pumila	
NCU00155193	Castanea pumila var. pumila	
NCU00155192	Castanea pumila var. pumila	
NCU00155202	Castanea pumila var. pumila	
NCU00155201	Castanea pumila var. pumila	
NCU00155200	Castanea pumila var. pumila	
NCU00155208	Castanea pumila var. pumila	
NCU00155207	Castanea pumila var. pumila	
NCU00155206	Castanea pumila var. pumila	
NCU00155205	Castanea pumila var. pumila	
NCU00155204	Castanea pumila var. pumila	
NCU00155171	Castanea pumila var. pumila	
NCU00155176	Castanea pumila var. pumila	
NCU00155175	Castanea pumila var. pumila	
NCU00155174	Castanea pumila var. pumila	
NCU00146417	Castanea pumila var. pumila	
NCU00027810	Castanea pumila var. pumila	
NCU00155168	Castanea pumila var. pumila	
NCU00155167	Castanea pumila var. pumila	

NCU00155166	Castanea pumila var. pumila
NCU00155165	Castanea pumila var. pumila
NCU00155164	Castanea pumila var. pumila
NCU00155172	Castanea pumila var. pumila
NCU00155189	Castanea pumila var. pumila
NCU00155188	Castanea pumila var. pumila
NCU00155187	Castanea pumila var. pumila
NCU00155145	Castanea pumila var. pumila
NCU00155150	Castanea pumila var. pumila
NCU00155148	Castanea pumila var. pumila
NCU00155149	Castanea pumila var. pumila
NCU00155152	Castanea pumila var. pumila
NCU00155135	Castanea pumila var. pumila
NCU00155137	Castanea pumila var. pumila
NCU00155136	Castanea pumila var. pumila
NCU00155141	Castanea pumila var. pumila
NCU00155140	Castanea pumila var. pumila
NCU00155139	Castanea pumila var. pumila
NCU00155131	Castanea pumila var. pumila
NCU00155130	Castanea pumila var. pumila
NCU00155133	Castanea pumila var. pumila
NCU00155128	Castanea pumila var. pumila
NCU00155389	Castanea sativa
NCU00155129	Castanea pumila var. pumila

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821 Appendix S4. Illumina sequencing summary. Samples highlighted in yellow were not used for genotyping due to a low number
822 (<100,000) of total reads. Samples highlighted in blue are technical replicates of *C. mollissima* and *C. crenata* cultivars that were not
823 genotyped because they produced the fewest retained reads of two technical replicates per cultivar.
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well ID	DNA tube name	Species	barcode	GBS ID	SRA filename	Total reads	NoRadTag	LowQuality	Retained reads	depth of coverage
A1	A1	C. p. var. pumila	TGACGCCA	1	cv_1	2369236	6978	3665	2358593	27
B1	CEF1	C. dentata	CAGATA	2	cv_2	3289014	12714	5165	3271135	38
C1	BF6	C. dentata	GAAGTG	3	cv_3	4022872	22782	6187	3993903	44
D1	669	C. dentata	TAGCGGAT	4	cv_4	464578	2178	663	461737	19
E1	A2	C. p. var. pumila	TATTCGCAT	5	cv_5	7041114	19157	10723	7011234	59
F1	CEF2	C. dentata	ATAGAT	6	cv_6	4644790	16713	7335	4620742	49
G1	CR1	C. dentata	CCGAACA	7	cv_7	6328058	14423	9999	6303636	61
H1	710	C. dentata	GGAAGACAT	8	cv_8	183218	3942	253	179023	17
A2	A3	C. p. var. pumila	GGCTTA	9	cv_9	5596302	14166	8700	5573436	48
B2	CEF3	C. dentata	AACGCACATT	10	cv_10	2745442	11419	4155	2729868	35
C2	CR2	C. dentata	GAGCGACAT	11	cv_11	5654802	12022	8831	5633949	56
D2	675	C. dentata	CCTTGCCATT	12	cv_12	1292256	6238	1907	1284111	27
E2	A4	C. p. var. pumila	GGTATA	13	cv_13	10175602	26508	15541	10133553	73
F2	CEF4	C. dentata	TCTTGG	14	cv_14	4731848	17074	7141	4707633	48
G2	CR3	C. p. var. pumila	GGTGT	15	cv_15	4269384	38568	6651	4224165	39
H2	KYADA1	C. dentata	GGATA	16	cv_16	1885530	20109	2789	1862632	28
A3	A5	C. p. var. pumila	CTAAGCA	17	cv_17	7324422	22242	11549	7290631	59
B3	G1	C. dentata	ATTAT	18	cv_18	6351844	28447	9745	6313652	54
C3	CR4	C. p. var. pumila	GCGCTCA	19	cv_19	8560246	17846	13200	8529200	66
D3	G2	C. p. var. pumila	ACTGCGAT	20	cv_20	8004334	15649	12144	7976541	67
E3	CR5	C. p. var. pumila	TTCGTT	21	cv_21	8540658	22051	13177	8505430	66

