



US00PP14839P3

(12) **United States Plant Patent**
Beineke

(10) **Patent No.: US PP14,839 P3**
(45) **Date of Patent: Jun. 1, 2004**

- (54) **BLACK WALNUT TREE NAMED ‘BEINEKE 10’**
- (50) Latin Name: *Juglans nigra L.*
Varietal Denomination: **‘Beineke 10’**
- (75) Inventor: **Walter Beineke**, West Lafayette, IN (US)
- (73) Assignee: **American Forestry Technologies, Inc.**, WestPoint, IN (US)
- (*) Notice: Subject to any disclaimer, the term of this patent is extended or adjusted under 35 U.S.C. 154(b) by 58 days.
- (21) Appl. No.: **10/141,063**
- (22) Filed: **May 8, 2002**
- (65) **Prior Publication Data**
US 2003/0213032 P1 Nov. 13, 2003
- (51) **Int. Cl.⁷** **A01H 5/00**
- (52) **U.S. Cl.** **Plt./154**
- (58) **Field of Search** **Plt./154**

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(57) **ABSTRACT**

A new and distinct cultivar of black walnut tree (*Juglans nigra L.*) which is distinctly characterized by extremely rapid growth rate, fairly strong central stem tendency, thereby producing good timber qualities. The new variety has poor nut bearing qualities. Nut crops are erratic. Nut-bearing begins late in the life of the tree. This new variety of black walnut tree was discovered by the applicant near West Lafayette, Ind. in a black walnut planting of seedling progeny from aa previously selected tree for outstanding timber producing potential. This selection has been designated as BW 504, a seedling progeny of BW 36 in records maintained by the applicant on the performance of the selection and grafts made from the selection and will be known henceforth as Beineke 10.

3 Drawing Sheets

Latin name of the genus and species: *Juglans nigra L.*

BACKGROUND OF THE INVENTION

This new variety of black walnut tree was discovered by the applicant near West Lafayette, Ind. in a black walnut planting of seedling progeny from previously selected trees for outstanding timber producing potential. This selection has been designated as BW 504, a seedling progeny of BW 36 (unpatented) in records maintained by the applicant on the performance of the selection and grafts made from the selection and will be known henceforth as ‘Beineke 10.’ The male parent is unknown, as is generally the case with black walnut trees (Beineke, 1989).

SUMMARY OF THE INVENTION

A new and distinct cultivar of black walnut tree (*Juglans nigra L.*) which is distinctly characterized by extremely

rapid growth rate, fairly strong central stem tendency, thereby producing good timber qualities, the trait of commercial interest. Beineke 10 was 22 years old when described at a location near West Lafayette, Ind.

After the original clone was selected, and assigned an identity number of BW 504 the aforesaid tree was reproduced by collecting scions from it and grafting these onto common black walnut rootstocks at American Forestry Technologies, Inc., West Point Ind. These asexual reproductions ran true to the originally discovered tree and to each other in all respects. However, direct comparisons could not be made between BW36 and Beineke 10 because no BW36 trees were planted at this site.

Color values used were from the Munsell Color Chart for Plant Tissues. However, color is too dependent on weather conditions and fertilization to be consistent or distinctive.

For example, leaves can be made a deeper green by applying nitrogen. Walnut tree leaves turn yellow as the season progresses, especially if there is a lack of rainfall. As black walnut meats dry, they become darker. Simply being on the ground for a week causes the outer shell to darken. Bark color involves many shades of gray through brown and black.

Beineke 6 is hardy in USDA zones 4, 5, 6, 7, and 8.

BRIEF DESCRIPTION OF THE DRAWINGS

FIG. 1 is a photograph showing the timber form of 'Beineke 10.'

FIG. 2 is a photograph showing the leaves of 'Beineke 10.'

FIG. 3 is a photograph showing the nuts of 'Beineke 10.'

BOTANICAL DESCRIPTION OF THE PLANTS

The botanical details of this new and distinct variety of walnut tree are as follows:

Tree:

Size.—Large, 68 ft. at 22 years, crown diameter of 22 ft.

