



Standard Test Method for Immunological Measurement of Four Principal Allergenic Proteins (Hev b 1, 3, 5 and 6.02) in Natural Rubber and Its Products Derived from Latex¹

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^{ε1} NOTE—Footnote 6 editorially corrected in January 2010.

1. Scope

1.1 This test method covers an immunological method known as an immunoenzymetric assay to quantify the amount of 4 principal *Hevea brasiliensis* [*Hev b*] allergenic proteins [Hev b 1, Hev b 3, Hev b 5 and Hev b 6.02] in natural rubber and its products² derived from latex using monoclonal antibodies specific for epitopes on these proteins. Since these assays quantify the levels of only 4 of the known 13 officially acknowledged allergens potentially present in natural rubber latex containing products, the sum of the four allergen levels shall be viewed as an indicator of the allergen burden and not as a measure of the total allergen content that can be released from the product.

1.2 For the purpose of this test method, the range of allergenic protein will be measured in terms of nanogram to microgram quantities per gram or unit surface area of a natural rubber containing product.

1.3 The test method is not designed to evaluate the potential of natural rubber containing materials to induce or elicit Type I (IgE-mediated) hypersensitivity reactions.

1.4 This test method should be used under controlled laboratory conditions to detect and quantify the level of 4 allergenic proteins found in natural rubber containing products. It should not be used to describe, appraise or assess the hazard or risk of these natural rubber containing materials or products under actual in use conditions.

1.5 The values stated in SI units are to be regarded as standard. No other units of measurement are included in this standard.

1.6 *This standard does not purport to address all of the safety concerns, if any, associated with its use. It is the responsibility of the user of this standard to establish appro-*

priate safety and health practices and determine the applicability of regulatory limitations prior to use.

2. Referenced Documents

2.1 *ASTM Standards:*³

D1193 Specification for Reagent Water

D4483 Practice for Evaluating Precision for Test Method Standards in the Rubber and Carbon Black Manufacturing Industries

D4678 Practice for Rubber—Preparation, Testing, Acceptance, Documentation, and Use of Reference Materials

D5712 Test Method for Analysis of Aqueous Extractable Protein in Natural Rubber and Its Products Using the Modified Lowry Method

D6499 Test Method for The Immunological Measurement of Antigenic Protein in Natural Rubber and its Products

E691 Practice for Conducting an Interlaboratory Study to Determine the Precision of a Test Method

3. Terminology

3.1 *Definitions:*

3.1.1 *accepted reference value (ARV)*—value that serves as an agreed upon reference for comparison and which is derived as (1) a theoretical or established value, based on scientific principles, (2) an assigned or certified value, based on experimental work of some national or international organization, or (3) a consensus or certified value, based on collaborative experimental work under the auspices of a scientific or engineering group.

3.1.1.1 *Discussion*—ARV is an average industrial reference material (IRM) property or parameter value established by way of a specified test program. In this standard, the ARV as defined in the IRMs for the reference antigens and capture and detection antibodies is determined by analyzing a high and low control in an inter-laboratory study and using the assigned

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² This procedure has not been validated for condoms, particularly lubricated condoms, which could contain surfactants or other ingredients that could interfere with the assay.

³ For referenced ASTM standards, visit the ASTM website, www.astm.org, or contact ASTM Customer Service at service@astm.org. For *Annual Book of ASTM Standards* volume information, refer to the standard's Document Summary page on the ASTM website.

values of these high and low controls to verify that the assay is in control and that the reagents are performing properly.

3.1.2 *accuracy*—the closeness of agreement between a test result and an accepted reference value.

3.1.3 *allergens*—protein antigens which induce allergic immune reactions typically mediated through IgE antibodies.

3.1.4 *analyte*—any element, ion, compound, substance, factor, infectious agent, cell, organelle, activity (enzymatic, hormonal, or immunological), or property the presence or absence, concentration, activity, intensity, or other characteristics of which are to be determined.

3.1.5 *antibody*—an immunoglobulin, a protein that is produced as a part of the humoral immune response which is capable of specifically combining with antigen.

3.1.5.1 *Discussion*—Any of numerous Y-shaped protein molecules produced by B lymphocytes as a primary immune response, each molecule and its clones having a unique binding site that can combine with the complementary site of an antigen, as on a virus or bacterium, thereby signaling other immune responses. (See *monoclonal antibody*.)

3.1.6 *antigen*—any substance that can stimulate the production of antibodies within an organism and combine specifically with them.

3.1.7 *background absorbance*—the absorbance reading in the solution resulting from non-specific interactions caused by the presence of chemicals, ions, etc., other than the analyte being measured.

