

CERTIFICATION

AOAC® Performance TestedSM

Certificate No.

112001

The AOAC Research Institute hereby certifies the test kit known as:

Solus One E. coli O157

manufactured by

Solus Scientific Solutions Ltd. 9 Mansfield Networkcentre Millennium Business Park Concord Way, Mansfield Nottinghamshire, NG19 7JZ

This method has been evaluated in the AOAC® *Performance Tested Methods*SM Program and found to perform as stated by the manufacturer contingent to the comments contained in the manuscript. This certificate means that an AOAC® Certification Mark License Agreement has been executed which authorizes the manufacturer to display the AOAC *Performance Tested* SM certification mark along with the statement - "THIS METHOD'S PERFORMANCE WAS REVIEWED BY AOAC RESEARCH INSTITUTE AND WAS FOUND TO PERFORM TO THE MANUFACTURER'S SPECIFICATIONS" - on the above mentioned method for a period of one calendar year from the date of this certificate (November 3, 2020 – December 31, 2021). Renewal may be granted at the end of one year under the rules stated in the licensing agreement.

Scott Coates, Senior Director
Signature for AOAC Research Institute

November 3, 2020

Date

METHOD AUTHORS

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KIT NAME(S)

Solus One E. coli O157

CATALOG NUMBERS

INDEPENDENT LABORATORY

SGS Vanguard Sciences Inc. 224 North Derby Lane North Sioux City, SD USA

AOAC EXPERTS AND PEER REVIEWERS

Thomas Hammack¹, Michael Brodsky², Wayne Ziemer³

- ¹US FDA CFSAN, College Park, MD, USA
- ² Brodsky Consultants, Thornhill, Ontario, CANADA
- ³ Independent Consultant, Loganville, GA, USA

APPLICABILITY OF METHOD

Analytes - Escherichia coli O157, including H7

Matrices – Fresh raw ground beef (~75% lean, 375 g) and fresh raw beef trim (~75% lean, 375 g).

Performance claims - No statistical difference was detected between Solus One *E. coli* O157 ELISA and the reference culture method of U. S. Department of Agriculture-Food Safety and Inspection Service *Microbiology Laboratory Guidebook* (MLG), 5C.00 (2019), Detection, Isolation and Identification of Top Seven Shiga Toxin-Producing Escherichia coli (STECs) from Meat Products and Carcass and Environmental Sponge (2) for fresh raw beef trim matrix. However, a statistically significant difference was observed between candidate and reference method results for fresh raw ground beef, where more positive results were detected with the candidate method at both high and low contamination levels.

REFERENCE METHOD

USDA FSIS MLG 5C.00 (2019), Detection, Isolation and Identification of Top Seven Shiga Toxin-Producing Escherichia coli (STECs) from Meat Products and Carcass and Environmental Sponge (2)

| ORIGINAL CERTIFICATION DATE November 3, 2020 | CERTIFICATION RENEWAL RECORD New Approval |
|---|---|
| METHOD MODIFICATION RECORD NONE | SUMMARY OF MODIFICATION NONE |
| Under this AOAC® <i>Performance Tested</i> SM License Number, 112001 this method is distributed by: NONE | Under this AOAC® <i>Performance Tested</i> SM License Number, 112001 this method is distributed as: NONE |

PRINCIPLE OF THE METHOD (1)

Solus One *E. coli* O157 is an antibody-based high sensitivity ELISA method paired with media and our proprietary media supplement – Solus One supplement: for the rapid and specific detection of *E. coli* O157, including H7 strains in select foods samples. Solus One *E. coli* O157 relies on antibodies attached to the wells of microplate strips that are highly specific to *E. coli* O157, including H7 antigens. Following enrichment, samples are heat treated and an aliquot is added to the antibody coated wells.

