Chapter 11

Identification of Microorganisms Encountered in the Upper Respiratory Tract

Iris L. Sun

Department of Biology
Purdue University
West Lafayette, Indiana 47907-1392
(317) 494-4984, FAX (317) 494-0876

Iris received her Ph.D. in Cell Biology from the Department of Biological Sciences, Purdue University. She then stayed in the department as a research associate working on the action mechanism of anti-cancer drugs and studying the effect of cytokines on transformed cells. Besides doing research, she also teaches Lab of Medical Microbiology and Introduction of Microbiology in the department.

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Background Information

Differentiation of Potential Pathogens from the Upper Respiratory Tract

A. Gram Positive Aerobic Cocci: *Staphylococci*

*Staphylococci* occur typically as grape-like clusters because cell division occurs in three planes and daughter cells tend to remain attached. They grow well on basic or enriched media at temperatures ranging from 25°C to 35°C forming opaque, smooth, and glistening colonies. On sheep blood agar, colonies of *S. aureus* appear gold or bronze.

*Staphylococci* produce the enzyme catalase which decomposes hydrogen peroxide (H₂O₂) into oxygen (O₂) and water (H₂O). They are most resistant to extreme conditions, and can survive temperatures of 60°C for 1 hour or be carried on dust particles and live for months on dried pus or sputum. Another mechanism for their survival is a tolerance to high salt concentration, such as salty food. *Staphylococci* are resistant to phenol, sulfonamides and many antibiotics, but are very sensitive to basic dyes. They produce a high number of toxins which account for the variety of diseases, in addition to infections of the respiratory tract.

*Hemolysins* are hemolytic exotoxins acting on the cell membrane and lysing several kinds of blood and tissue cells.

*Leucocydins* are non hemolytic exotoxins which lyse lymphocytes, neutrophils and macrophages.

*Enterotoxin* is produced by about 50% of coagulase positive strains of *Staphylococci*. It is a common cause of food poisoning.

*Exfoliatin* damages skin cells resulting in peeling of large sheets of lining cells. It affects primarily young infants causing the scalded skin syndrome.

*Coagulase* causes the blood plasma to clot. It is only present in virulent strains and its property is the base for the coagulase test.

*Lipase* breaks down lipids in skin structure and lipoproteins in the blood. *Staphylococci* utilizes the metabolites derived from lipid breakdown, and this may explain the high number of *Staphylococci* on the skin surface.

*Hyaluronidase* increases tissue permeability to *Staphylococci*.

*Staphylokinase* dissolves clots and contributes to the spread of local infections.

Pathogenicity:

The abscess is the type lesion caused by *Staphylococci*, a consequence of its pus-forming ability and limited capacity for spread.
S. aureus is the most common pathogen while S. epidermidis is not pathogenic but may cause infections in some circumstances. Since both are inhabitants of the skin, most lesions are superficial and develop as boils, pustules, pimples, furuncles, carbuncles, or impetigo contagiosa. Staphylococcal pneumonia results from multiple abscesses in the lungs. The invasion of the body by Staphylococci (septicemia) results from secondary invasion of the bloodstream by organisms from a localized skin infection.

Bacteriological Diagnosis:

Gram-stained smears and cultures on blood agar are the first step toward Staphylococcal identification. The enzyme catalase distinguishes Staphylococci (catalase +) from Streptococci (catalase –). The Coagulase test distinguishes pathogenic staphylococci (coagulase +) from nonpathogenic (coagulase –). Virulent Staphylococci produce yellow pigment visible in cultures, ferment the sugar mannitol and produce deoxyribonucleases (DNAses).

Phage typing: Bacteriophages are viruses that specifically attach to bacteria, invade them and cause their lysis. When an agar plate is inoculated with bacteria from a pure culture and subsequently covered with a specific phage suspension, the organisms susceptible to the phage will be lysed and cause agar areas of no growth known as plaques.

S. aureus of human origin have been “typed” using this technique and classified into types of increasing pathogenicity. Bacteriophages are given identification numbers which also designate strains of Staphylococci related to them.

B. Gram Positive Facultative Anaerobic Cocci: Streptococci

Streptococci are facultative anaerobic saprophytes and parasites that normally occur on a wide variety of body surfaces. Because of their ability to ferment lactose, the Streptococci are also important agents of milk spoilage. Streptococci may be distinguished from Staphylococci as follows: (1) cocci are arranged in short chains or in pairs rather than as irregular clusters; (2) colonies exposed to the air are small, drop-like, mucoid, and unpigmented; (3) catalase is not produced; (4) NaCl concentration exceeding 6.5% is not tolerated; and (5) growth is not inhibited by sodium azide (Streptococci lack cytochromes).

