



مقدم من فريق أفق الطبي



ch3 - Growth of Bacteria

Bacteria → binary fission → process by one parent cell divides to form two progeny cells

exponential growth / logarithmic
 ex → N.O of cell → 1 2 4 8 16

after 4 generations

generation time or doubling time →

ex → escherichia coli 20 min

Mycobacterium 24h

3h 30 min

rapid production of very large numbers of bacteria

short doubling time

① amount of nutrients ② T, pH ③ environmental factors / doubling time

Growth Curve Stage

① lag phase → no divide / metabolic activity

② log phase → rapid cell division
 ex → penicillin / cephalosporin / cell wall

③ stationary phase → no of cells new → no of cells that die
 steady state / nutrient depletion / toxic product → growth to slow

④ death phase → decline in the N.O of viable bacteria
 LAG / stationary / Death / time

Aerobic and anaerobic growth

most organisms

adequate supply of O_2 metabolism + growth

oxygen $\xrightarrow{\text{flavoproteins}}$ hydrogen receptor (final step O_2)

flavoproteins

energy coupled

H_2O_2 hydrogen peroxide + oxygen $\rightarrow$ free radical superoxide O_2^- toxic

superoxide dismutase

water + $H_2O_2 \Rightarrow$ catalyzes $2H_2O_2 \rightarrow 2H_2O + O_2$

O_2 requirements

① ex $\Rightarrow$ M. tuberculosis $\Rightarrow$ obligate aerobes $\Rightarrow$ hydrogen receptor $O_2 \xrightarrow{\text{ATP-generating system}}$ growth in O_2

② E. coli $\Rightarrow$ facultative anaerobes $\Rightarrow$ respiration $\Rightarrow$ ATP gain O_2 plus ATP gain fermentation $\Rightarrow$ grow O_2 or no O_2

③ obligate anaerobes

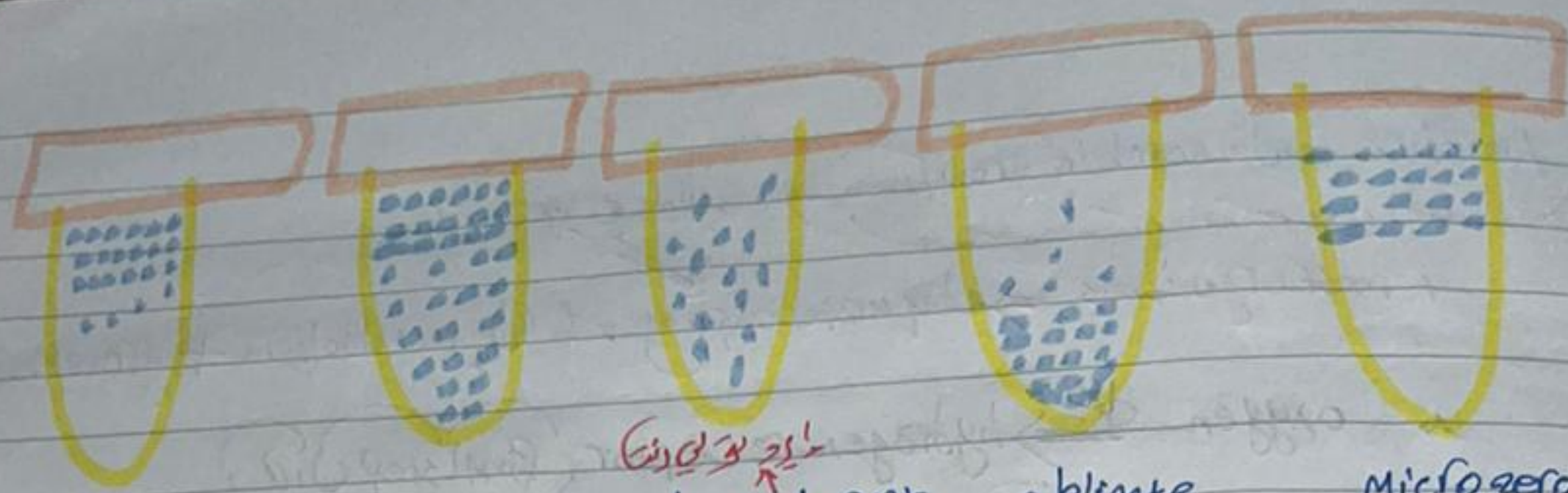
cannot grow in presence O_2
superoxide dismutase, catalase

