

### Clinical bacteriology numeric grading scheme guideline

Grade	Interpretation	Definition and examples
4	<b>Full value</b>	Accepted by the committee as the correct answer either in terms of current nomenclature or in terms of appropriate clinical relevancy, including listing pathogen-specific negative results, correct Antimicrobial Profile Reporting and/or descriptive reporting, e.g. MRSA, ESBL producer, VRE, and Notification of Public Health , i.e., <i>Salmonella</i> from a stool sample.
3	<b>Essentially correct or Acceptable</b>	A nomenclature or susceptibility error, generally at the species level, not technically correct but would have little or no clinical impact. A deviation from what is considered the most clinically relevant result, but one which would pose little difficulty in interpretation of the sample's report. For example: <i>Staphylococcus hominis</i> vs. <i>Staphylococcus epidermidis</i> <i>Enterobacter aerogenes</i> vs. <i>Enterobacter cloacae</i> Susceptible vs. intermediate Excessive over-reporting of susceptibility testing results (calculated as minus-1 from the full value).
2	<b>Separator</b>	To augment the difference between the two grading groups.
1	<b>Incorrect or Unacceptable</b>	A nomenclature error that would be wrong at the species level, but by reporting may have an impact on clinical interpretation and potentially a treatment error. A major susceptibility error. A clinical relevancy result that could lead to a diagnosis or treatment error. For example: <i>Corynebacterium jeikeium</i> vs. diphtheroids; <i>Staphylococcus aureus</i> vs. <i>Staphylococcus epidermidis</i> Identify VSE as VRE. Reporting the presence of <i>Neisseria meningitidis</i> from a throat swab.
0	<b>Very incorrect or Very unacceptable</b>	A nomenclature error that would be wrong at either the genus or species level or a very major susceptibility error that could result in a significant interpretation or treatment error. A clinical relevancy result that could lead to a major diagnosis or treatment error. For example: <i>Salmonella</i> species vs. <i>Citrobacter</i> species <i>Escherichia coli</i> vs. <i>Shigella dysenteriae</i> <i>Burkholderia cepacia</i> vs. <i>Pseudomonas aeruginosa</i> Identify <i>Neisseria meningitidis</i> in a blood culture as a contaminant Identify VRE as VSE. Reporting <i>Staphylococcus aureus</i> and <i>Escherichia coli</i> in a mixed blood culture as 'probable contaminants'.