With regard to differentiation of strains of Streptococci:

Pyogenic Streptococci such as S. pyogenes are β-hemolytic. β-Hemolytic Streptococci may become virulent pathogens under certain conditions, digesting deep tissues and destroying phagocytes. Some strains synthesize a hyaluronic acid which is identical to that in human connective tissue. The bacteria, therefore, are not coated with antibodies and are able to evade phagocytes. Most of the clinically important β-hemolytic Streptococci are in serologic Group A. Group A strains are uniformly sensitive to low concentrations (0.02 units) of bacitracin. Only a small percentage of other serologic groups are susceptible. In practice, a disc impregnated with the drug is placed on a lawn of bacteria. Group A Streptococci fail to grow around the disc where bacitracin has diffused.

Hemolytic Streptococci are typed according to two immunologically active components of the cell wall: the “M” protein and an extractable glucan, N-acetylglucosamine (the “Lancefield carbohydrate”). The Lancefield scheme (Table 4) is most widely used. Groups A, C, and G are implicated in a wide variety of infections. Group B Streptococci may be carried asymptptomatically by adults, but they can cause meningitis and other infections in the newborn. They may also be pathogenic in diabetic women, often as part of a syndrome with thromboembolism and necrosis of
the skin. Identification of Group A *Streptococci* can be confirmed by assaying serum for one of several streptococcal enzymes: streptolysin O, streptococcal hyaluronase, or streptococcal DNAse.

*Viridans Streptococci* are hemolytic. They cause partial hemolysis of red blood cells resulting in greening of colony surroundings. Some strains produce large polysaccharide capsules that probably have significance in pathogenesis. The capsules of *S. mutans* result in deposits around teeth (dental plaque) that enhance tooth decay. The capsules of *S. pneumoniae* (sometimes called diplococci) help it adhere to the epithelium of the lung. Because of the role of *S. pneumoniae* in lobar pneumonia, middle-ear infections, meningitis, and other infections, its diagnostic characteristics are clinically important. Pneumococci may form extremely mucoid colonies, are lysed by bile salts or by sodium deoxycholate, and are inhibited by the antibiotic optochin (ethylhydrocupreine hydrochloride). Biotypes of pneumococci are distinguished serologically. The polysaccharides making up the capsule have specific antigenic properties, and reaction with homologous antisera results in capsular swelling (Quellung reaction).

C. Gram-positive rods: Coryneform Bacteria

Typical *Corynebacteria* are small, Gram positive, club shaped rods arranged in V-shaped pairs, “Chinese letters”, or palisade arrangements. They are all nonmotile and aerobic. The distinctive arrangements of *Corynebacteria* may be attributed to the fact that the cell walls do not completely separate and the cells remain attached to each other at an angle. Other distinctive characteristics of *Corynebacteria* are bluish-purple metachromatic granules that are visible after staining cells with methylene blue, and the formation of black colonies on blood agar containing potassium tellurite. The granules are negatively charged particles of polyphosphate which absorb the methylene blue cations more strongly than other components of the cell. They are more visible in *Corynebacterium diphtheriae* than other *Corynebacteria*. The potentially pathogenic *C. diphtheriae* can be distinguished from nonpathogenic species on the basis of fermentation reactions: *C. diphtheriae* ferments maltose whereas other species do not. Virulence is generally verified by *in vitro* tests for toxin production such as the Elek virulence test. For this test, a strip or filter paper is soaked in *diphtheria* antitoxin then is embedded in an agar plate. Once the agar has hardened, the organism is streaked perpendicular to the filter paper. A positive test is indicated by a zone of precipitation.

D. Gram-Negative Cocci: The Family *Neisseriaceae*

The *Neisseriaceae* include four parasitic genera: *Neisseria, Branhamella, Moraxella* and *Acinetobacter*. All are Gram negative cocci or coccobacilli, nonspore forming, catalase positive and nonmotile. Members of the genus *Neisseria* are kidney shaped diplococci with flat adjacent sides. Aerobic or facultatively anaerobic, they are oxidase positive.

The Gram stain shows typical extra- or intracellular diplococci. In some instances, they stain very deep pink and in clinical specimens are present with a high number of polymorphonuclear cells. *Neisseria meningitidis* grows well on enriched media such as chocolate agar at 37°C, with 6% CO₂ and 50% humidity. After 18 hours of incubation on chocolate agar, colonies of *N. meningitidis* are round, grey and convex with a smooth moist glistening surface and an entire edge. The medium around the colonies may exhibit a greenish appearance.

The genus *Branhamella* contains one species important for human: *Branhamella catarrhalis*. Gram stain and colonies are identical to *N. meningitidis*. *B. catarrhalis* can be differentiated from *N. meningitidis* in the following way: *B. catarrhalis* grows at room temperature on blood or chocolate agar while *N. meningitidis* does not. In addition, *B. catarrhalis* does not ferment glucose, maltose, lactose, sucrose nor fructose while *N. meningitidis* ferments glucose and maltose.
E. Small Gram Negative Bacilli: The Genus *Haemophilus*

Bacterial species of genus *Haemophilus* require special growth factors that are found only in blood and other blood fluid. The name *Haemophilus* means “blood-loving” (Greek, *haemo* = blood; *philus* = loving). *Haemophilus influenzae* is the major disease-producing species of this genus. Bergey’s classification of bacteria lists *Haemophilus* among the facultative anaerobic Gram negative rods. *Haemophilus* are nonmotile, non-sporeforming bacteria surrounded by a capsule. Most of them are oxidase and catalase positive, ferment glucose and require additional factors such as Hemin (called Factor X) and/or NAD (called Factor V) to grow and develop colonies on trypticase soy agar. They grow well on chocolate agar since it contains both factors.

