

Technique	Application examples	Sensitivity	Specificity	Advantages	Disadvantages
Conventional cell-based	culture in/on several media, microscopy, Gram-staining, biochemical tests	good-excellent	good	<ul style="list-style-type: none"> <li>quite inexpensive</li> <li>qualitative and quantitative data</li> </ul>	<ul style="list-style-type: none"> <li>pathogens have to grow in artificial media</li> <li>labour-intensive and time-consuming</li> </ul>
Immunological	ELISA, serological assay, Microarray	moderate-good	moderate-good	<ul style="list-style-type: none"> <li>high-throughput capacity</li> <li>rapid</li> <li>relatively low-cost</li> <li>ease-of-use</li> <li>qualitative and quantitative data</li> </ul>	<ul style="list-style-type: none"> <li>detection limit for organisms/antigens with low abundance</li> <li>difficulties in generating selective antibodies</li> </ul>
Nucleic-acid based	Hybridisation ( <i>in situ</i> , Sandwich, Southern blot, Northern blot)	moderate-good	excellent	<ul style="list-style-type: none"> <li>rapid</li> <li>ease-of-use</li> </ul>	<ul style="list-style-type: none"> <li>detection limit for organisms with low abundance</li> <li>pre-cultivation necessary</li> <li>limit number of probes in one experiment</li> <li>unknown species cannot be detected</li> </ul>
	PCR (nested, multiplex, real time, competitive quantitative)	excellent	excellent	<ul style="list-style-type: none"> <li>rapid</li> <li>ease-of-use</li> <li>low amounts necessary</li> <li>qualitative and quantitative data</li> </ul>	<ul style="list-style-type: none"> <li>limited capacity for multiplexing</li> <li>unknown species cannot be detected</li> </ul>
	sequencing	good-excellent	excellent	<ul style="list-style-type: none"> <li>provides most detailed, unbiased information</li> <li>reveals novel organisms</li> </ul>	<ul style="list-style-type: none"> <li>detection limit for organisms with low abundance</li> <li>expensive and time-consuming</li> </ul>
	DNA/RNA-Microarray	good	excellent	<ul style="list-style-type: none"> <li>high-throughput capacity</li> <li>ease-of-use</li> </ul>	<ul style="list-style-type: none"> <li>detection limit for organisms with low abundance</li> <li>medium expensive and time-consuming (when labelled)</li> <li>unknown species cannot be detected</li> </ul>
Other	MALDI-TOF mass spectrometry	good	good	<ul style="list-style-type: none"> <li>high-throughput capacity</li> <li>rapid</li> <li>generates easily interpretable spectra</li> <li>qualitative and quantitative data</li> <li>low overall operating costs</li> </ul>	<ul style="list-style-type: none"> <li>detection limit for organisms with low abundance</li> <li>host proteins and normal flora might overlap mass spectra</li> <li>high initial investments and maintenance costs</li> <li>lacking differentiation of closely related species</li> </ul>