AOAC SMPR 2010.004

Standard Method Performance Requirements for Immunological-Based Handheld Assays (HHAs) for Detection of *Bacillus anthracis* Spores in Visible Powders

Intended Use: Field use by first responders for analysis of visible powders

Method Developer and Independent Validation Studies

Probability of Detection at the Acceptable Minimum Detection Level

1 Definitions

Probability of detection (POD) is the proportion of positive analytical outcomes for a qualitative method for a given matrix at a given agent level or concentration. POD is concentrationdependent. The acceptable minimum detection level (AMDL) is the predetermined minimum level of a biological threat agent, which must be detected by the candidate method with an estimated 5% lower confidence limit on the POD of 0.95 or higher. The AMDL is dependent on the intended use.

2 Test Conditions

AMDL is 10⁷ CFU/mL *Bacillus anthracis* Ames spores in candidate method sample collection buffer.

Table 1. Bacilius anthracis HHA method: inclusivity pa
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3 Acceptance Criteria

Estimated 5% lower confidence limit on the POD must be 0.95 or higher. (No more than one failure in 96 replicates.)

Inclusivity

1 Definition

Strains or isolates or variants of the target agent(s) that the method can detect (Table 1).

2 Test Conditions

Test spores of each member of the *Bacillus anthracis* inclusivity panel at AMDL.

3 Acceptance Criteria

100% positive results.

Note: In the case of a negative result, retest that strain 96 times with no failures allowed to demonstrate an estimated 5% lower confidence limit on the POD of 0.95 or higher.

Exclusivity

1 Definition

Nontarget agents, which are potentially cross-reactive, that are not detected by the method (Table 2).

2 Test Conditions

Test spores of each member of the *Bacillus anthracis* HHA exclusivity panel at 10 times AMDL.

3 Acceptance Criteria

100% negative results.

Cluster	Genotype	Strain	MRI No.ª	Origin	Characteristics
A1a	7	Canadian bison	107448	Wood bison	pX01+, pX02+, VNTR genotype group A1a
АЗа	45 ^b	V770-NP-1R	107240	Vaccine (USA)	pX01+, pX02-, VNTR genotype group A3a
A2	29	PAK-1	107518	Sheep (Pakistan)	pX01+, pX02+, VNTR genotype group A2
A3a	51	BA1015	107446	Bovine (MD)	pX01+, pX02+, VNTR genotype group A3a
A3b	62	Ames	107517	Bovine (Texas)	pX01+, pX02+, VNTR genotype group A3b
A3c	67	K3	107497	South Africa	pX01+, pX02+, VNTR genotype group A3c
A3d	68	Ohio ACB	107339	Pig	pX01+, pX02+, VNTR genotype group A3d
A4	69	SK-102 (Pakistan)	107449	Imported wool (Pakistan)	pX01+, pX02+, VNTR genotype group A4
A4	77	Vollum 1B	107539	USAMRIID ^a	pX01+, pX02+, VNTR genotype group A4
B1	82	BA1035	107451	Human (South Africa)	pX01+, pX02+, VNTR genotype group B1
B2	80	RA3	107520	Bovine (France)	pX01+, pX02+, VNTR genotype group B2
С	Unk⁰	2002013094 (240)	124030	Louisiana	pX01+, pX02+, VNTR genotype group C
A1a	8	Pasteur	107171	USAMRIID	pX01-, pX02+, VNTR genotype group A1a
A3b	59, 61 ^{<i>b</i>}	Sterne	107453	USAMRIID	pX01+, pX02-, VNTR genotype group A3b
A1b	23	Turkey No. 32	107255	Human (Turkey)	pX01+, pX02+, VNTR genotype group A1b
	Cluster A1a A3a A3a A3b A3c A3d A4 A4 B1 B2 C A1a A3b A1b	Cluster Genotype A1a 7 A3a 45 ^b A2 29 A3a 51 A3b 62 A3c 67 A3d 68 A4 69 A4 77 B1 82 B2 80 C Unk ^c A1a 8 A3b 59, 61 ^b A1b 23	Cluster Genotype Strain A1a 7 Canadian bison A3a 45 ^b V770-NP-1R A2 29 PAK-1 A3a 51 BA1015 A3b 62 Ames A3c 67 K3 A3d 68 Ohio ACB A4 69 SK-102 (Pakistan) A4 77 Vollum 1B B1 82 BA1035 B2 80 RA3 C Unk ^c 2002013094 (240) A1a 8 Pasteur A3b 59, 61 ^b Sterne A1b 23 Turkey No. 32	Cluster Genotype Strain MRI No. ^a A1a 7 Canadian bison 107448 A3a 45 ^b V770-NP-1R 107240 A2 29 PAK-1 107518 A3a 51 BA1015 107446 A3b 62 Ames 107517 A3c 67 K3 107497 A3d 68 Ohio ACB 107339 A4 69 SK-102 (Pakistan) 107449 A4 77 Vollum 1B 107539 B1 82 BA1035 107451 B2 80 RA3 107520 C Unk ^c 2002013094 (240) 124030 A1a 8 Pasteur 107171 A3b 59, 61 ^b Sterne 107453 A1b 23 Turkey No. 32 107255	Cluster Genotype Strain MRI No. ^a Origin A1a 7 Canadian bison 107448 Wood bison A3a 45 ^b V770-NP-1R 107240 Vaccine (USA) A2 29 PAK-1 107518 Sheep (Pakistan) A3a 51 BA1015 107446 Bovine (MD) A3b 62 Arnes 107517 Bovine (Texas) A3c 67 K3 107497 South Africa A3d 68 Ohio ACB 107339 Pig A4 69 SK-102 (Pakistan) 107449 Imported wool (Pakistan) A4 69 SK-102 (Pakistan) 107449 Imported wool (Pakistan) A4 77 Vollum 1B 107539 USAMRIID ^a B1 82 BA1035 107451 Human (South Africa) B2 80 RA3 107520 Bovine (France) C Unk ^c 2002013094 (240) 124030 Louisiana A1a <t< td=""></t<>