Vigor.—Vigorous.

Growth rate.—Very rapid, 22% larger in diameter than the average of parental Purdue 1 (U.S. Plant Pat. No. 4,543) grafts, planted the same year on the same land. Diameter growth rate (at 4½ feet above the ground) averages 0.555 inches per year over 22 years 12.2 inches.

Form.—Good timber form, not as good as Purdue 1 (U.S. Plant Pat. No. 4,543),—34% poorer than average of the entire planting. Stem form was obtained by subjectively rating the straightness of the main stem on a scale of 1 to 5 with 1 representing a perfectly straight stem; 2, slight crook or deviation of the central stem; 3, about average straightness; 4, several severe crooks or a single fork; and 5, a very crooked, forked and/or leaning central stem. Beineke 10 averages 2. It has the same straightness as the parent tree, BW 36. The trees of the present invention are grown in plantations, not open fields (not natural stands). In plantations, trees are upright and have no distinctive or characteristic crown shape because all branches are seeking to grow upwards.

Branches: Diameter depends on age and size of tree, varies from ½" to 12", bark color varies from grays to browns.

Leaves:

Compound leaves.—Size — Much shorter than average; average length — 10.38".

Leaflets.—Size — Smaller than average; average length — 3.70", average width — 2.85", average number of leaflets — 14.0 — lanceolate; acutely pointed, rounded base. Petioles — short.

Thickness.—Thin; Texture — smooth; Margin — serrated; Color — Topside — dark green (2.5 G 4/4 on the Munsell Color Chart for Plant Tissues); Under-side — light green. (5 GY 5/4 on the Munsell Color Chart for Plant Tissues).

Anthracnose resistance.—Good.

Nut:

Size.—Small; average length — 1.27"; average diameter in suture plane — 1.10", average diameter cheek to cheek — 1.33".

Uniformity of size.—Not much variation.

Form.—Rounded, flattened in suture plane. See FIG. 3.

Blossom end.—Rounded.

Basal end.—Slightly pointed to rounded.

Thickness of shell.—Thick.

Ridges.—Rounded off; not sharp.

Color.—Mottled, 5 YR 3/2 and 2/5 YR 3/4 on the Munsell Color Chart for Plant Tissues.

Flowering habit:

Age at which trees start producing catkins.—Late (about 6–7 years).

Number of catkins produced.—Average.

Age at which tree starts producing pistillate flowers.—Late (about 6–7 years).

Number of pistillate flowers produced by young trees.—Few.

Number of pistillate flowers produced by mature trees.—Few.

Lateral shoots producing pistillate flowers.—None.

Number of pistillate flowers per inflorescence.—2 to 4.

Flowering season: Flowers typically in May in Indiana.

There are probably 1—million pollen per catkin. Female flowers are about ¼" long and grow to two "pollen pick up points" which subsequently break apart. Pollen exits as "dust" which is not feasible to quantitate.

Nut crop:

Bearing.—Erratic.

Productivity.—Low.

Ripening period.—Late.

Evenness of maturity (period between time when first and last nuts are ready for harvest).—Even.

Quality.—Good.

Distribution of nuts on tree.—Throughout.

GENETIC METHOD OF IDENTIFICATION

DNA "fingerprint" for Identification of 'Beineke 10.'

DNA was isolated from the leaves of Beineke 10. For purposes of DNA fingerprinting, nine highly polymorphic loci from a suite of microsatellites developed by Woeste et al. (2002) were chosen. Microsatellites sizes were checked against previously published standards and verified by a second independent analysis. The "fingerprint" is the collection of microsatellite allele sizes at each locus for 'Beineke 10.'

DNA was isolated from the leaves of 10 black walnut trees obtained from Walter Beineke using CTAB extraction buffer (50 mM TRIS-HCL, pH 8.0, 20 mM EDTA, pH 8.0, 0.7M NaCl, 0.4M LiCl, 2% SDS, 2% TAB, nd 1% PVP). After isolation the DNA from each tree was quantified and diluted with nanopure distilled water to a final concentration of 5 ng/μL. The samples were stored in 96-well plates at 20° C.