obligate anaerobe vary in response to O_2 exposure

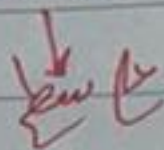
killed rapidly

survive

but not able to grow



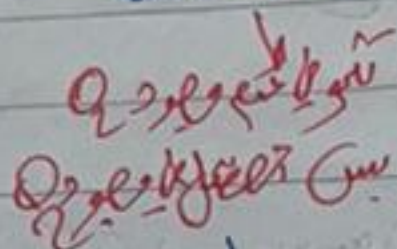
obligate
aerobe



high conc of O_2

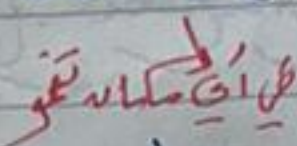
presence of
enz superoxide
dismutase (SOD) +
catalase

facultative
anaerobe



best where most
 O_2 is present

Aerotolerant
anaerobe

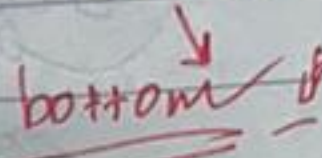


Growth occurs
evenly, O_2 has no effect

presence of some
enzyme

allows harmful forms of O_2
to be partially neutralized

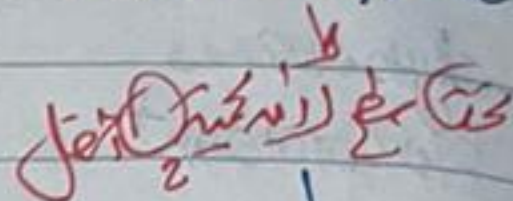
obligate
anaerobe



growth when
no O_2

lacks of enzymes

microaerophile



Growth when
low conc of O_2

produce lethal
amounts of toxic

fermentation of sugars

identification, classification, differentiation

ex $\Rightarrow$ Neisseria gonorrhoeae, Neisseria meningitidis

maltose, glucose fermentation are used to distinguish

meningitides $\Leftarrow$ fermentation of maltose, glucose, & neisseria

$\times$ E. coli $\Rightarrow$ fermentation of lactose

$\times$ salmonella & shigella $\Rightarrow$ non-lactose fermentation

glucose
fermentation

(glycolytic)

def $\Rightarrow$ Break down of sugar to pyruvic acid, $\xrightarrow{\text{w/o O}_2}$ lactic acid
mono saccharide $\Rightarrow$ glucose / maltose
galactose.

(note) lactose $\Rightarrow$ disaccharide $\sim$ glucose + galactose

$\xrightarrow{\text{E. coli } \beta\text{-galactosidase}}$ cleaved

process by facultative bacteria generate ATP
 $\downarrow$
grow faster in the presence of O_2

facultative + anaerobic $\Rightarrow$ fermentation, la / aerobic $\Rightarrow$ fermentation

deamination $\Rightarrow$ (Pseudomonas) $\Rightarrow$ c.a.
Alkaline PH $\downarrow$

How to PH indicator phenol red, $\Rightarrow$ red $\Rightarrow$ acid, yellow $\Rightarrow$ neutral, pink $\Rightarrow$ basic

(fermentation) $\Rightarrow$ red $\Rightarrow$ fermentation

fermentation $\Rightarrow$ clinical laboratory
 $\Rightarrow$ pyruvate + lactate acid turns the medium acid

iron metabolism

important for production of enzymes
+ growth factor

iron in form of ferric ion $\Rightarrow$ growth ~~not~~

- iron mostly sequestered ^{by} transferrin
 $\rightarrow$ ~~regulation~~

- siderophores ^{new} $\Rightarrow$ capture iron (chelating agent)
~~release~~ $\rightarrow$ ~~transferrin~~

ch-4

Genetics

single circular DNA
weight $\rightarrow 2 \times 10^9$
base pairs $\rightarrow 5 \times 10^6$

escherichia coli

human

MW 5×10^8 (mycoplasma) $\leftarrow$ DNA of the smallest

cell wall

1000 genes $\rightarrow$ 1000 protein
+ base pairs 3×10^9

two sets of chromosomes

(2n)

bacteria $\rightarrow$ haploid / eukaryotic $\rightarrow$ diploid

recessive / dominant

haploid $\rightarrow$ mutation $\rightarrow$ not expressed $\rightarrow$ cell lost that trait.