If *E. coli* O157 specific antigen is present in the samples, it will bind immunologically to the antibody. After washing to remove unbound material, an enzyme-labelled antibody will bind to the captured proteins and thus to the well. After a second wash step to remove any unbound enzyme-antibody, the enzyme substrate is added. The substrate reacts in the presence of the enzyme producing a blue color change in the sample well. The substrate reaction is stopped after 30 minutes with the addition of dilute sulfuric acid changing any blue color present in the wells to yellow.

Optical densities resulting from this color change are read within 10 minutes in a generic plate reader using a 450 nm filter (e.g. a microplate reader or a Dynex DS2 instrument plate reader), where a result of an $OD_{450} < 0.200$ is considered to be negative for the target pathogen and $OD_{450} \ge 0.200$ is considered to be positive for the target pathogen.

DISCUSSION OF THE VALIDATION STUDY (1)

Solus One *E. coli* O157 methods successfully recovered *E. coli* O157, including H7 species from select food matrixes analyzed. Using POD analysis, no statistically significant differences were observed for un-inoculated samples by the candidate methods (both manual and automated) and the respective MLG 5C.00 reference method for both food matrixes tested. Whilst an un-inoculated raw ground beef test portion returned a presumptive positive result on the candidate automated sample preparation method, it was later confirmed negative for *E. coli* O157:H7. However, statistically significant differences were observed between the candidate methods (both manual and automated) and the MLG 5C.00 reference method, tested on raw ground beef matrix samples; where significantly higher numbers of positive samples for both low- and high-inoculation levels were detected by the candidate method. In addition, for the raw beef trim matrix, no statistically significant differences were observed between the number of positive samples detected by the automated sample preparation candidate method and the MLG 5C.00 reference method.

POD analysis of Solus One *E. coli* O157 method Robustness indicated no statistically significant differences were observed between nominal and experimental combinations using the automated method.

The results of the inclusivity and exclusivity evaluation demonstrated 100% agreement with expected results for the test panels and confirmed the high specificity and selectivity of the method to *E. coli* O157, including H7 species.

The method offers the benefit of the use of either a manual sample preparation or automated sample preparation to obtain results. Each method was quick and simple to perform, providing results in 2 h post incubation of the selective enrichment. The small footprint of both methods offers the ability to test in various laboratories. The Dynex DS2 software is user friendly with the ability to track lot information and sample identification quickly and with ease. Additionally, the Dynex DS2 software and instrument also offer the ability to run multiple assays at one time and has an open platform.

| Tab | le 1: So | lus One <i>E. col</i> | <i>i</i> O157 Inc | lusivity Resu | ilts (| (1) | |
|-----|----------|-----------------------|-------------------|---------------|--------|-----|--|
|-----|----------|-----------------------|-------------------|---------------|--------|-----|--|