For purposes of DNA fingerprinting, nine highly polymorphic loci from a suite of microsatellites developed by Woeste et al. (2002) were chosen. Amplification of each locus was performed with an MJ Research Tetrad Thermocycler (Waltham, Mass.) using 10 μL reactions in 96-well plates. The PCR reaction mix contained 2 μL of the aforementioned black walnut DNA, 5 μL Sigma Taq ReadyMix (Sigma Aldrich, St. Louis, Mo.), 0.4 μL of a 20 pmol mixture of forward and reverse fluorescence labeled primer, and 3 μL PCR grade water supplied with the Sigma ReadyMix. PCR amplification was for 30 cycles of 94° C. for 20 sec, 55° C. for 30 sec, and 72° C. for 1 min. All primers were annealed at 55° C. The products were then held at 4° C. until aliquots could be loaded into 6% Long Ranger (polyacrylamide) denaturing gels (BMA, Rockland, Me.). For each individual

0.5 μ L PCR product was added to 0.75 μ L blue dextran and 0.25 μ L of CXR 350 bp Ladder Standard (Promega, Fitchburg Center, Wis.) in a new 96-well 1 late. The samples were denatured for 2 min at 95° C. and loaded onto a CAL96 96-well laminated membrane comb (The Gel Company, San Francisco, Calif.). Electrophoresis was at 3,000 V, 60 mA, 200 Watts, 50° C. for 2 hours using an ABI 377 (Perkin Elmer) with 36 cm plates and 0.2 mm spacers. The resulting data was analyzed using ABI's GeneScan 3.1.2 and Genotyper 2.5 (Perkin Elmer). Microsatellite sizes were checked against previously published standards and verified by a second independent analysis. The “fingerprint” is the collection of microsatellite allele sizes at each locus for each tree.

Locus	Forward	Reverse
WGA2	GACGACGAAGGTGTACGGAT (SEQ ID NO: 1)	GTACGGCTCTCCTTGCGATC (SEQ ID NO: 10)
WGA6	CCATGAAACTTCATGCGTTG (SEQ ID NO: 2)	CATCCCAAGCGAAGGTTG (SEQ ID NO: 11)
WGA24	TCCCCCTGAAATCTTCTCCT (SEQ ID NO: 3)	TTCTCGTGGTGCTTGTTGAG (SEQ ID NO: 12)
WGA32	CTCGGTAAGCCACACCAATT (SEQ ID NO: 4)	ACGGGCAGTGTATGCATGTA (SEQ ID NO: 13)
WG33	TGGTCTGCGAAGACACTGTC (SEQ ID NO: 5)	GGTTCGTCGTTTGTTGACCT (SEQ ID NO: 14)
WGA86	ATGCCTCATCTCCATTCTGG (SEQ ID NO: 6)	TGAGTGGCAATCACAAGGAA (SEQ ID NO: 15)
WGA89	ACCCATCTTTCACGTGTGTG (SEQ ID NO: 7)	TGCCTAATTAGCAATTTCCA (SEQ ID NO: 16)

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Locus	Forward	Reverse
WGA90	CTTGTAATCGCCCTCTGCTC (SEQ ID NO: 8)	TACCTGCAACCCGTTACACA (SEQ ID NO: 17)
WGA97	GGAGAGGAAAGGAATCCAAA (SEQ ID NO: 9)	TTGAACAAAAGGCCGTTTTTC (SEQ ID NO: 18)

The best interpretation of the current data indicates that the probability that any other black walnut tree would have the collection of microsatellite allele sizes listed is less than 1 in 10⁻¹⁷.
Sizes (bp) of microsatellites at 9 loci used to fingerprint ‘Beineke 10’ (2 alleles at each locus).