*Table 11.1.* Normal bacterial flora of the mouth.

<table>
<thead>
<tr>
<th>Microorganism</th>
<th>Range of Incidence (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Staphylococcus epidermidis</em> (coagulase negative)</td>
<td>75–100</td>
</tr>
<tr>
<td><em>Staphylococcus aureus</em> (coagulase positive)</td>
<td>Common</td>
</tr>
<tr>
<td><em>Streptococcus miis</em> and other α-hemolytic streptococci</td>
<td>100</td>
</tr>
<tr>
<td><em>Streptococcus salivarius, S. mutans, S. sanguis</em></td>
<td>100</td>
</tr>
<tr>
<td><em>Peptostreptococci</em></td>
<td>Prominent</td>
</tr>
<tr>
<td><em>Veillonella alcalescens</em></td>
<td>100</td>
</tr>
<tr>
<td><em>Lactobacilli</em></td>
<td>95</td>
</tr>
<tr>
<td><em>Actinomyces israeli</em></td>
<td>Common</td>
</tr>
<tr>
<td><em>Enterobacteriaceae</em></td>
<td>65</td>
</tr>
<tr>
<td><em>Haemophilus influenzae</em></td>
<td>25–100</td>
</tr>
<tr>
<td><em>Bacteroides fragilis</em></td>
<td>Common</td>
</tr>
<tr>
<td><em>Bacteroides melanogenicus</em></td>
<td>Common</td>
</tr>
<tr>
<td><em>Bacteroides oralis</em></td>
<td>Common</td>
</tr>
<tr>
<td><em>Fusobacterium nucleatum</em></td>
<td>15–90</td>
</tr>
<tr>
<td><em>Candida albicans</em></td>
<td>6–50</td>
</tr>
<tr>
<td><em>Treponema denticola, refringens, orale</em></td>
<td>Common</td>
</tr>
</tbody>
</table>

*Obligate anaerobe
**Table 11.2. Normal bacterial flora of the nasopharynx.**

<table>
<thead>
<tr>
<th>Microorganism</th>
<th>Range of Incidence (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Staphylococcus aureus</em></td>
<td>20–85</td>
</tr>
<tr>
<td><em>Staphylococcus epidermidis</em></td>
<td>90</td>
</tr>
<tr>
<td>Aerobic Corynebacteria (diphtheroids)</td>
<td>5–80</td>
</tr>
<tr>
<td><em>Streptococcus pneumoniae</em></td>
<td>0–17</td>
</tr>
<tr>
<td><em>Streptococcus pyogenes</em> (Group A)</td>
<td>0.1–5</td>
</tr>
<tr>
<td><em>Branhamella catarrhalis</em></td>
<td>12</td>
</tr>
<tr>
<td><em>Haemophilus influenzae</em></td>
<td>12</td>
</tr>
<tr>
<td>Gram negative bacteria (<em>Enterobacteriaceae</em>)</td>
<td>Rare</td>
</tr>
</tbody>
</table>

**Table 11.3. Normal bacterial flora of the oropharynx (throat).**

<table>
<thead>
<tr>
<th>Microorganism</th>
<th>Range of Incidence (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Staphylococcus aureus</em></td>
<td>35–40</td>
</tr>
<tr>
<td><em>Staphylococcus epidermidis</em></td>
<td>30–70</td>
</tr>
<tr>
<td>Aerobic corynebacteria (diphtheroids)</td>
<td>50–90</td>
</tr>
<tr>
<td><em>Streptococcus pyogenes</em> (Group A)</td>
<td>0–9</td>
</tr>
<tr>
<td><em>Streptococcus pneumoniae</em></td>
<td>0–50</td>
</tr>
<tr>
<td>α- and nonhemolytic Streptococci</td>
<td>25–99</td>
</tr>
<tr>
<td><em>Branhamella catarrhalis</em></td>
<td>10–97</td>
</tr>
<tr>
<td><em>Neisseria meningitidis</em></td>
<td>0–15</td>
</tr>
<tr>
<td><em>Haemophilus influenzae</em></td>
<td>5–20</td>
</tr>
<tr>
<td><em>Haemophilus parainfluenzae</em></td>
<td>20–35</td>
</tr>
</tbody>
</table>
Table 11.4. Lancefield groupings of streptococci.