| Organism species + isolates | Characterization | Source | Origin | Result |
|---|----------------------|-------------------------------|--------------------------------|----------------|
| 5 1 11 110157.117 | | | | |
| Escherichia coli O157:H7 | Not available | VGSCI ^a 1 | Beef wild type | + ^b |
| Escherichia coli O157:H7 | Not available | VGSCI 2 | Beef wild type | + |
| Escherichia coli O157:H7 | stx- eae+ | USDA FSIS ^c 465-97 | FSIS control | + |
| (fluorescent control strain) | 51150 | T14/d 000 C 4 | | |
| Escherichia coli O157:H7 | EHEC, stx1,2 | TW 08264 | Human | + |
| Escherichia coli O157:H7 | EHEC, stx1,2 | TW 08612 | Human | + |
| Escherichia coli O157:H7 | EHEC, stx1,2 | TW 10022 | Human | + |
| Escherichia coli O157:H7 | EHEC, stx1,2 | TW 08616 | Human, HUS | + |
| Escherichia coli O157:H7 | EHEC, stx1,2 | TW 10012 | Human | + |
| Escherichia coli O157:H7 | Not available | ATCC ^e 35150 | Feces | + |
| Escherichia coli O157:H7 | Not available | ATCC 43894 | Feces, HUS | + |
| Escherichia coli O157:H7 | Not available | ATCC 43895 | Hamburger, outbreak associated | + |
| Escherichia coli O157:H7 | Not available | ATCC BAA-1882 | Bovine feces | + |
| Escherichia coli O157:H7 | Not available | ATCC 43889 | Feces, HUS | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP ^f -3 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-4 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-8 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-9 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-10 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-11 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-12 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-13 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-15 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-16 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-17 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-19 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-20 | Beef | + ^b |
| Escherichia coli O157:H7 | Not available | MDP-21 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-22 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-23 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-24 | Beef | + |
| Escherichia coli O137:H7 | Not available | MDP-25 | Beef | + |
| Escherichia coli O157:H7 | Not available | | Beef | + |
| | | MDP-26 | | |
| Escherichia coli O157:H7 | Not available | MDP-27 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-28 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-29 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-31 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-32 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-33 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-34 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-37 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-38 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-39 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-40 | Beef | + |
| Escherichia coli O157:H7 | EHEC eae+ stx+ gamma | MDP-42 | Beef | + |
| Escherichia coli O157:H7 | Not available | MDP-43 | Beef | + |
| Escherichia coli O157:Non-Motile | Not available | TW 10238 | Clinical | + |
| Escherichia coli O157:- | Not available | TW 06555 | Unknown | + |
| Escherichia coli O157:Non-Motile | Not available | TW 09096 | Unknown | + |
| Escherichia coli O157:Non-Motile | Not available | TW 02883 | Unknown | + |
| Escherichia coli O157:Non-Motile | Not available | TW 07984 | Clinical | + |
| ^a SGS Vanguard Sciences, Inc., North Sig | oux City. SD. | | | |

[°]SGS Vanguard Sciences, Inc., North Sioux City, SD.

^b+ = The target analyte was detected by Solus One *E. coli* O157.

^cU.S. Department of Agriculture, Food Safety and Inspection Services, Meat Safety & Quality Research Unit, Clay Center, NE.

^dSTEC Center at Michigan State University = Thomas S. Whittam STEC Center at Michigan State University, East Lansing, MI.

^eAmerican Type Culture Collection, Manassas VA.

 $^{^{\}it f}$ USMARC O157:H7 Diversity Panel, U.S. Meat Animal Research Center, Clay Center, NE.

