

Q1 2018 Throat – Urine Culture



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Specimen 1 Urine - 76 year old Male, Post TURP, Foley catheter

Organisms	Extent 1	2	3	4	5	Total
943 - Aerobe found; but referred for ID	9	0	1	0	1	11
993 - Growth of gram-negative organisms	4	0	0	0	0	4
814 - <i>Klebsiella pneumoniae</i>	2	0	2	0	0	4
991 - <i>Klebsiella</i> or <i>Enterobacter</i>	0	0	3	0	0	3
811 - <i>Klebsiella</i> sp.; NOS	0	0	2	0	0	2
989 - <i>Klebsiella</i> ; <i>Staphylococcus</i> or <i>Streptococcus</i>	0	0	1	0	0	1
TOTAL PARTICIPANTS						25

Flagging appears for failure to report 811, 814, 942, 919, 942, 943, 983, 989, 991 or 993.

In addition to the required organism, participants in all extents may report (No additional codes)

This sample contained *Klebsiella pneumoniae*.

Specimen 2 Urine - 81 year old Female, Incontinency

Organisms	Extent 1	2	3	4	5	Total
943 - Aerobe found; but referred for ID	9	0	1	0	1	11
987 - <i>E.coli</i> ; <i>Citrobacter</i> or <i>Enterobacter</i>	1	0	3	0	1	5
993 - Growth of gram-negative organisms	4	0	0	0	0	4
799 - <i>Escherichia coli</i>	1	0	3	0	0	4
798 - <i>Escherichia</i> sp.; NOS	0	0	1	0	0	1
718 - Normal flora found, not normally reported	0	0	1	0	0	1
994 - Growth of gram-positive organisms	0	0	0	0	1	1
TOTAL PARTICIPANTS						27

Flagging appears for failure to report 798, 799, 919, 942, 943, 987 or 993.

In addition to the required organism, participants in all extents may report 718, 777, and 994

This sample contained *Escherichia coli* and *Corynebacterium* sp.

Specimen 3 Urine - 20 year old Male, Fever pain

Organisms	Extent 1	2	3	4	5	Total
943 - Aerobe found; but referred for ID	10	0	1	0	1	12
994 - Growth of gram-positive organisms	4	0	1	0	0	5
989 - <i>Klebsiella</i> ; <i>Staphylococcus</i> or <i>Streptococcus</i>	0	0	2	0	0	2
873 - <i>Staphylococcus</i> sp.; NOS	0	0	2	0	0	2
879 - <i>Staphylococcus saprophyticus</i>	0	0	2	0	0	2
942 - Primary culture only, refer for ID	1	0	0	0	0	1
874 - <i>Staphylococcus</i> sp.; coagulase-negative; NOS	0	0	1	0	0	1
TOTAL PARTICIPANTS						25

Flagging appears for failure to report 873, 874, 879, 942, 943, 989 or 994.

In addition to the required organism, participants in all extents may report (No additional codes)

This sample contained *Staphylococcus saprophyticus*.

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Specimen 4 Throat - 22 year old Female, Cough, fever, sore throat

Organisms	Extent	1	2	3	4	5	Total
922 - Neg for Grp A strep screen by culture		20	17	8	0	2	47
975 - Neg for strep Group A antigen		1	2	2	0	0	5
919 - Neg for beta-hemolytic strep screen		1	1	0	0	0	2
923 - Pos for Grp A strep screen by culture		0	0	1	0	0	1
718 - Normal flora found, not normally reported		0	1	0	0	0	1
		TOTAL PARTICIPANTS					56

Flagging appears for failure to report 891, 922, 975 or 994.

In addition to the required organism, participants in all extents may report 718.

This sample contained *Streptococcus* sp., Group B and alpha *Strep* & *Neisseria* sp.

Specimen 5 Throat - 6 year old Male, Mild sore throat

Organisms	Extent	1	2	3	4	5	Total
923 - Pos for Grp A strep screen by culture		20	16	9	0	2	47
976 - Pos for strep Group A antigen		1	2	2	0	0	5
921 - Pos for beta-hemolytic strep screen		1	1	0	0	0	2
922 - Neg for Grp A strep screen by culture		0	1	0	0	0	1
886 - <i>Streptococcus</i> sp.; beta-hemolytic Grp A (<i>S. pyogenes</i>)		0	1	0	0	0	1
		TOTAL PARTICIPANTS					56

Flagging appears for failure to report 886, 921, 923 or 976.

In addition to the required organism, participants in all extents may report 883.

This sample contained *Streptococcus pyogenes*, Group A and alpha *Strep*.