<table>
<thead>
<tr>
<th>Group</th>
<th>Species</th>
<th>Hemolysis</th>
<th>Significance</th>
</tr>
</thead>
<tbody>
<tr>
<td>A</td>
<td><em>Streptococcus pyogenes</em></td>
<td>β</td>
<td>Important human diseases; group sensitive to penicillin</td>
</tr>
<tr>
<td>B</td>
<td><em>Streptococcus agalactiae</em></td>
<td>β (?)</td>
<td>Bovine mastitis</td>
</tr>
<tr>
<td>C</td>
<td><em>Streptococcus equi</em></td>
<td>β</td>
<td>Animal diseases; mild human</td>
</tr>
<tr>
<td></td>
<td><em>Streptococcus zooepidemicus</em></td>
<td>β</td>
<td>Respiratory infections</td>
</tr>
<tr>
<td></td>
<td><em>Streptococcus equisimilis</em></td>
<td>β</td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>Streptococcus dysgalactiae</em></td>
<td>–</td>
<td></td>
</tr>
<tr>
<td>D</td>
<td><em>Streptococcus faecalis</em></td>
<td></td>
<td>Enterococci; genitourinary tract infections, endocarditis, human wound infections, found in dairy products</td>
</tr>
<tr>
<td></td>
<td><em>Streptococcus faecalis subsp. liquefaciens</em></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>Streptococcus faecalis subsp. zymogenes</em></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td><em>Streptococcus faecium</em></td>
<td></td>
<td></td>
</tr>
<tr>
<td>E</td>
<td></td>
<td></td>
<td>Disease of swine, found in normal milk</td>
</tr>
<tr>
<td>F</td>
<td><em>Streptococcus minutus</em></td>
<td></td>
<td>Found in human respiratory tract</td>
</tr>
<tr>
<td>G</td>
<td><em>Streptococcus anginosus</em></td>
<td></td>
<td>Mild respiratory infections in human; genital infections in dogs</td>
</tr>
<tr>
<td>H</td>
<td><em>Streptococcus sanguis</em></td>
<td>α</td>
<td>Found in human respiratory tract</td>
</tr>
<tr>
<td>K</td>
<td><em>Streptococcus salivarius</em></td>
<td>α</td>
<td>Found in human respiratory tract</td>
</tr>
<tr>
<td>L</td>
<td></td>
<td></td>
<td>Genital tract infections in dogs</td>
</tr>
<tr>
<td>M</td>
<td></td>
<td></td>
<td>Genital tract infections in dogs</td>
</tr>
<tr>
<td>N</td>
<td><em>Streptococcus lactis</em></td>
<td>α</td>
<td>Lactic group; found in dairy products</td>
</tr>
<tr>
<td></td>
<td><em>Streptococcus cremoris</em></td>
<td></td>
<td>Viridans group; subacute bacterial endocarditis; found in human respiratory tract; Microaerophilic streptococci; Anaerobic streptococci–13 species</td>
</tr>
</tbody>
</table>
Materials

A. BSS
The formula for the 10^∞ stock of phosphate balanced salts solution (BSS) is:
\[\begin{align*}
K_2HPO_4 & \quad 0.40 \text{ g} \\
KH_2PO_4 & \quad 0.10 \text{ g} \\
KC1 & \quad 8.0 \text{ g} \\
\text{Distilled H}_2\text{O} & \quad 100 \text{ ml}
\end{align*}\]
Filter sterilize and store 4°C – 8°C

B. Bile Esculin Agar
1,000 ml H_2O
64 g bile esculin agar

C. Blood & Chocolate Blood Agar
1,000 ml H_2O
40 g T.S.A (Trypticase Soy Agar, see below).
Pour 570 ml of media into 1 liter flask with stir bar. Autoclave it then add:
–For chocolate blood: Add 30 ml of sheep blood into the media right after autoclaving.
–For Blood: Add 30 ml of sheep blood into the media when the media cool down to 50°C.

D. C.T.A. (cystine tryptic agar) with glucose, lactose, maltose or sucrose
1,000 ml H_2O
29.5 g C.T.A.
To each 1,000 ml of media, add:
–For glucose: 10 g glucose/liter
–For lactose: 10 g lactose/liter
–For maltose: 5 g maltose/liter
–For sucrose: 10 g sucrose/liter

E. Coagulase Plasma EDTA
Difco Laboratories
P.O. Box 331058
Detroit, Michigan 48232
1-800-521-0851

F. DNAse Test Agar
1,000 ml H_2O
42 g DNAse test agar

G. Gelatin Agar
1,000 ml distilled H_2O
8 g nutrient broth
30 g gelatin
15 g agar
H. ID2 Triplate and ID QUAD Plate
REMEL
P.O. Box 14428
12076 Santa Fe Drive
Lenexa, Kansas 66215
1-800-255-6730

I. Mannitol Salt Agar
1,000 ml H₂O
111 g mannitol salt agar

J. Nutrient Agar
1,000 ml H₂O
31 g nutrient agar
or:
8 g nutrient broth
15 g agar

K. 6.5% NaCl Broth
1,000 ml H₂O
65 g NaCl (sodium chloride)
25 g heart infusion broth
3 g glucose
5 ml bromocresol purple indicator (0.2 g bromocresol purple + 12.5 ml ETOH)

