

PCR-/NAT *Mycoplasma pneumoniae*
(RV 541) November 2019



Tabelle 1: Probenzusammensetzung und erwartetes Ergebnis.
Sample composition and expected results.

	<i>Erwartet / expected</i>		<i>Probenzusammensetzung / Sample composition</i>
1925411	++	61	<i>Mycoplasma pneumoniae</i> (~ 5x10 ⁴ genome copies/mL)
1925412	∅	62	<i>Escherichia coli</i> K12
1925413	+++	61	<i>Mycoplasma pneumoniae</i> (~ 5x10 ⁵ genome copies/mL)
1925414	+	61	<i>Mycoplasma pneumoniae</i> (~ 5x10 ³ genome copies/mL)

Tabelle 2: Häufigkeit der Mitteilung verschiedener Befunde.
Absolute numbers of reported individual results.

<i>n = 139</i>	<i>Probennummer (Sample no.)</i>					<i>Inhibition</i>			
	1925411	1925412	1925413	1925414		1925411	1925412	1925413	1925414
Befund <i>Result</i>									
Positiv	138	2	138	133	n.d.	0	0	0	0
Negativ	1	137	1	6 ¹⁾	nein <i>no</i>	139	139	139	139
Fraglich <i>Questionable</i>	0	0	0	0	ja <i>yes</i>	0	0	0	0

Tabelle 3: Häufigkeit richtig positiver und richtig negativer NAT-Befunde bei Anwendern verschiedener Methoden.

Absolute numbers and relative frequency of reported true positive and true negative results among various NAT methods.

NAT-Methode [Code] (total number)	NAT richtig positiv <i>True positive results</i>						NAT richtig negativ	
	1925411		1925413		1925414		1925412	
	Absolut <i>Absolute</i>	%	Absolut <i>Absolute</i>	%	Absolut <i>Absolute</i>	%	Absolut <i>Absolute</i>	%
LightMix <i>M.pneumoniae</i> [20] (n = 15)	14	93	15	100	15	100	15	100
AID CAP Bacteria [21] (n = 5)	5	100	5	100	5	100	5	100
AmpliGnost MP PCR Kit [23] (n = 6)	6	100	6	100	6	100	6	100
Diagenode MP/CP [24] (n = 3)	3	100	3	100	3	100	2	67
r-Biopharm RIDAGENE Mp [25] (n = 6)	6	100	6	100	6	100	6	100
GeneProof <i>M. pneumoniae</i> [26] (n = 8)	8	100	8	100	7	88	8	100
Commercial assay / kit [27] (n = 58)	58	100	57	98	54	93	57	98
<i>In house</i> PCR assay [28] (n = 38)	38	100	38	100	37	97	38	100

Comments: ¹⁾ As sample #1925414 contained a low number of *Mycoplasma pneumoniae* target organisms, negative PCR results were not rated "false negative" in this EQAS distribution.