| Table 2: Solus One <i>E. coli</i> O157 Exclusivity Results (1) | | | | | | | | |
|--|------------------|--------------------------|---------------------------|------------|--|--|--|--|
| Organism | Characterization | Source | Origin | Result | | | | |
| Escherichia coli | Not available | ATCC ^a 11229 | Clinical | _ <i>b</i> | | | | |
| Escherichia coli | Not available | ATCC 35218 | Canine | - | | | | |
| Escherichia coli O103:H11 | EHEC eae+, stx+ | CDC ^c 06-3008 | Clinical | - | | | | |
| Escherichia coli O103:K:H8 | stx- eae- | ATCC 23982 | Feces | - | | | | |
| Escherichia coli O111:H8 | stx+ eae+ | ATCC BAA-179 | Stool from human with HUS | - | | | | |
| Escherichia coli O111:K58(B4):H- | stx- eae+ | ATCC 33780 | Clinical | - | | | | |
| Escherichia coli O121:H19 | EHEC eae+, stx+ | CDC 02-3211 | Clinical | - | | | | |
| Escherichia coli O121:H19 | EHEC, stx1,2 | TW ^d 08004 | Clinical | - | | | | |
| Escherichia coli O145:H25 | EHEC, stx2 | TW 09153 | Not available | - | | | | |
| Escherichia coli O145:Non-Motile | EHEC eae+, stx+ | CDC 99-3311 | Clinical | - | | | | |
| Escherichia coli O26:H11 | EHEC eae+, stx+ | CDC 03-3014 | Clinical | - | | | | |
| Escherichia coli O26:H32 | non-STEC | TW 01209 | Not available | - | | | | |
| Escherichia coli O45:H10 | non-STEC | TW 03052 | Not available | - | | | | |
| Escherichia coli O45:H2 | EHEC eae+, stx+ | CDC 00-3039 | Clinical | - | | | | |
| Escherichia coli 069:Non-Motile | Not available | TW 07942 | Clinical, diarrhea | - | | | | |
| Escherichia coli O111:H8 | Not available | ATCC 3114 | Clinical | - | | | | |
| Acinetobacter baumannii | Not available | ATCC 19606 | Urine | - | | | | |
| Alcaligenes faecalis | Not available | ATCC 8750 | Not available | - | | | | |
| Bacillus cereus | Not available | ATCC 11778 | Not available | - | | | | |
| Bacillus pumilus | Not available | ATCC 700814 | Not available | - | | | | |
| Carnobacterium maltaromaticum | Not available | ATCC 27865 | Raw milk | - | | | | |
| Citrobacter braakii | Not available | ATCC 51113 | Snake | - | | | | |
| Citrobacter freundii | Not available | ATCC 8090 | Not available | - | | | | |
| Citrobacter koseri | Not available | ATCC 27156 | Not available | - | | | | |
| Cronobacter sakazakii | Not available | ATCC 29544 | Rabbit | - | | | | |
| Edwardsiella tarda | Not available | ATCC 15947 | Human feces | - | | | | |
| Enterobacter aerogenes | Not available | ATCC 13048 | Sputum | - | | | | |
| Enterobacter cloacae | Not available | ATCC 23355 | Not available | - | | | | |
| Enterococcus faecalis | Not available | ATCC 19433 | Not available | - | | | | |
| Hafnia alvei | Not available | ATCC 51815 | Milk | - | | | | |
| Klebsiella oxytoca | Not available | ATCC 43165 | Clinical | - | | | | |
| Listeria monocytogenes | Not available | ATCC 19111 | Poultry | - | | | | |
| Microbacterium testaceum | Not available | ATCC 15829 | Paddy | - | | | | |
| Proteus hauseri | Not available | ATCC 13315 | Not available | - | | | | |
| Proteus mirabilis | Not available | ATCC 25933 | Clinical | - | | | | |
| Proteus vulgaris | Not available | ATCC 6380 | Not available | - | | | | |
| Pseudomonas aeruginosa | Not available | ATCC 15442 | Animal water bottle | - | | | | |
| Shigella boydii | Not available | ATCC 9207 | Not available | - | | | | |
| Staphylococcus aureus | Not available | NCTC ^e 12493 | Not available | - | | | | |
| Streptococcus pyogenes | Not available | ATCC 12384 | Not available | - | | | | |

^aAmerican Type Culture Collection, Manassas VA. ^b- = The target analyte was not detected by Solus One *E. coli* O157.

 $^{{}^{}c}\textsc{Centers}$ for Disease Control and Prevention, Atlanta, GA.

^dSTEC Center at Michigan State University = Thomas S. Whittam STEC Center at Michigan State University, East Lansing, MI.

^eNational Collection of Type Cultures, Porton Down, Salisbury, UK.

| Table 3. Solus One E. coli O157 Results: Presumptive vs. Cor | nfirmed (1 | 1) |
|--|------------|----|
|--|------------|----|