L. Trypticase Soy Agar (T.S.A.)
1,000 ml H₂O
30 g trypticase soy broth
15 g agar

M. Tellurite Chocolate Blood Agar
1,000 ml H₂O
20 g proteose peptone #3
2 g dextrose
5 g NaCl
13 g agar

Pour 540 ml of media into 1 liter flask, autoclave it, cool down to 50°C., then add:
30 ml of sheep blood
30 ml tellurite solution (1%)
Note for the Instructor

This experiment is mainly for students to learn the procedures of identifying each type of bacteria encountered in the upper respiratory tract. Due to the serious pathogenicity of some of the bacteria, such as *N. meningitidis*, *S. aureus* and *C. diphtheriae*, it is suggested to give students unknown species with no or minimal pathogenicity, such as *S. epidermidis*, *S. agalactiae*, *N. lactamica* and *H. parainfluenzae*. Students need to go through all the procedures in order to identify their unknown species. You have not given them pathogenic species, but they have learned the procedures for the identification.

Student Outline

Objective

The objective of this exercise is to become familiar with the diversity of the microorganisms residing in the mouth, nose and throat. You will attempt to identify potential pathogens by comparing them with known cultures, and performing several tests to determine various physical and physiological properties of the organisms. These will allow you to identify most of the genera encountered. You will also be introduced to simple, presumptive procedures for determining the presence of *Corynebacteria* and *Neisseriae*.

1. Observation of known organisms

A. Observe on the microscope, under 400∞ magnification as well as oil-immersion (1,000∞), stained slides of *Staphylococci*, *Streptococci*, *Branhamella*, *Neisseria*, and *Haemophilus*. For each of them, record in your notebook, all characteristics which would allow you to recognize them and differentiate them from each other, *i.e.*, their color, shape, arrangement, presence of granules, etc. Rinse slides thoroughly with xylene to remove immersion oil.

B. Observe each known culture under direct illumination and under the dissecting scope, and record their characteristics in your notebook. The objective of this part is to become familiar with the appearance and characteristics of several important groups of bacteria you will later encounter in unknown cultures.

2. Differentiation of major groups of bacteria based upon the observation of Gram stained organisms and the performance of biochemical tests

To become familiar with the use of simple biochemical procedures, I suggest that you perform each test using first a known species, then repeating it using your unknown and known organisms as positive or negative controls. The unknown may be either your URT isolate or one provided to you. These exercises will allow you to verify that manipulations are performed correctly and that they give the expected result.

A. Gram Staining

Choose two known and two unknown pure cultures. On each plate, look for a well isolated colony and locate it by circling and numbering the bottom of the plate around it. Gram stain a loopful of organisms, and observe them under oil immersion.
Procedure:

a. Cover a heat-fixed smear with crystal violet for one minute.
b. Briefly wash off the excess stain with water from the squeeze bottle after the one minute above.
c. Cover with Gram’s iodine for one minute.
d. Wash very briefly with water.
e. Hold the slide at a 45° angle. Gently pour alcohol over the slide and let it run off the slide just until it runs colorless. This is the critical step! Do not use the disinfecting alcohol for this.
f. Quickly wash with water again for a brief period.
g. Counter stain with safranin for one minute.
h. Wash briefly.
i. Carefully blot dry with paper towels. Be very careful not to press down on top of the bacteria. Simply let the paper towel absorb the excess water.
j. View the slide at 1000∞ magnification again.

B. Biochemical Testing

a. Differentiation of Staphylococci from other gram positive cocci, namely Streptococci, by means of the catalase test

Principle: Catalase is an enzyme that decomposes hydrogen peroxide (H₂O₂) into oxygen (O₂) and water (H₂O), bubbling out the oxygen. Excluding the Streptococci, most aerobic and facultative anaerobic bacteria possess catalase activity.

Procedures: Put a loopful of bacteria on a slide. Add a few drops of 3% hydrogen peroxide to the bacterial sample. The presence of catalase is shown by effervescence (a “fizzing” or bubbling of the liquid).

b. Differentiation of S.aureus from S.epidermidis

Further differentiation between two common staphylococcal species can be achieved by inoculating them into rabbit plasma, streaking on mannitol salt agar, gelatin slant and DNA slant. The results are shown in the following table.

Table 11.5. Differentiation of S. aureus from S. epidermidis.

<table>
<thead>
<tr>
<th>Test</th>
<th>S. aureus</th>
<th>S. epidermidis</th>
</tr>
</thead>
<tbody>
<tr>
<td>Coagulase</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>Acid from mannitol</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>Liquefies gelatin</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td>DNAse positive</td>
<td>+</td>
<td>–</td>
</tr>
</tbody>
</table>
**Gelatin Liquefaction.** In order to determine whether an organism produces gelatinase, a gelatin stab can be inoculated with the organism. A gelatin stab contains nutrient broth with 10–20% gelatin added. This amount of gelatin causes the material to “gel” or become semi-solid at temperatures below 25°C. The bacterial culture to be tested is “stabbed” into the nutrient gelatin tube with an inoculating needle. If the organism produces gelatinase, then the enzyme hydrolyzes the gelatin, and the gelatin subsequently loses its ability to gel.