^a MRI = MRI Global; USAMRIID = The United States Army Medical Research Institute for Infectious Diseases.

Approved by AOAC SPADA on April 24, 2007.

^b Organism contains only seven of eight MLVA markers due to the lack of pX02. Genotypes listed are consistent with seven of the eight markers. (*Note*: Footnote applies to BA2 and BA14 genotype designations.)

^c Unk = Unknown.

Table 2. Bacillus anthracis HHA method: Exclusivity panel

No.	Species	Strain	Plasmid status
BANN1	B. cereus	S2-8	pXO1-, pXO2-
BANN2	B. cereus	ЗА	pXO1-, pXO2-
BANN3	B. thuringiensis	HD1011	pXO1-, pXO2-
BANN4	B. thuringiensis	97-27	pXO1-, pXO2-
BANN5	B. thuringiensis	HD682	pXO1-, pXO2-
BANN6	B. cereus	E33L	pXO1-, pXO2-
BANN7	B. cereus	D17	pXO1-, pXO2-
BANN8	B. thuringiensis	HD571	pXO1-, pXO2-
BANN9	B. cereus	Al Hakam	pXO1-, pXO2-
BANN10	B. cereus	ATCC 4342	pXO1-, pXO2-
BANN11	B. cereus	FM1	pXO1-, pXO2-
BANN12	B. cereus	G9241	pBCXO1+ª, pXO2-
BANN13	B. cereus	03BB102	pXO1+, capA+, capB+, capC+ ^b
BANN14	B. cereus	03BB108	pXO1+, capA+, capB+, capC+ ^b
BANN15	B. thuringiensis	subsp. israelensis HD 1002	pXO1-, pXO2-
BANN16	B. thuringiensis	subsp. <i>kurstaki</i> HD 1	pXO1-, pXO2-
BANN17	B. thuringiensis	subsp. <i>morrisoni</i> HD 600	pXO1-, pXO2-
BANN18	B. coagulans	ATCC 7050	pXO1-, pXO2-
BANN19	B. mycoides	ATCC 6462	pXO1-, pXO2-
BANN20	B. megaterium	ATCC 14581	pXO1-, pXO2-
BANN21	B. cohnii		pXO1-, pXO2-
BANN22	B. psychrosaccharolyticus		pXO1-, pXO2-
BANN23	B. benzoevorans		pXO1-, pXO2-
BANN24	B. megaterium		pXO1-, pXO2-
BANN25	B. orikoshii		pXO1-, pXO2-
BANN26	B. macroides		pXO1-, pXO2-
BANN27	B. clausil	DSM 8716T	
BANN28	B. vedderi	DSM 9768T	
BANN29	B. subtilis	ATCC 6051T	
BANN30	B. licheniformis		
BANN31	B. amyloliquefaciens		
BANN32	B. circulans		
BANN33	B. pumilus		
BANN34	Clostridium difficile		
BANN35	Clostridium sardiniense		
BANN36	Clostridium perfringens		