| | | ELISA | | | Solu | s One <i>E. coli</i> O | 157 presumptive | Solu | us One <i>E. coli</i> | O157 confirmed | | |
|-------------------------------------|--|---------------------|--------------------------------|-------|-------|------------------------|-----------------|------|-----------------------|----------------|---------------|---------------------|
| Matrix | Strain | method ^a | MPN ^b /test portion | N^c | X^d | POD_{CP}^e | 95% CI | х | POD_{CC}^f | 95% CI | $dPOD_{CP}^g$ | 95% CI ^h |
| Fresh Raw | Escherichia coli | | N/A ^j | 5 | 1 | 0.20 | 0.00, 0.62 | 0 | 0.00 | 0.00, 0.44 | 0.20 | -0.36, 0.76 |
| Ground Beef (~75% lean, | O157:H7, MDP ⁱ -28 | Automated | 0.30 (0.13, 0.54) | 20 | 12 | 0.60 | 0.39, 0.78 | 14 | 0.70 | 0.48, 0.86 | -0.10 | -0.28, 0.08 |
| 375g) | MDP-28 | | 0.49 (0.22, 1.11) | 5 | 5 | 1.00 | 0.57, 1.00 | 5 | 1.00 | 0.57, 1.00 | 0.00 | -0.47, 0.47 |
| Fresh Raw | Escherichia coli O157:H7, MDP-28 | O157:H7, Manual | N/A | 5 | 0 | 0.00 | 0.00, 0.44 | 0 | 0.00 | 0.00, 0.44 | 0.00 | -0.47, 0.47 |
| Ground Beef (~75% lean, 375g) | | | 0.30 (0.13, 0.54) | 20 | 12 | 0.60 | 0.39, 0.78 | 14 | 0.70 | 0.48, 0.86 | -0.10 | -0.28, 0.08 |
| | | | 0.49 (0.22, 1.11) | 5 | 5 | 1.00 | 0.57, 1.00 | 5 | 1.00 | 0.57, 1.00 | 0.00 | -0.47, 0.47 |
| Fresh Raw | - 1 . 1 . 1 | | N/A | 5 | 0 | 0.00 | 0.00, 0.44 | 0 | 0.00 | 0.00, 0.44 | 0.00 | -0.47, 0.47 |
| Beef Trim (~75% lean, | O157:H7, | • | 0.40 (0.21, 0.69) | 20 | 8 | 0.40 | 0.22, 0.61 | 8 | 0.40 | 0.22, 0.61 | 0.00 | -0.13, 0.13 |
| 375g) | ATCC* 43895 | | 0.95 (0.49, 1.55) | 5 | 5 | 1.00 | 0.57, 1.00 | 5 | 1.00 | 0.57, 1.00 | 0.00 | -0.47, 0.47 |

[&]quot;The Solus One E. coli O157 ELISA sample preparation method was manual and/or automated using the Dynex DS2. All were using the DS2, with the manual read using the instrument set on "Plate Read Only".

^bMPN = Most Probable Number is calculated using the LCF MPN calculator provided by AOAC RI, with 95% confidence interval.

 $^{^{}c}N$ = Number of test portions.

 $^{^{}d}x$ = Number of positive test portions.

^ePOD_{CP} = Candidate method presumptive positive outcomes divided by the total number of trials.

^fPOD_{CC} = Candidate method confirmed positive outcomes divided by the total number of trials.

 $[^]g$ dPOD_{CP} = Difference between the candidate method presumptive result and candidate method confirmed result POD values.

^h95% CI = If the confidence interval of a dPOD does not contain zero, then the difference is statistically significant at the 5% level.

ⁱMDP = USMARC O157:H7 Diversity Panel, U.S. Meat Animal Research Center, Clay Center, NE.

 $^{^{}j}N/A = Not applicable.$

^kATCC = American Type Culture Collection, Manassas, VA.