**The DNAse Test.** DNA test agar contains DNA which can be precipitated by 1N HCL. Where an organism has degraded the DNA, no precipitation will occur. The result is a clear halo around the colonies on the agar. Other formulations use indicator dyes such as methyl green or toluidine blue in the medium. Methyl green is green when complexed with DNA and colorless when the DNA has been hydrolyzed. Toluidine blue is blue with the DNA and pink when it is hydrolyzed. The mannitol salt agar can differentiate between mannitol fermenters and non-fermenters. The fermenter produces acids and changes the PH indicator, phenol red, from red to yellow. The non-fermenter does not change the red color of the agar plate.

**The Coagulase Test:**

**Principle:** Coagulase is an enzyme converting fibrinogen into fibrin and resulting in the formation of a visible clot. The test can be performed either on a slide with organisms taken from an agar colony, or in a tube with cells from a broth culture.

**Procedure:**
- **Tube test:** Aseptically add 0.5 ml of rabbit plasma in a small test tube by means of a Pasteur pipette. With a 1 ml pipet, add 0.5 ml of a broth culture of the organisms. Mix gently (no vortex!) and place in a 37°C incubator for at least 4 hours. If positive, a clot will be present in the bottom. Never forget your controls, especially when working with an unknown, perform the test simultaneously with *S. aureus* (positive control) and *S. epidermidis* (negative control).

**c. Subsequent tests and observations to differentiate among Streptococci**

The pathogenic *Streptococci* grow readily in the laboratory, but the best development requires blood or serum media and body temperature, namely 37°C. Their colonies on blood agar reveal several types of hemolysis around them when examined by reflected light. α-hemolytic *Streptococci* leave a zone of partially lysed RBC (red blood cells) around the colony, surrounding it by a darker halo (brown or greenish). β-hemolytic *Streptococci* lyse all RBC around and are thus surrounded by a clear yellowish or transparent halo-hemolysis, represents the absence of hemolysis around the colony.

To distinguish between those different types of hemolysis, look at the plates either under the dissecting microscope where you may be able to see RBC in blood agar although it may be difficult to focus, or simply through the light microscope. Compare the tint of the halo: *S. pneumoniae* is α-hemolytic, *S. pyogenes*, β-hemolytic. Compare unknowns with control plates.

**d. Differentiation of Streptococcus pneumoniae by means of the bile solubility test**

**Principle:** Bile salts like sodium deoxycholate have the capability to selectively lyse *Streptococcus pneumoniae* when added to actively growing bacterial cells in a culture medium. The bile solubility test can be performed either with a broth culture (pH = 7.0) or with colonies on agar. Only the latter test will be explained here.

**Procedure:** To a well isolated colony of the test organism on blood agar, add a drop of 5% sodium deoxycholate and place the plate at 37°C for half an hour. Bile soluble colonies of
S. pneumoniae disappear leaving a hemolyzed area. Negative controls, which would be any other α-hemolytic Streptococci, will remain intact.

e. Differentiation of group A Streptococci from other serological types

All β-hemolytic Streptococci can be divided into 18 or more serological groups, (Table 11.4) called Lancefield groups (after their discoverer Rebecca Lancefield) on the basis of carbohydrates present in the cell wall. Most pathogenic β-hemolytic Streptococci belong to group A, among them S. pyogenes. Group B are for the most part harmless to humans, Group C and G are less pathogenic.

To distinguish Group A Streptococci from the other groups, antibiotic susceptibility tests are used. The procedure is to pick up cells from a colony we wish to identify and streak the entire surface of a plate to create a lawn of colonies. A disc impregnated with a particular antibiotic is then placed on the agar and the culture incubated for 24 hours at 37°C in the CO₂ incubator. The antibiotic contained in the paper disc will diffuse into the medium and if the bacteria are sensitive to it, a zone of no growth will appear around the disc. Group A Streptococci are sensitive to Bacitracin (labeled commercially here as Taxo-A discs), Streptococcus pneumoniae (which is α-hemolytic) is sensitive to Optochin (Taxo-P-discs). Streak a few plates with unknown organisms and with known controls. Place the two discs at a reasonable distance from each other. Press the disc gently on agar with a flamed forceps or a needle. Label your plates, incubate them at 37°C and check them tomorrow, or the day after. Remember to keep a record of your manipulations. Another way to differentiate S. pneumoniae is the bile solubility test explained before.

f. Identification of group B Streptococci by means of the CAMP test

A test allowing identification group B β-hemolytic Streptococci is called the CAMP test (an acronym of its four discoverers).