WGA2		WGA6		WGA24		WGA32		WGA90	
136	140	142	142	234	246	169	181	160	162
WGA86		WGA97		WGA33		WGA89			
218	226	165	171	220	220	181	181		

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I claim	which has good timber quality, is fast growing, has fairly
1. A new and distinct variety of black walnut tree named	strong central stem tendency.
‘Beineke 10’, substantially as illustrated and described,	* * * * *



FIG. 1

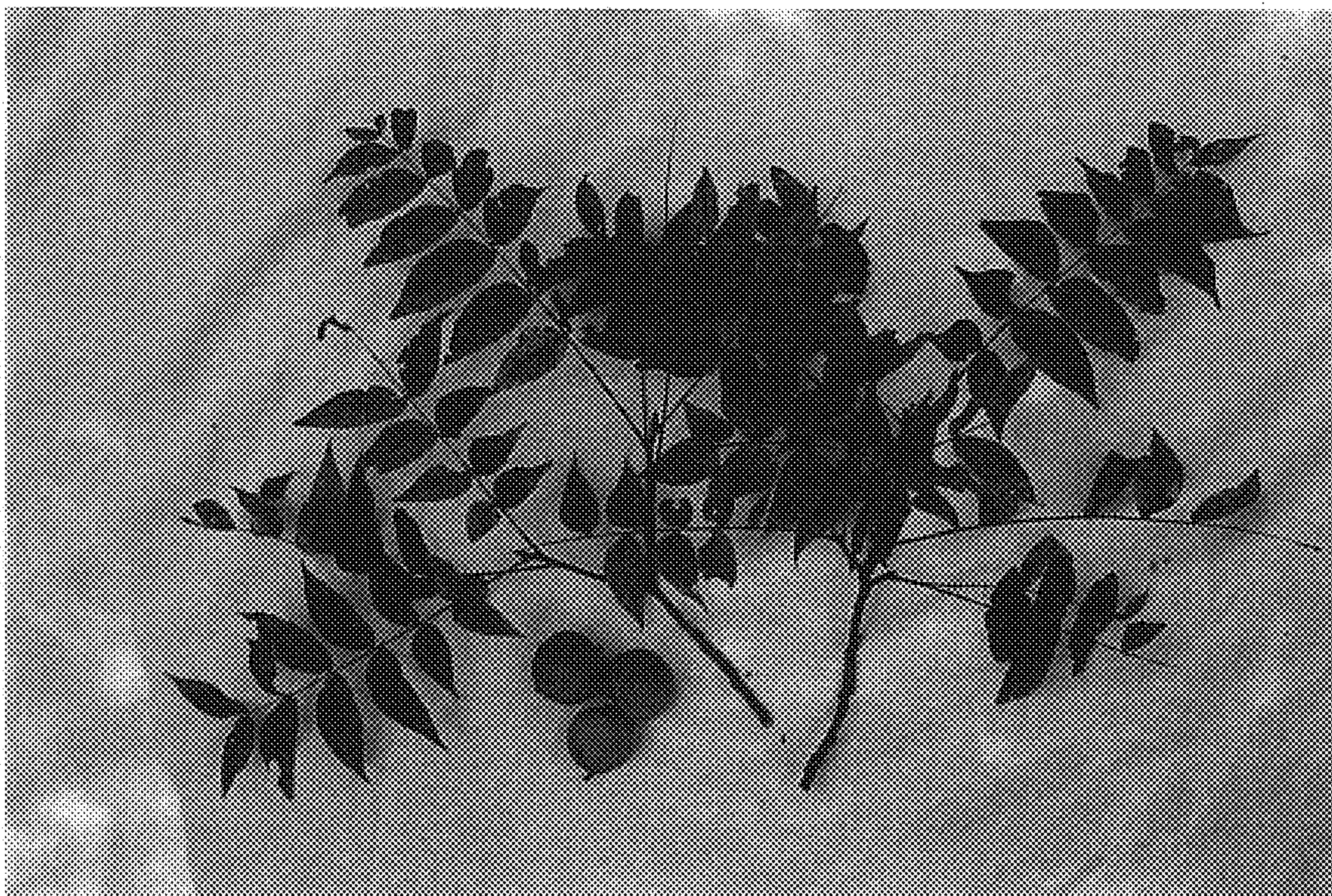


FIG. 2

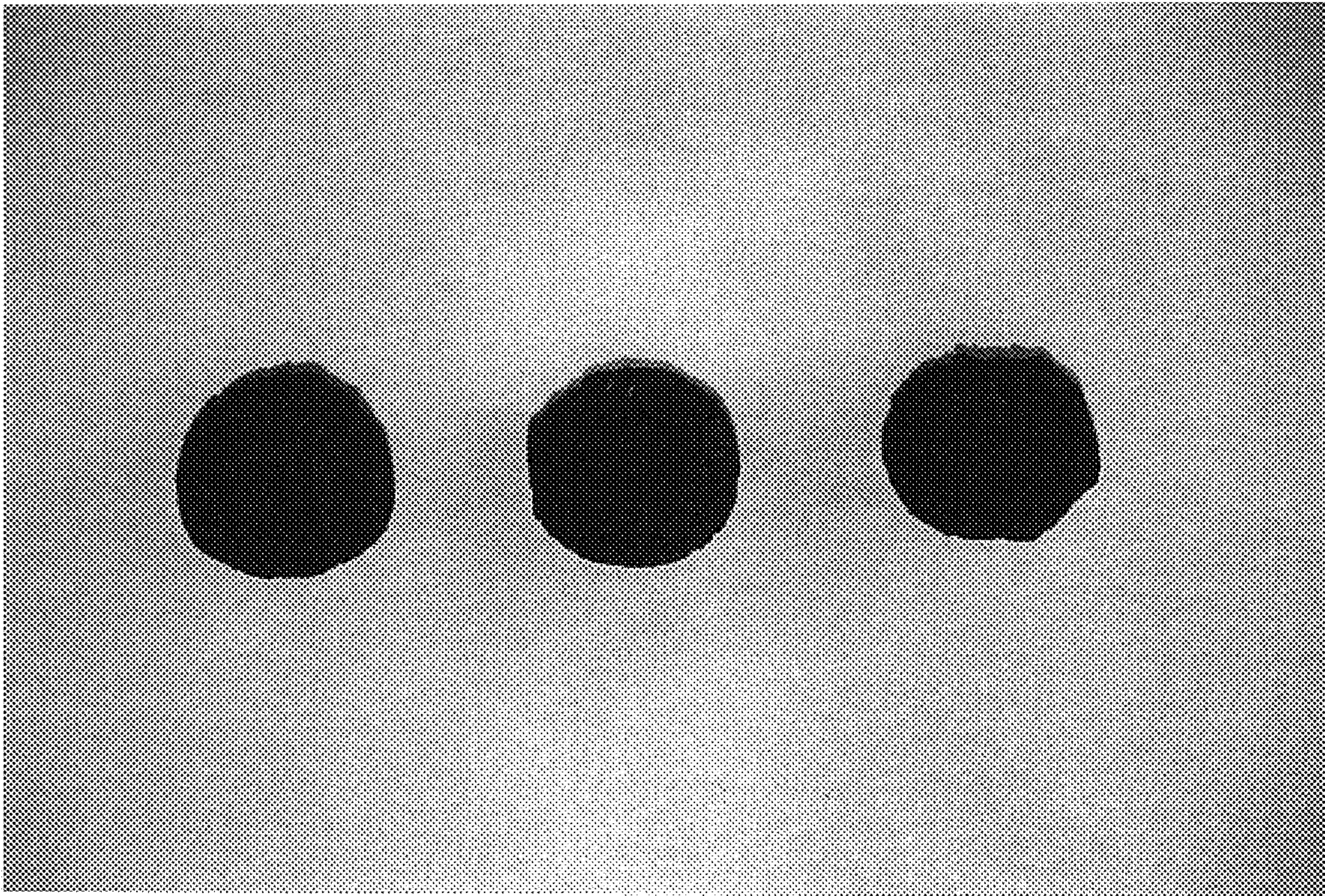


FIG. 3