one set of chromosomes

Def $\rightarrow$ mutation $\rightarrow$ change in the base sequence of DNA $\rightarrow$ insertion of different amino acid into a protein, appearance of a altered phenotype

mutations result from three types of molecular changes.

Base substitution $\rightarrow$ base for base / insertion / deletion
DNA replication $\rightarrow$ "one base is inserted in place of another"

base / C/G hydrogen bonding $\rightarrow$ error $\rightarrow$ DNA polymerase $\rightarrow$ wrong base is inserted

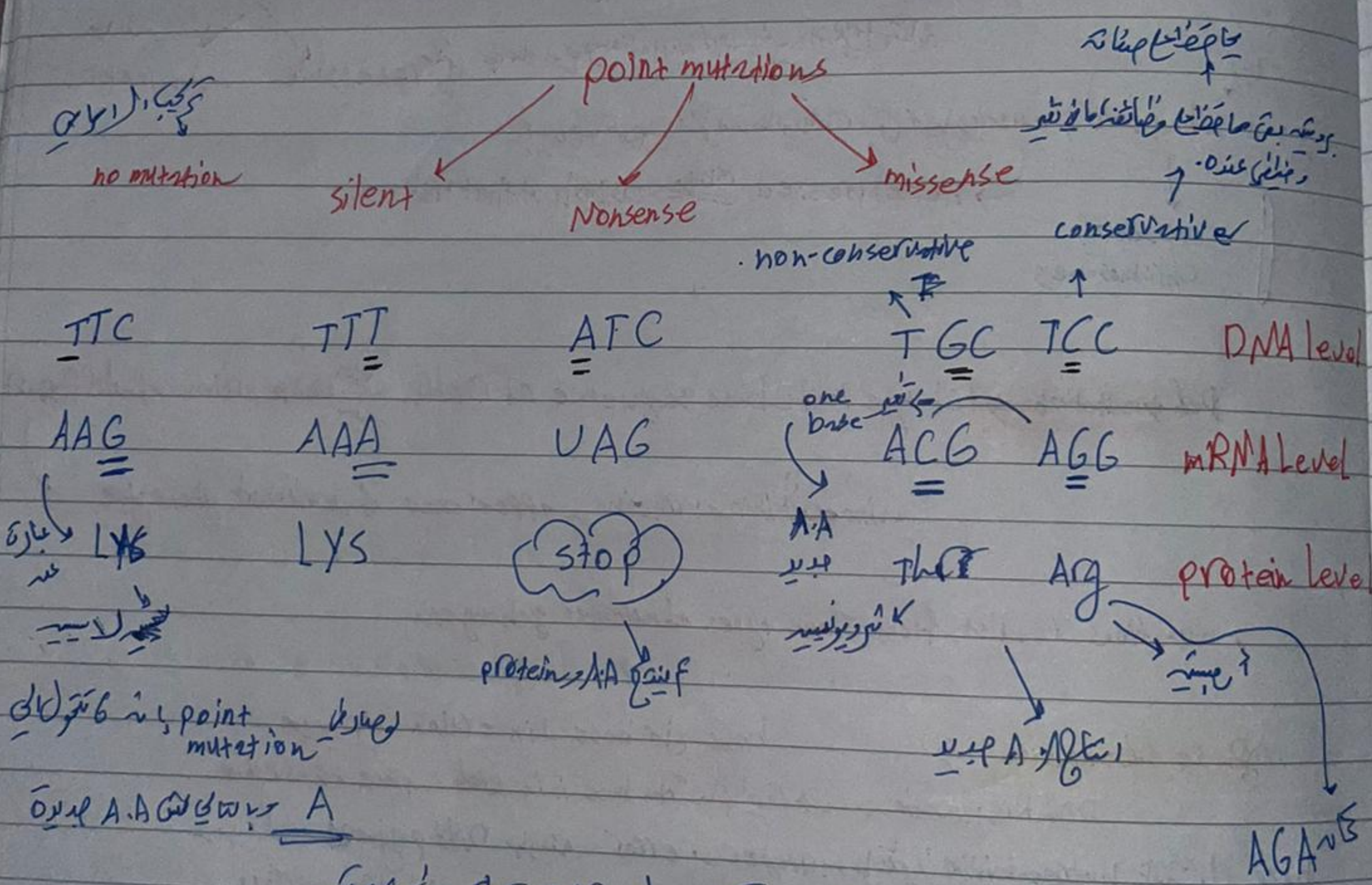
missense mutation $\rightarrow$ base substitution $\rightarrow$ code that simply cause a different A.A to be inserted
A.A $\rightarrow$ codes the protein