3.1.8 *binding capacity*—within the context of this document, refers to the number of Hev b allergen molecules that a primary capture antibody can bind reproducibly under standardized assay conditions (pH, ionic strength, protein matrix, time, temperature).

3.1.9 *blocking solution*—a non-reactive protein solution used to prevent nonspecific antibody adsorption and to reduce background absorbance.

3.1.10 *calibration*—the standardization of an instrument setting or an assay configuration.

3.1.11 *calibration material/calibrator*—a material (for example, solution) of known quantitative/qualitative characteristics (for example, concentration, activity, intensity, reactivity) used to calibrate, graduate, or adjust a measurement procedure or to compare the response obtained with the response of a test specimen/sample.

3.1.12 *concentration range*—the recommended analyte concentration range in nanograms per mL to micrograms per mL that produces an absorbance reading from 0.1 to 2.0–3.0 units (depending on the instrument).

3.1.13 *data reduction algorithm*—a mathematical process that converts assay-response data (for example, absorbance units) into interpolated dose results.

3.1.13.1 *Discussion*—The dose–response relationship in the assay is defined by the standard, reference, or calibration curve.

3.1.14 *detection limit/limit of detection*—the smallest quantity of an analyte that can be reproducibly and a statistically significant manner distinguished from the variance of the background, or a zero calibrator in a given assay system.

3.1.14.1 *Discussion*—It is usually defined at the 95 % confidence interval and has also been called the lower detection

limit or positive threshold of the assay; this term is not synonymous with analytical sensitivity.

3.1.15 *enzyme linked immunosorbent assay (ELISA)*—an immunological test method to quantify antigen or antibody levels using an enzyme as the detection mechanism.

3.1.16 *epitope/determinant*—(1) the minimum molecular structure of the antigenic site that will react with an antibody; (2) any site on an antigen molecule at which an antibody can bind; the chemical structure of the site determining the specific combining antibody.

3.1.17 *IgE*—human IgE is an immunoglobulin of the approximate molecular weight of 190 000, which exists normally in monomeric form and constitutes approximately 0.0005 % of the total serum immunoglobulins.

3.1.17.1 *Discussion*—It binds with high affinity to FcεR1 receptors on mast cells and basophils and FcεR2 receptors on a number of cells. IgE mediates the production and release of vasoactive mediators following the binding of allergen.

3.1.18 *immunoenzymetric assay (IEMA)*—a two-site non-isotopic immunological test method that employs two antibodies, a primary antibody to capture and a secondary enzyme conjugated antibody to detect the analyte of interest.

3.1.19 *immunoglobulin*—a glycoprotein composed of two heavy and two light chains that functions as an antibody. Human immunoglobulins have been subdivided into different isotypes (IgM, IgG, IgA, IgD, IgE), each of which possess a unique set of antigenic markers, physiochemical properties, and each of which produce a different pattern of effector functions (receptor binding, complement activation, opsonization).

3.1.19.1 *Discussion*—All antibodies are immunoglobulins, but it is not certain that all immunoglobulins possess antibody function.

3.1.20 *industry reference materials (IRM)*—materials that have been prepared according to a specified production process to generate a uniform lot; the parameters that define the quality of the lot are evaluated by a specified measurement program.

3.1.20.1 *Discussion*—IRMs are divided into two types according to the production process for generating the material.

3.1.21 *linearity*—the ability (within a given range) of an assay to provide results that are directly proportional to the concentration [amount] of the analyte in the test sample.

3.1.22 *monoclonal antibody*—antibody produced by cells created through the fusion of an antibody producing cell (B-lymphocyte) with immortal cancer cells.

3.1.22.1 *Discussion*—This process produces a hybrid (hybridoma) that expresses properties of both cells. The cells are all identical since they derive from a single cell and are called “monoclonal.”

3.1.23 *parallelism*—extent to which the dose–response relationship between two materials (that is, calibrator versus unknown specimens) is constant for the examined range of concentrations.

3.1.23.1 *Discussion*—Parallelism is a property (and a requirement) of quantitative immunoassays in which the calibrator and test sera produce parallel dose–response curves.

3.1.24 *precision*—the closeness of agreement between independent test results obtained under prescribed conditions; agreement between replicate measurements.

3.1.24.1 *Discussion*—Precision has no numerical value but is expressed in terms of imprecision—the standard deviation (SD) or the coefficient of variation (CV: SD/mean) of the results in a set of replicate measurements.

3.1.25 *precision profile*—the precision of an assay across the analyte concentration range of interest.