F3	412	C. alabamensis	ATATAA	22	cv_22	652780	8800	1023	642957	23
G3	A7	C. p. var. pumila	TGGCAACAGA	23	cv_23	2983238	7788	4758	2970692	36
H3	G3	C. p. var. pumila	CTCGTCG	24	cv_24	10297574	19305	15794	10262475	75
A4	CR6	C. p. var. pumila	GCCTACCT	25	cv_25	4413006	10728	6775	4395503	52
B4	735	C. p. var. pumila	CACCA	26	cv_26	184696	33774	204	150718	17
C4	645	C. dentata	AATTAG	27	cv_27	1585046	12563	2406	1570077	28
D4	A8	C. p. var. pumila	GGAACGA	28	cv_28	5037102	13672	7953	5015477	50
E4	BF1	C. p. var. pumila	ACAACT	29	cv_29	7777390	45883	11685	7719822	65
F4	CR7	C. p. var. pumila	ACTGCT	30	cv_30	8891970	39077	13829	8839064	72
G4	646	C. dentata	CGTGGACAGT	31	cv_31	2191268	10454	3459	2177355	33
H4	A9	C. p. var. pumila	TGGCACAGA	32	cv_32	8534758	18418	13267	8503073	74
A5	BF2	C. p. var. pumila	TGCTT	33	cv_33	8839182	28483	13472	8797227	80
B5	671	C. dentata	GCAAGCCAT	34	cv_34	1394902	5541	2242	1387119	28
C5	647	C. p. var. pumila	CGCACCAATT	35	cv_35	320568	7879	467	312222	17
D5	A10	C. p. var. pumila	CTCGCGG	36	cv_36	6173688	17069	9276	6147343	56
E5	BF3	C. p. var. pumila	AACTGG	37	cv_37	12554228	38341	19181	12496706	66
F5	420	C. alabamensis	ATGAGCAA	38	cv_38	387424	3119	604	383701	19
G5	672	C. dentata	CTTGA	39	cv_39	2862578	12325	4294	2845959	43
H5	648	C. p. var. pumila	GCGTCCT	40	cv_40	2444192	18776	3696	2421720	35
A6	WB1	C. p. var. pumila	ACCAGGA	41	cv_41	3358210	12294	5071	3340845	38
B6	BF4	C. p. var. pumila	CCACTCA	42	cv_42	3518126	8355	5469	3504302	39
C6	649	C. p. var. pumila	TCACGGAAG	43	cv_43	485544	11104	754	473686	18
D6	WB2	C. p. var. pumila	TATCA	44	cv_44	4396572	70444	6770	4319358	43