^a pBCXO1 is pXO1-like, but not identical.

^b capA, B, and C are contained within the pXO2 plasmid of *Bacillus anthracis*; however, only the capA, B, and C sequences are found in 03BB102 and 03BB108. *Approved by AOAC SPADA on April 15, 2009.* *Note*: In the case of a positive result, retest that strain 96 times with no failures allowed to demonstrate a 95% upper confidence limit on the POD of 0.05 or lower.

Environmental Interference

1 Definition

Ability of the assay to detect target organism in the presence of environmental substances and to be free of cross-reaction from environmental substances (*Annex A*).

2 Test Conditions

Test powders as liquid suspensions or solutions in the presence and absence of *Bacillus anthracis* Ames spores at the AMDL. Test swab materials in the presence and absence of *Bacillus anthracis* Ames spores at the AMDL.

3 Acceptance Criteria

No false positives and no false negatives observed.

Note: In the case of a false-positive or false-negative result, retest the material 96 times with no failures allowed.

Collaborative Validation Study

Reproducibility

1 Definition

Precision under conditions where independent test results are obtained with the same methods on equivalent test items in different laboratories with different operators using separate instruments.

2 Test Conditions

Test *Bacillus anthracis* Ames spores at AMDL and near neighbor organism at 10 times AMDL. At least 12 replicates per material per collaborator with 12 collaborators (four collaborators at each of three test sites).

3 Acceptance Criteria

Must produce at least 10 valid data sets. Report standard deviation of reproducibility (s_v) .

POD at the AMDL Under Reproducibility Conditions (formerly termed System False-Negative Rate)

1 Definition

Rate of positive system results in a population of known positive test portions.

2 Test Conditions

Test *Bacillus anthracis* Ames spores at AMDL. At least 12 replicates per collaborator with 12 collaborators (four collaborators at each of three test sites).

3 Acceptance Criteria

Data for target agent must demonstrate an estimated 5% lower confidence limit on the CPOD of 0.95 or higher, where CPOD is the probability of detection calculated from pooled valid collaborative data.

POD in the Absence of Analyte Under Reproducibility Conditions (formerly termed System False-Positive Rate)

1 Definition

Rate of positive system results in a population of known negative test portions.

2 Test Conditions

Test near neighbor organism at 10 times AMDL. At least 12 replicates per collaborator with 12 collaborators (four collaborators at each of three test sites).

3 Acceptance Criteria

Data for near neighbor must demonstrate a 95% upper confidence limit on the CPOD of 0.05 or lower, where CPOD is the probability of detection calculated from pooled valid collaborative data.

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ANNEX A Environmental Factors Panel for Validating HHAs for Biothreat Agents

1 Powders and Chemicals

Bacillus thuringiensis powders (e.g., Dipel) Powdered milk Powdered infant formula (Fe fortified) Powdered infant formula (low Fe formulation) Powdered coffee creamer Powdered sugar Talcum powder Wheat flour Baking soda Chalk dust Brewer's yeast Dry wall dust Cornstarch Baking powder GABA (Gama aminobutyric acid) L-Glutamic acid Kaolin Chitin Chitosan MgSO₄ Boric acid Powdered toothpaste Popcorn salt 2 Swab Materials

Cotton swab with plastic shaft Rayon swab with plastic shaft Macrofoam swab with plastic shaft Method Developer sample collection device

Approved by AOAC SPADA on August 3, 2009.