UAA UAG UGA
= stop codon

non sense mutation → when a base substitution generates termination codon stop protein synthesis

silent mutation → mutation that does not change the amino acid sequence

GUV → CAA (VAL)
GUA ← CAT → (VAL)



missense mutation → when a base substitution generates a different amino acid

2) frame shift mutation

when one or more base pairs are added or deleted
 modify the base sequence

shifts reading frame on the ribosome
 incorporation of the wrong A.A. "downstream"

from the mutation + production of inactive protein

TAA CTG CAG GT → original sequence

TAA COT GCA GGT insertion

insertion of

extra A.A. at this codon

3) transposons

jumping between pieces of DNA in cells between bacterial cells, mice

transposons or insertion sequences are integrated in to the DNA

insertion of genes is profound change, new insertion next adjacent gene

mutations can be caused by chemicals/radiation/viruses

① nitrous acid + alkylating agent $\rightarrow$ hydrogen bond
Ochale existing base
cytosine pairs with thymine pairs with adenine $\leftarrow$ wrong base $\leftarrow$ preferentially pair

② 5-bromouracil $\rightarrow$ base analogue $\rightarrow$ thymine $\rightarrow$ resemble normal base

5-bromouracil $\rightarrow$ methyl group $\rightarrow$ polar radius $\rightarrow$ bromine atom

5-methyluracil $\leftarrow$ thymine $\rightarrow$ insertion mutation

note $\rightarrow$ 5-bromouracil $\rightarrow$ thymine

hydrogen bonding fidelity

guanine

$\rightarrow$ A-T base pair $\rightarrow$ G-C base pair

antiviral drug $\rightarrow$ mutation

③ benZpyrene $\rightarrow$ tobacco smoke

frameshift mutations $\rightarrow$ existing DNA bases

mutagens, carcinogens

mutation

intercalate

intercalate between the adjacent base $\rightarrow$ DNA sequence

- X-rays have high energy and can damage DNA in three ways

- ① break down the covalent bond
DNA $\rightarrow$ ribose P $\rightarrow$ bases
- ② bases $\rightarrow$ free radical $\rightarrow$ electron
- ③ hydrogen bonding $\rightarrow$ bases $\rightarrow$ electron

- Ultraviolet radiation

thymine dimer
ribosome (thymine is dimerisation) $\rightarrow$ DNA $\rightarrow$ inability of the DNA to replicate properly

- mutator bacteriophage $\rightarrow$ mutation

bacteriophage $\rightarrow$ mutation

binary diffusion $\rightarrow$ bacterial chromosome $\rightarrow$ insertion $\rightarrow$ DNA

mutation $\rightarrow$ DNA $\rightarrow$ mutation $\rightarrow$ mutation $\rightarrow$ mutation

frameshift or deletions

4

transfer of DNA within bacterial cells.

transposons

↑ transfer DNA from one site on the bacterial chromosome to be another site or to a plasmid

→ inserted in bacterial chromosome or plasmid

medically important antibiotic resistance

between bacterial cells within bacterial cells

conjugation

transposons

mechanism

copy

paste

transfer of DNA within bacteria

programmed rearrangements

programmed rearrangements

→ Defect → movement of genes from silent storage site to active expression site.

