Part C. BIT Summary Template



| Verification of Validat tested by (AST Method) | | | | | | | | | |
|-----------------------------------------------------------------------------------------|------------------------------|-----------------------|-------------------------|----------------------|-----------------|-------------|----------|-----------|-----------------------------------|
| Studies performed (dates): | | | | | | | | | |
| I. Purpose Verify or Validate perfor For organism or organis Reference/Comparator resu | mance of mance of management | (Name of Me | thod or Con | nmercial AST | Device) _ | | | | |
| For Antimicrobial(s) and Brea | | | | | | | | | |
| | Old Breakpoints (MIC μg/ml) | | | Nev | / Breakpoin | ts (MIC μg | | | |
| Antimicrobial(s) | S | SDD | 1 | R | S | SDD | 1 | R | Breakpoint Source (FDA/CLSI) |
| | | | | | | | | | |
| | | | | | | | | | |
| | | | | | | | | | |
| Abbreviations: I, intermediate; MIC, mini | imal inhibitory (| concentration; R, res | sistant; S, suscept | ible; SDD, susceptil | ble dose depend | lent. | | | |
| II. Verification/Validation St | tudv | | | | | | | | |
| A. AST System | , | | | | | | | | |
| Panel/Card | | | | Software ver | sion | | | | |
| B. Accuracy | | | | | | | | | |
| Number of isolates | | | | | | | | | |
| Isolate source(s) (eg, CDC & FDA Antibio | | | | | | | | | |
| Reference result source (eg, CDC & FDA AR Isol | | AICs, in-hous | e reference | broth microd | dilution, re | ference lab | oratory) | | |
| | | | | | | | | tor AST r | method that is verified/validated |



for the new breakpoints or preestablished using a reference (eg, CDC & FDA AR Isolate Bank) or verified/validated comparator method.



| C. Reproducibility (precision) | |
|--------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------|
| Number of isolates | |
| Isolate source(s) | |
| | ank, clinical isolates quality control strains) |
| D. Quality Control | |
| Isolate(s) | Testing frequency(eg. per run) |
| (ie, name/strain number) | (eg, per run) |
| E. Analysis | |
| 1. Interpret MIC results man | ually utilizing new breakpoints as listed above (see I. Purpose). |
| 2. Compare interpretive cate comparator results. | egory results (eg, S, SDD, I, R) obtained from test system to the interpretive category obtained from the reference, |
| 3. General guidance for acce | ptable accuracy |
| Categoric Agreement (CA) | ≥90% |
| Very Major Errors (VME) | <3% |
| Major Errors (ME) | <3% |
| Minor Errors (MiE) | Determined by the laboratory director. |
| 4. Note: A category agreeme (ie, within ±1 two-fold dil | ent of <90% may be acceptable if the majority of errors are minor and the minor errors have essential agreement ution). |
| 5. Acceptable reproducibility 95% of replicate results for | y or a single antimicrobial agent/organism fall into either an S, I, SDD, or R category. |
| III. Procedure | |
| A. Materials and testing proced | dure for system to be verified/validated |
| Described in SOP | (this Laboratory's SOP #) |
| B. Record results on Appendix I | 2 |





- **C.** Options for discrepancy resolution (following a check for transcription error or other possible human error that could lead to resolution without retesting)
 - **1.** Repeat in triplicate.
 - 2. Test using another method that has been verified/validated for new breakpoints (eg, disk diffusion)
 - **3.** Send isolate to a reference laboratory.
- **D.** Update data table (Part F or Part G) and perform analyses with resolved results.

IV. Calculation of Accuracy, Reproducibility, and Error Rates

- **A.** Accuracy: Calculate each agent separately.
 - 1. CA (%) = Number of isolates with same category results/total isolates x 100
 - 2. VMEs (%) = Number of "S" isolates (test system results)/number of "R" isolates (reference results) x 100
 - 3. MEs (%) = Number of "R" isolates (test system results)/number of "S" isolates tested (reference results) x 100
 - **4.** MiEs (%) = Number of isolates where one result (either test or reference) is "I or SDD" and the other is "S" or "R"/total isolates x 100
- B. Reproducibility

Number of results that are reproducible for each organism/drug combination/total number of isolates

V. Summary of Results Obtained

A. Accuracy

| | # of Isolates* | | | | CA | VME | ME | MiE | |
|----------|----------------|---|-----|---|----|-------|-------|-------|-------|
| Agent(s) | Total | S | SDD | 1 | R | # (%) | # (%) | # (%) | # (%) |
| | | | | | | | | | |
| | | | | | | | | | |
| | | | | | | | | | |
| | | | | | | | | | |

^{*}Numbers represent a summary of the reference results.

Abbreviations: CA, categorical agreement; I, intermediate; ME, major errors; MiE, minor errors; S, susceptible; SDD, susceptible dose dependent; R, resistant; VME, very major errors.





| B. Reproducibility | |
|--------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------|
| % of | (agent/organism) results were reproducible |
| I have reviewed theaccuracy and precision, for the (AST test method)method is considered acceptable for patient testing. | (verification/validation) data for, and the performance of the |
| Reviewed by: | Date: |
| Signature: | |

This is Part C of the of the 2023 Breakpoint Implementation Toolkit. To access the entire BIT Toolkit, visit https://clsi.org/bit-toolkit/.