E6	BF5	C. dentata	TAGCCAA	45	cv_45	5960384	54571	9082	5896731	54
F6	650	C. p. var. pumila	ATATCGCCA	46	cv_46	818876	5146	1210	812520	21
G6	651	C. p. var. pumila	CTCTA	47	cv_47	1798116	11713	2695	1783708	29
H6	652	C. p. var. pumila	GGTGCACATT	48	cv_48	625570	9729	942	614899	19
A7	653	C. p. var. pumila	CTCTCGCAT	49	cv_49	8848	675	7	8166	14
B7	507	C. dentata	CAGAGGT	50	cv_50	3613706	15383	5525	3592798	48
C7	513	C. dentata	GCGTACAAT	51	cv_51	2153114	5350	3402	2144362	34
D7	514	C. dentata	ACGCGCG	52	cv_52	2873730	11423	4383	2857924	39
E7	124	C. p. var. pumila	GTCGCCT	53	cv_53	3416422	10973	5181	3400268	46
F7	525	C. p. var. pumila	AATAACCAA	54	cv_54	3719032	15301	5694	3698037	48
G7	126	C. p. var. pumila	AATGAACGA	55	cv_55	3127094	9484	4951	3112659	43
H7	527	C. p. var. pumila	CGTCGCCACT	56	cv_56	1746	527	2	1217	10
A8	528	C. p. var. pumila	ATGGCAA	57	cv_57	1723710	7468	2780	1713462	31
B8	529	C. p. var. pumila	GAAGCA	58	cv_58	157746	24598	216	132932	16
C8	531	C. dentata	AACGTGCCT	59	cv_59	159562	1636	234	157692	16
D8	532	C. p. var. pumila	CCTCG	60	cv_60	2484538	64197	3632	2416709	35
E8	535	C. p. var. pumila	CTCAT	61	cv_61	4260888	60936	6472	4193480	49
F8	149	C. p. var. pumila	ACGGTACT	62	cv_62	2042698	6087	3181	2033430	32
G8	694	C. p. var. ozarkensis	GCGCCG	63	cv_63	5953966	73081	8866	5872019	57
H8	705	C. p. var. ozarkensis	CAAGT	64	cv_64	1454382	74121	2129	1378132	28
A9	673	C. dentata	TCCGAG	65	cv_65	3028130	10212	4608	3013310	42
B9	674	C. dentata	TAGATGA	66	cv_66	2085178	11179	3206	2070793	32
C9	H1	C. alabamensis	TGGCCAG	67	cv_67	5726678	20229	8575	5697874	53
D9	H2	C. dentata	GCACGAT	68	cv_68	4051206	20702	6386	4024118	44

E9	H3	C. dentata	TTGCTG	69	cv_69	2144734	9074	3271	2132389	34
F9	H4	C. dentata	CGCAACCAGT	70	cv_70	2846048	8484	4285	2833279	40
G9	F1	C. dentata	TCACTG	71	cv_71	5066636	23183	7716	5035737	56
H9	F5	C. dentata	ACAGT	72	cv_72	7189348	49805	11066	7128477	72
A10	AG1	C. alabamensis	GGAGTCAAG	73	cv_73	4533302	10877	7160	4515265	48
B10	AG2	C. dentata	TGAAT	74	cv_74	4696900	88410	7177	4601313	49
C10	AG3	C. alabamensis	CATAT	75	cv_75	6806594	480687	9691	6316216	57
D10	AG6	C. alabamensis	GTGACACAT	76	cv_76	8693168	20173	13627	8659368	63
E10	AG7	C. alabamensis	TATGT	77	cv_77	9244322	26806	14408	9203108	80
F10	L14	C. dentata	CAGTGCCATT	78	cv_78	6036182	13627	9387	6013168	60
G10	L19	C. dentata	ACAACCAACT	79	cv_79	5539000	22750	8480	5507770	57
H10	SS2	C. p. var. pumila	TGCAGA	80	cv_80	4056690	22128	6154	4028408	51
A11	SS3	C. p. var. pumila	CATCTGCCG	81	cv_81	708390	2772	1095	704523	21
B11	SS4	C. p. var. pumila	GGACAG	82	cv_82	5588968	21131	8641	5559196	63
C11	LIN1	C. dentata	ATCTGT	83	cv_83	7461978	19509	11699	7430770	71
D11	F3	C. dentata	AAGACGCT	84	cv_84	3353160	10272	5270	3337618	37
E11	AG4	C. alabamensis	GAATGCAATA	85	cv_85	6908794	24796	10926	6873072	60
F11	H9	C. alabamensis	TAGCAG	86	cv_86	491558	27366	698	463494	13
G11	AG5	C. alabamensis	ATCCG	87	cv_87	118870	25961	139	92770	13
H11	L10	C. dentata	CTTAG	88	cv_88	3987598	13931	6114	3967553	47
A12	H6	C. alabamensis	TTATTACAT	89	cv_89	1985086	8208	2995	1973883	31
B12	CH21	C. alabamensis	GCCAACAAGA	90	cv_90	514854	2455	743	511656	19
C12	CH22	C. dentata	TGCCGCAT	91	cv_91	6034174	199398	9196	5825580	56
D12	CH23	C. alabamensis	CGTGTC	92	cv_92	1193850	9430	1709	1182711	24
E12	CH24	C. alabamensis	CAACCACACA	93	cv_93	2972860	10879	4486	2957495	37
F12	CH25	C. alabamensis	GCTCCGA	94	cv_94	3299294	7773	4888	3286633	39
G12	CH26	C. alabamensis	TCAGAGAT	95	cv_95	2696792	45481	4293	2647018	37