Principle: Group B Streptococci secrete a factor, the CAMP factor, which enhances the hemolytic activity of Staphylococcus aureus β-lysin on erythrocytes. Wherever the two reactants overlap on a sheep blood agar plate, accentuation of the β-hemolytic reaction occurs.

Procedure: On a blood agar plate, streak with the flamed needle a single line of Staphylococcus aureus. Flame the needle. Then pick up your unknown Streptococci and streak similarly a single line perpendicular to the previous one and with 5 mm distant from each other. The two lines must not touch each other. The plates are incubated at 37°C in CO₂ incubator for 24 hours. A positive test will show a triangular zone of enhanced clearing in the zone of closest proximity between the two streaks. Streptococcus agalactiae is the positive control.

g. Differentiation of group D Streptococci by means of Bile-esculin test

Principle: Bile-esculin (BE) test (for the identification of Group D Streptococci):

\[
\text{Group D} \\
\text{Bile-esculin medium} \rightarrow \text{Esculetin + Glucose} \\
\text{Streptococci} \\
\text{Esculetin + Ferric Citrate} \rightarrow \text{Black complex 174} \\
\text{(positive)}
\]

Procedure: Stab the bacteria into the bile-esculin slant with the flamed loop, then streak a zig-
zag line on the slant surface. The group D *Streptococci* will turn the slant into dark black. A positive BE test tells you that you have a group D *Streptococcus*; differentiation of the two types of group D *streptococci* depends on the salt-tolerance test.

3. Gram negative cocci: *Neisseria* and *Branhamella*

The upper respiratory tract may harbor various species of *Neisseria* which being morphologically similar, are distinguished by carbohydrate utilization. *Neisseria gonorrhoeae* normally parasitizes the genito-urinary tract but may be recovered from the URT in rare occasions. *Neisseria lactamitica* is a commensal. Both are to be distinguished from *Neisseria meningitidis* and *Branhamella catarrhalis*. All species develop small, opaque, gray-white colonies with a glistening surface after 24 to 48 hour incubation on chocolate agar in a moist atmosphere at 35–37°C in 5% CO₂.

*Exercise Procedure:*

a. Observe known colonies of *Neisseria sp.* and *Branhamella* in cultures and on Gram stained slides.

b. Take some unknowns from chocolate agar and try to identify possible *Neisseria* by observing colony morphology and performing a Gram stain.

c. After having recognized at least two *Neisseria* sp. try to identify the species using the Brown test or CTA media for carbohydrate utilization.

*Brown Test Procedure for Carbohydrate Utilization*

For each species to be identified, take 5 tubes with 0.1 ml balanced salt solution with 1% phenol red (BSS tubes), four solutions of carbohydrates namely: glucose, maltose, sucrose and lactose and pure colonies of unknown and known controls. Harvest a few loopfuls of colonies in one of the BSS tubes and mix well as to obtain a heavy suspension. Label the four other tubes for each carbohydrate. Add one drop of each respective carbohydrate to each BSS tube with a Pasteur pipet, using a different pipet for each tube, then one drop of bacterial suspension. Mix well. Close the tubes and incubate in a 35°C waterbath for 4 hours. Record the results on a chart. Positive tests in which the bacteria have utilized the sugar have turned yellow (acidic). Negative tests remain red-orange, the color of phenol red indicator at neutral pH. Compare your results with Table 11.6 below.

*Procedure for CTA Carbohydrate-Containing Media*

CTA (Cystine Tryptic Agar) is a semisolid basal medium used for fermentation studies with the addition of various carbohydrates. Also, its low agar content makes it well suited for motility studies. In this exercise, inoculate the following CTA carbohydrate-containing media immediately below the surface: glucose, lactose, maltose, and sucrose. Tighten lids and incubate without increasing CO₂. Examine daily up to five days or until acid production is seen. A positive result may be yellow color for acid production or diffuse growth for motility.
Table 11.6. Results of carbohydrate utilization tests.

<table>
<thead>
<tr>
<th>Species</th>
<th>Glucose</th>
<th>Maltose</th>
<th>Sucrose</th>
<th>Lactose</th>
</tr>
</thead>
<tbody>
<tr>
<td>N. gonorrhoeae</td>
<td>+</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>N. meningitidis</td>
<td>+</td>
<td>+</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>N. lactamica</td>
<td>+</td>
<td>+</td>
<td>–</td>
<td>+</td>
</tr>
<tr>
<td>B. catarrhalis</td>
<td>–</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
</tbody>
</table>

4. Gram positive rods, Corynebacterium diphtheriae

*Corynebacterium diphtheriae* are aerobic, non-sporeforming, nonmotile Gram positive rods with a typical microscopic appearance: the cells contain granules and frequently show club shaped swellings typical of the genus. Snapping divisions result in a typical palisade (picket-fence) and angular arrangement of the cells.

*Corynebacteria* grow on blood agar where they form tiny grayish colonies but are mostly isolated on highly selective media such as tellurite agar plates or Loeffler’s Serum Slants. Non-toxigenic *Corynebacterium diphtheriae* are used in our cultures. Observe the morphology of Gram stained *Corynebacteria* noting the club-shaped end, palisade arrangement (or “Chinese letters”) and possible granules.