→ e.g. → nisseria gonorrhoea / Borrellia recurrentis

→ Benefit → evade immune response.

Transfer of DNA between bacterial cells.

note → most important consequence of DNA transfer

antibiotic resistance

Def

[1] conjugation

⇒ mating of 2 bacterial cells in which DNA is transferred from donor to recipient.

→ **requirements** → f factor in the donor (f⁺)
~~receptor~~ in the recipient (f⁻) receptor.

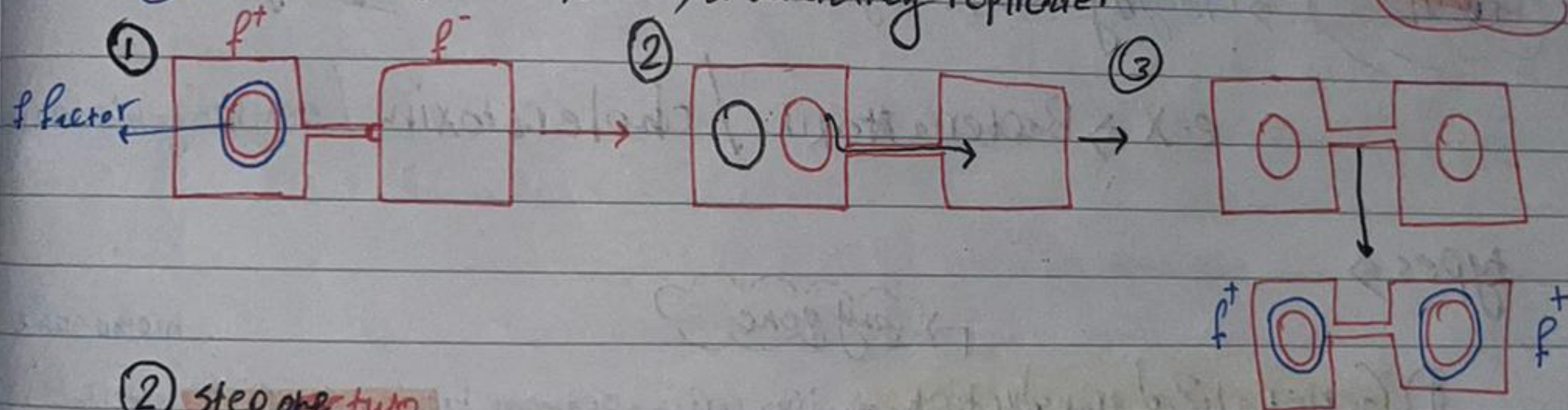
→ **f factor** → genes → proteins required for conjugation such as **pilin** (sex pili)

→ **types** → plasmid transfer (1) (2) bacterial chromosomal DNA transfer

Mechanisms for:

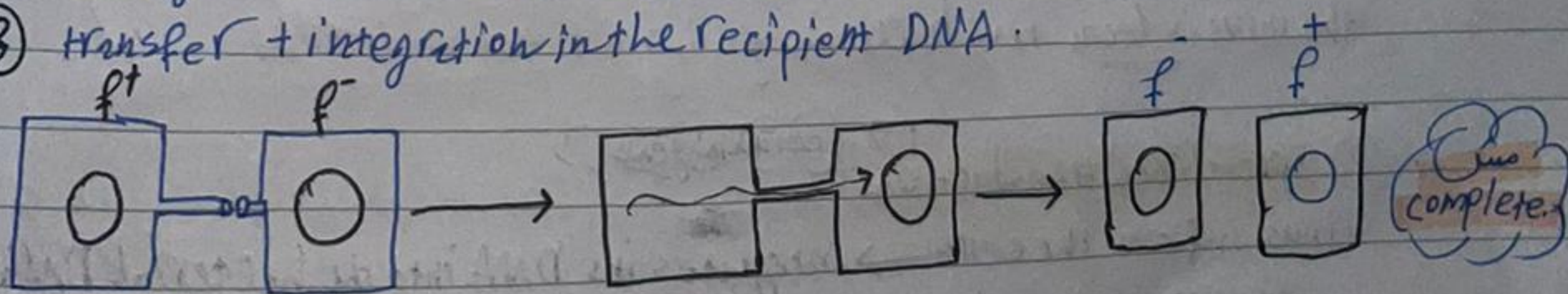
step one

- ① the pilus of f⁺ attaches to the receptor of f⁻
- ② enzymatic cleavage of the factor
- ③ one strand is transferred, and then they replicate.