H12	CH27	C. p. var. pumila	CGTTCA	96	cv_96	3903682	19055	6025	3878602	40
A11		C. dentata AD98	CATCTGCCG	97	cv_97	1858140	8912	5260	1843968	21
B11		C. molissima Mahogany	GGACAG	98	cv_98	13652574	75331	37598	13539645	54
C11		C. mollissima Mahogany	ATCTGT	99	cv_99	15041542	66372	41249	14933921	#N/A
D11		C. molissima Nanking	AAGACGCT	100	cv_100	11359132	50407	31615	11277110	56
E11		C. mollissima Nanking	GAATGCAATA	101	cv_101	5894876	46111	16604	5832161	#N/A
F11		C. dentata NC DOT	TAGCAG	102	cv_102	11053870	2042580	20919	8990371	56
G11		C. dentata Cranberry	ATCCG	103	cv_103	6282890	63302	16997	6202591	34
H11		C. crenata Fort Defiance	CTTAG	104	cv_104	15531680	1419895	36228	14075557	56
A12		C. crenata Fort Defiance	TTATTACAT	105	cv_105	6862274	33527	19145	6809602	#N/A
B12		C. crenata Morrow Mountain	GCCAACAAGA	106	cv_106	3412182	18856	9703	3383623	#N/A
C12		C. crenata Morrow Mountain	TGCCGCAT	107	cv-107	4919620	149206	13435	4756979	41
D12		C. crenata Rita #1	CGTGTCA	108	cv_108	4964916	34518	13384	4917014	43
E12		C. dentata Haun (PennState)	CAACCACACA	109	cv_109	4530	369	12	4149	#N/A
F12		C. crenata Rita #2	GCTCCGA	110	cv_110	7019724	29611	19432	6970681	58
Plants excluded from genotyping due to <100000 reads (highlighted in yellow)					49, 56, 109					
Lower quality replicates of Asian cultivars excluded from genotyping (highlighted in blue)					99,101,105,106					
Total reads for 106 plants including all technical replicates of E. Asian Castanea cultivars					489935694					
Total reads for 106 plants (only counting only the best replicate of each E. Asian cultivar used for genotyping)					458724820					
Total individual plants sequenced					106					

Total retained reads for 106 plants	451514582					
total samples retained	103					
Percentage reads retained	98.4281997					
Average clean reads per retained individual	4383505.34					
average coverage depth per retained individual (103 tot)	43.688835					

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Appendix S5. Summary of datasets exported by Stacks 1.45.

Dataset name	File name on GitHub	Taxa included	Total individuals	Total SNPs
NAC+EAC	batch_1.vcf	<i>C. alabamensis</i> , <i>C. dentata</i> , <i>C. pumila</i> var. <i>pumila</i> , <i>C. pumila</i> var. <i>ozarkensis</i> , <i>C. crenata</i> , <i>C. mollissima</i>	103	103,616
NAC	out98copy.vcf	<i>C. alabamensis</i> , <i>C. dentata</i> , <i>C. pumila</i> var. <i>pumila</i> , <i>C. pumila</i> var. <i>ozarkensis</i>	96	190,656
NAC+EAC583	allsamples_popcodes.txt	<i>C. alabamensis</i> , <i>C. dentata</i> , <i>C. pumila</i> var. <i>pumila</i> , <i>C. pumila</i> var. <i>ozarkensis</i> , <i>C. crenata</i> , <i>C. mollissima</i>	103	583

843 Appendix S6. Maximum likelihood phylogenetic tree showing identities of individual samples at
 844 branch tips. Samples representing *C. alabamensis* are highlighted in red. Numbers at nodes
 845 indicate bootstrap support. Scale bar is proportional to 0.07 substitutions/site.

846 0.07

Appendix S7. Weir and Cockerham's *Fst* between taxa pairs, summary of STRUCTURE analyses, and list of principal components.