*Exercise Procedure:*

Streak the bacteria on the blood tellurite agar plates and observe the colony phenotype.

5. Gram negative rods, *Haemophilus*

Identification of *Haemophilus* species by streaking the bacteria on ID2 Triplate and ID Quad Plate.

a. *ID2 Triplate*

Section 1 contains horse blood and will determine hemolytic activity of *Haemophilus*. Section 2 (chocolate agar) contains reagent necessary for the porphyrin test. Section 3 contains hemin.

b. *ID Quad Plate:*

The *Haemophilus* ID Quad Plate is intended to demonstrate growth factor requirements of species within genus *Haemophilus*. The plate contains Hemin, Beta-DPN (NAD), Horse Blood and Hemin-Beta DPN combination, each sectioned into various compartments. The combination of Hemin and Beta-DPN will speciate *Haemophilus* by means of “Growth” or “No Growth.” Horse Blood added to Quadrant Four will detect the hemolytic activity of the *Haemophilus* species.
# Table 11.7. Identification of *Haemophilus* species on ID2 Triplate.

<table>
<thead>
<tr>
<th>Organism</th>
<th>Hemolysis (sect. 1)</th>
<th>Porphyrin* (sect. 2)</th>
<th>Hemin (sect. 3)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>H. influenzae</em></td>
<td>none</td>
<td>no fluorescence</td>
<td>no growth</td>
</tr>
<tr>
<td><em>H. parainfluenzae</em></td>
<td>none</td>
<td>orange fluorescence</td>
<td>no growth</td>
</tr>
<tr>
<td><em>H. hemolyticus</em></td>
<td>beta</td>
<td>no fluorescence</td>
<td>no growth</td>
</tr>
<tr>
<td><em>H. parahemolyticus</em></td>
<td>beta</td>
<td>orange fluorescence</td>
<td>no growth</td>
</tr>
<tr>
<td><em>H. aphrophilus</em></td>
<td>none</td>
<td>orange fluorescence</td>
<td>growth</td>
</tr>
</tbody>
</table>

*Examine the plate under UV light*

# Table 11.8. Identification of *Haemophilus* species on ID Quad Plate.

<table>
<thead>
<tr>
<th>Organism</th>
<th>Growth in Quadrant</th>
<th>Growth Factor Requirements</th>
<th>Hemolysis (Horse Blood)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>I  II  III  IV</td>
<td>X-Factor (Hemin) V-Factor (β-DPN)</td>
<td></td>
</tr>
<tr>
<td><em>H. influenza</em></td>
<td>–  –  +  +</td>
<td>+</td>
<td>–</td>
</tr>
<tr>
<td><em>H. parainfluenza</em></td>
<td>–  +  +  +</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td><em>H. haemolyticus</em></td>
<td>–  –  +  +</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td><em>H. parahaemolyticus</em></td>
<td>–  +  +  +</td>
<td>–</td>
<td>+</td>
</tr>
<tr>
<td><em>H. aphrophilus</em></td>
<td>+  –  +  +</td>
<td>V</td>
<td>–</td>
</tr>
<tr>
<td><em>H. aegypticus</em></td>
<td>–  –  +  +</td>
<td>+</td>
<td>–</td>
</tr>
</tbody>
</table>
Microorganisms in URT

Literature Cited (arranged by diagnostic test)

1. **Bacitracin (Taxo-A) Inhibition Test.**

2. **The Bile Solubility Test.**

3. **The CAMP Test.**

4. **The CTA Carbohydrate Utilization Test.**

5. **The Coagulase Test.**

6. **Diagnostic Tests for Group A S. Pyogenes.**

7. **The DNAse Test.**

8. **Optochin (Taxo-P) Inhibition Test.**

9. **The Oxidase Test.**
APPENDIX A
A Flow Chart of Expected Results for Staphylococcal and Streptococcal Species

Throat or Nasal Sample

Gram+ Cocci

Catalase test

Staphylococcus

Catalase test

S. aureus

S. epidermidis

or others

Coalulase test

S. epidermidis

or others

S. bovis

or S. equinus

Tolerance 7% NaCl

Enterococci

S. mitis

S. sanguis

S. milleri

S. mutans

Optochin sensitive

S. pneumoniae

(3)

Viridans Group

S. mitis

S. sanguis

S. milleri

S. mutans

S. pneumoniae

(3)

Bile-esculin hydrolysis

S. pneumoniae

(3)

CAMP Test

(2)

Group A

S. pyogens

β hemolytic streptococci

Group B

S. agalactiae

Resistant

Sensitive

Resistant

Sensitive

Resistant

Sensitive

β

Bactracin sensitive

α

Hemolytic character

All other groups

(1) Not in respiratory tract

(2) or Hippurate hydrolysis

(3) Confirmed with the bile solubility test, S. pneumoniae is bile soluble