3.1.25.1 *Discussion*—A precision profile is constructed by determining the standard deviation (or coefficient of variation) of replicate measurements (within assays, between assays, or between specimen dilutions within an assay) spanning the entire analyte concentration range, albeit without the exact knowledge of the true analyte concentration that is contained in the serum specimens. When the CV_{dose} (*Y*-axis) is graphed against the dose (*X*-axis), a precision profile plot is generated. The precision profile is also referred to as the “imprecision profile” by some investigators.

3.1.26 *primary antibody*—the antibody used first in an assay sequence that is specific for the antigen and is sometimes referred to as the capture antibody that binds the analyte of interest from a biological specimen.

3.1.27 *proficiency testing (PT)*—an independent (non-manufacturer sponsored) program in which challenge specimens are sent to participating laboratories to be evaluated in assays that measure a spectrum of analytes.

3.1.28 *qualitative assay*—an assay system that produces an indication of the presence or absence of an analyte but does not provide a precise estimate of the concentration of that analyte.

3.1.28.1 *Discussion*—A positive test result implies only that the assay signal exceeds the analytical threshold or positive cutoff point that has been set to obtain an arbitrary combination of diagnostic sensitivity and specificity.

3.1.29 *quantitative assay*—an assay system that produces an accurate and reproducible estimate of the concentration of an analyte in the test specimen.

3.1.29.1 *Discussion*—Its analysis involves interpolation from a calibration curve, which is referenced to a readily available standard reference preparation.

3.1.30 *quality control response*—level of analyte produced by an assay for a quality control specimen that has a previously defined analyte concentration range as defined by the manufacturer.

3.1.30.1 *Discussion*—Assay performance was evaluated by determining the agreement in Hev b 1, 3, 5 or 6.02 levels obtained for two quality control extracts containing a high or low level of each Hev b allergen, following analysis in multiple laboratories participating in the multi-center study.

3.1.31 *reference solution*—the solution against which the test sample is being compared.

3.1.32 *relative standard deviation (RSD)*—the coefficient of variation which is the standard deviation divided by the mean.

3.1.33 *repeatability*—precision under conditions where independent test results are obtained with the same method on identical test items in the same laboratory by the same operator using the same equipment within short intervals of time.

3.1.34 *repeatability limit (r)*—the value below which the absolute difference between two individual test results obtained under repeatability conditions may be expected to occur with a probability of approximately 0.95 (95 %).

3.1.34.1 *Discussion*—The repeatability limit is 2.8 (~1.96 · square root of 2) times the repeatability standard deviation. This multiplier is independent of the size of the inter-laboratory study.

3.1.35 *reproducibility*—precision obtained under conditions where test results are obtained with the same method on identical test items in different laboratories with different operators using different equipment.

3.1.36 *reproducibility limit (R)*—the value below which the absolute difference between two test results obtained under reproducibility conditions may be expected to occur with a probability of approximately 0.95 (95 %).

3.1.36.1 *Discussion*—The reproducibility limit is 2.8 (~1.96 · square root of 2) times the reproducibility standard deviation. This multiplier is independent of the number of laboratories participating.

3.1.37 *secondary antibody—in an IEMA*, it is the enzyme conjugated antibody used second in a sequence that is specific for the analyte or interest and that completes the sandwich of the analyte.

3.1.38 *standard solution*—the preparation containing a standard analyte that is used as a reference to which the unknown sample being measured is compared.

3.1.39 *substrate*—the material or substance upon which an enzyme reacts.

3.1.40 *titer*—the strength of an antibody in solution that takes into consideration its concentration and affinity.

4. Summary of Test Method

4.1 This standard defines a general laboratory method called the immunoenzymetric assay. The first step in the method involves the extraction of a latex-containing product using a procedure previously described in Test Methods **D5712** and **D6499**. In brief, the latex product is extracted for 2 h in an aqueous buffer, typically at 5 to 10 mL per gram with agitation such as rotation or shaking. Following the extraction process, the extract is recovered and the level of 4 Hev b allergens is quantified using a two-site, non-competitive binding immunoenzymetric assay (IEMA) **(1)**.⁴

4.2 The general IEMA design is based on the use of a monoclonal antibody which is specific for a single Hev b allergenic protein that is attached to a microtiter plate by adsorption. This “capture” antibody has also been called the primary antibody in the assay system. Following blocking of the microtiter plate, the “unknown or test” extracts of the latex product are pipetted into replicate wells in the plate. In a different part of the plate, calibrators with pre-defined levels of the relevant Hev b allergenic protein are pipetted in replicate to produce a calibration curve. These calibrators can be either reference proteins or working calibrators that have been cross-validated against the reference proteins. Additionally,

⁴ The boldface numbers in parentheses refer to the list of references at the end of this standard.