Table S7a. Weir and Cockerham's *Fst* output from SNPrelate Output: "Fst" = weighted *Fst* estimate

	mollissima	crenata	dentata	pumila	ozarkensis	alabamensis
mollissima		0.5271483	0.6433073	0.6378966	0.7069509	0.6633516
crenata	0.5271483		0.6409639	0.6345136	0.6765049	0.6569612
dentata	0.6433073	0.6409639		0.3846462	0.4182757	0.3985978
pumila	0.6378966	0.6345136	0.3846462		0.1668146	0.1078042
ozarkensis	0.7069509	0.6765049	0.4182757	0.1668146		0.204663
alabamensis	0.6633516	0.6569612	0.3985978	0.1078042	0.204663	

Table S7b. Output of summary statistics from *Structure* analysis of *C. dentata*, *C. pumila sensu lato*, *C. mollissima*, and *C. crenata*. Statistics were calculated using the program STRUCTURE HARVESTER (Earl and vonHoldt 2012). The number of clusters present in the dataset ($K = 3$) was determined using the method of Evanno et al. (2005). $K = 3$ and associated ΔK value are highlighted in yellow.

K	Reps	Mean LnP(K)	Stdev LnP(K)	Ln'(K)	Ln''(K)	Delta K
1	10	-16771.91	1.349444	—	—	—
2	10	-13981.87	288.972021	2790.04	887.69	3.071889
3	10	-12079.52	15.195599	1902.35	1906.02	125.4324
4	10	-12083.19	216.654187	-3.67	974.83	4.499475
5	10	-13061.69	2928.160213	-978.5	3100.42	1.058829
6	10	-17140.61	8393.275267	-4078.92	6007.02	0.715694
7	10	-15212.51	5278.254233	1928.1	3060.45	0.579822
8	10	-16344.86	7472.795325	-1132.35	717.02	0.095951
9	10	-16760.19	9230.403714	-415.33	—	—

Table S7c. Output of summary statistics from *Structure* analysis of chinquapin samples.

Statistics were calculated using the program STRUCTURE HARVESTER (Earl and vonHoldt 2012). The number of clusters present in the dataset ($K = 2$) was determined using the method of Evanno et al. (2005). $K = 2$ and associated ΔK value are highlighted in yellow.

K	Reps	Mean LnP(K)	Stdev LnP(K)	Ln'(K)	Ln''(K)	Delta K
1	10	-6426.61	0.479467	—	—	—
2	10	-6041.17	2.833941	385.44	296.32	104.5611
3	10	-5952.05	137.78048	89.12	31.65	0.229713
4	10	-5831.28	39.063219	120.77	449.57	11.50878
5	10	-6160.08	1208.25738	-328.8	644.53	0.533438
6	10	-5844.35	182.341421	315.73	265.61	1.456663
7	10	-5794.23	223.654088	50.12	222.98	0.996986
8	10	-5967.09	647.43212	-172.86	301.03	0.46496
9	10	-5838.92	294.650689	128.17	423.32	1.436684
10	10	-6134.07	1064.66472	-295.15	525.14	0.493244
11	10	-5904.08	372.702293	229.99	285.78	0.766778
12	10	-5959.87	336.013763	-55.79	154.31	0.459237
13	10	-5861.35	275.391916	98.52	128.35	0.466063
14	10	-5891.18	306.478028	-29.83	93.25	0.304263
15	10	-6014.26	285.24903	-123.08	286.41	1.00407
16	10	-5850.93	306.721405	163.33	—	—

Table S7d. Proportions of total variance explained by the 32 principal components identified from principal components analysis of North American *Castanea* samples. The 32 principal components combined explain 56.5% of the total variance.

Principal components	Proportion variance
PC1	0.10957356
PC2	0.02748213
PC3	0.02287274
PC4	0.0209522
PC5	0.02088753
PC6	0.01773292
PC7	0.01741902
PC8	0.01648643
PC9	0.01636512
PC10	0.01508126
PC11	0.01485904
PC12	0.01454827
PC13	0.01430424
PC14	0.01416125
PC15	0.01406701
PC16	0.01339484
PC17	0.01296989
PC18	0.01289406
PC19	0.0127412
PC20	0.01270266
PC21	0.01257472
PC22	0.01237297
PC23	0.01229619
PC24	0.01221846
PC25	0.01204935
PC26	0.01201129
PC27	0.01188767
PC28	0.01183929
PC29	0.01173212
PC30	0.0116869
PC31	0.01160848
PC32	0.01147692