| | | ELISA | | | So | olus One <i>E. coli</i> O157 results MLG 5C.00 | | 0 results | _ | | | | |
|--|------------------------------|------------------------|--------------------------------|----------------|-------|--|------------|------------|-----------|------------|------------|---------------------|-------------|
| Matrix | Strain | method ^a | MPN ^b /test portion | N ^c | X^d | POD_{C}^e | 95% CI | х | POD_R^f | 95% CI | $dPOD_c^g$ | 95% CI ^h | |
| Fresh Raw Ground Beef (~75% lean, 375g) | Escherichia coli | | N/A ^j | 5 | 0 | 0.00 | 0.00, 0.44 | 0 | 0.00 | 0.00, 0.44 | 0.20 | -0.44, 0.44 | |
| | O157:H7, | 7, Automated | 0.30 (0.13, 0.54) | 20 | 12 | 0.60 | 0.39, 0.78 | 5 | 0.25 | 0.11, 0.47 | 0.35 | 0.04, 0.58 | |
| | MDP ⁱ -28 | | 0.49 (0.22, 1.11) | 5 | 5 | 1.00 | 0.57, 1.00 | 2 | 0.40 | 0.12, 0.77 | 0.60 | 0.03, 0.88 | |
| Fresh Raw | Factor delice and | | N/A | 5 | 0 | 0.00 | 0.00, 0.44 | 0 | 0.00 | 0.00, 0.44 | 0.00 | -0.44, 0.44 | |
| Ground Beef (~75% lean, | Escherichia coli O157:H7, | | 0.30 (0.13, 0.54) | 20 | 12 | 0.60 | 0.39, 0.78 | 5 | 0.25 | 0.11, 0.47 | 0.35 | 0.04, 0.58 | |
| 375g) | MDP-28 | | 0.49 (0.22, 1.11) | 5 | 5 | 1.00 | 0.57, 1.00 | 2 | 0.40 | 0.12, 0.77 | 0.60 | 0.03, 0.88 | |
| Fresh Raw | English Salita and | Raw Fach a sinh in and | | N/A | 5 | 0 | 0.00 | 0.00, 0.44 | 0 | 0.00 | 0.00, 0.44 | 0.00 | -0.44, 0.44 |
| Beef Trim (~75% lean, 375g) | O157:H7, | • | 0.40 (0.21, 0.69) | 20 | 8 | 0.40 | 0.22, 0.61 | 6 | 0.30 | 0.15, 0.52 | 0.20 | -0.10, 0.45 | |
| | ATCC ^k 43895 | | 0.95 (0.49, 1.55) | 5 | 5 | 1.00 | 0.57, 1.00 | 4 | 0.80 | 0.38, 1.00 | 0.20 | -0.28, 0.63 | |

[&]quot;The Solus One E. coli O157 ELISA sample preparation method was manual and/or automated using the Dynex DS2. All were using the DS2, with the manual read using the instrument set on "Plate Read Only".

REFERENCES CITED

- 1. Kelly, S., Higgins, D., Tonner, E., Anthony, J.P., Bradley, G., Illingworth, S., Perera, N., Clemens, N., and Thompson-Strehlow, L., Validation of the Solus One *E. coli* O157 Test Method for Detection of *E. coli* O157 in Select food Matrixes, AOAC® *Performance Tested*^{5M} certification number 112001.
- 2. U. S. Department of Agriculture-Food Safety and Inspection Service *Microbiology Laboratory Guidebook*, 5C.00 (2019), Detection, Isolation and Identification of Top Seven Shiga Toxin-Producing Escherichia coli (STECs) from Meat Products and Carcass and Environmental Sponge (Accessed October 2020) https://www.fsis.usda.gov/wps/wcm/connect/7ffc02b5-3d33-4a79-b50c-81f208893204/mlg-5.pdf?MOD=AJPERES

^bMPN = Most Probable Number is calculated using the LCF MPN calculator provided by AOAC RI, with 95% confidence interval.

^cN = Number of test portions.

 $^{^{}d}x$ = Number of positive test portions.

^ePOD_c = Candidate method presumptive positive outcomes confirmed positive divided by the total number of trials.

^fPOD_R = Reference method confirmed positive outcomes divided by the total number of trials.

^gdPOD_C = Difference between the candidate method and reference method POD values.

^h95% CI = If the confidence interval of a dPOD does not contain zero, then the difference is statistically significant at the 5% level.

ⁱATCC = American Type Culture Collection, Manassas, VA.

 $^{^{}j}N/A = Not applicable.$

^kATCC = American Type Culture Collection, Manassas, VA.