② step one two

- ① pilus of f⁺ attaches to the receptor of f⁻
- ② cleavage of bacterial chromosome.
- ③ transfer + integration in the recipient DNA.



transfer of DNA (1) (2) (3) step two, step one

type of cell involved (prokaryotic)

note → The time for complete transfer = 100 min → ② only
→ usually the conjugation is broken and only part of the chromosome is transferred
→ newly acquired DNA integrates into the recipient's chromosome DNA, becoming part of genetic material
→ the most frequently transferred gene is the one adjacent to the oriT piece.
- Donor is first

② Transduction → Def: transfer of cell DNA by bacterial virus

mechanism: ① during viral growth, bacterial DNA (part) is incorporated into the virus

② in the recipient cell, viral DNA is integrated into bacterial DNA giving it new traits (lysogenic)

result → changing non pathogenic bacteria into pathogenic ones.

e.g. → Bacterial toxin / cholera toxin / erythrogenic toxin

types →

① Generalized transduction → any gene
bacterial DNA is fragmented → virus carries a segment from any part of bacterial chromosome
→ a fragment is incorporated into virus
the virus infects new bacteria giving it the new traits

② specialized transduction → certain gene
virus infects the cell → integrates its DNA into the bacterial DNA
excises DNA and a part of the adjacent bacterial DNA
→ infects new cell giving it new traits.

* type of cell involved (prokaryotic)

note $\Rightarrow$ some virus integrate at specific sites of bacterial DNA, so the adjacent genes that are transferred are specific to that virus.

③ transformation, transfer of DNA one cell to another

Methods $\rightarrow$ natural $\rightarrow$ ^{lysogenic / lytic} the dying cell release its DNA, other recipient cells take it up. ^{DNA $\rightarrow$ ϕ}

$\rightarrow$ Laboratory $\rightarrow$ ^{lysogenic} extract DNA from one type and introduce it to other types. ^{transfection} ^{ins (or) me} human ^{lysogenic}

transformation of DNA from encapsulated to Non-encapsulated
 $\rightarrow$ (Non-encapsulated $\rightarrow$ encapsulated)
non-pathogenic $\rightarrow$ pathogenic

Recombination $\rightarrow$ integration of the transferred DNA to the host DNA
 $\rightarrow$ requirements: ligases + endonucleases.

Types

DNA $\downarrow$ ϕ

① homologous: 2 pieces have extensive homologous regions pair up
exchanging piece $\Rightarrow$ breakage + reunion (note $\rightarrow$)

② Non-homologous: little if any, homology is necessary

type of cell involved (prokaryotic + eukaryotic)

Nature of DNA transferred (Any DNA)

ch-6 | Classification of medically important bacteria.

④ principles of classification.

① Cultural characteristics

③ Biochemical

② Morphology + staining

④ Molecular

⑤ Serological.

* nature of cell wall

* Thick, rigid cell wall

/ flexible thin cell wall

/ wall-less cells

ex: spirochete

① Thick cell wall.

→ extracellular bacteria

① free living bacteria → can live outside cells in laboratory media

ex: human + animal cells

② non-free living bacteria → requires cells to survive (obligate intracellular parasites)

ex: Chlamydia / Rickettsia

subdivided according to shape + staining

into

G^{-ve} cocci

- G^{ve} cocci

G^{ve} Bacilli

G^{ve} Bacilli

rods

③ have different O₂ requirements.

spore forming abilities

* G^{ve} Cocci

Arrangement

→ pneumonia / pharyngitis

→ streptococci

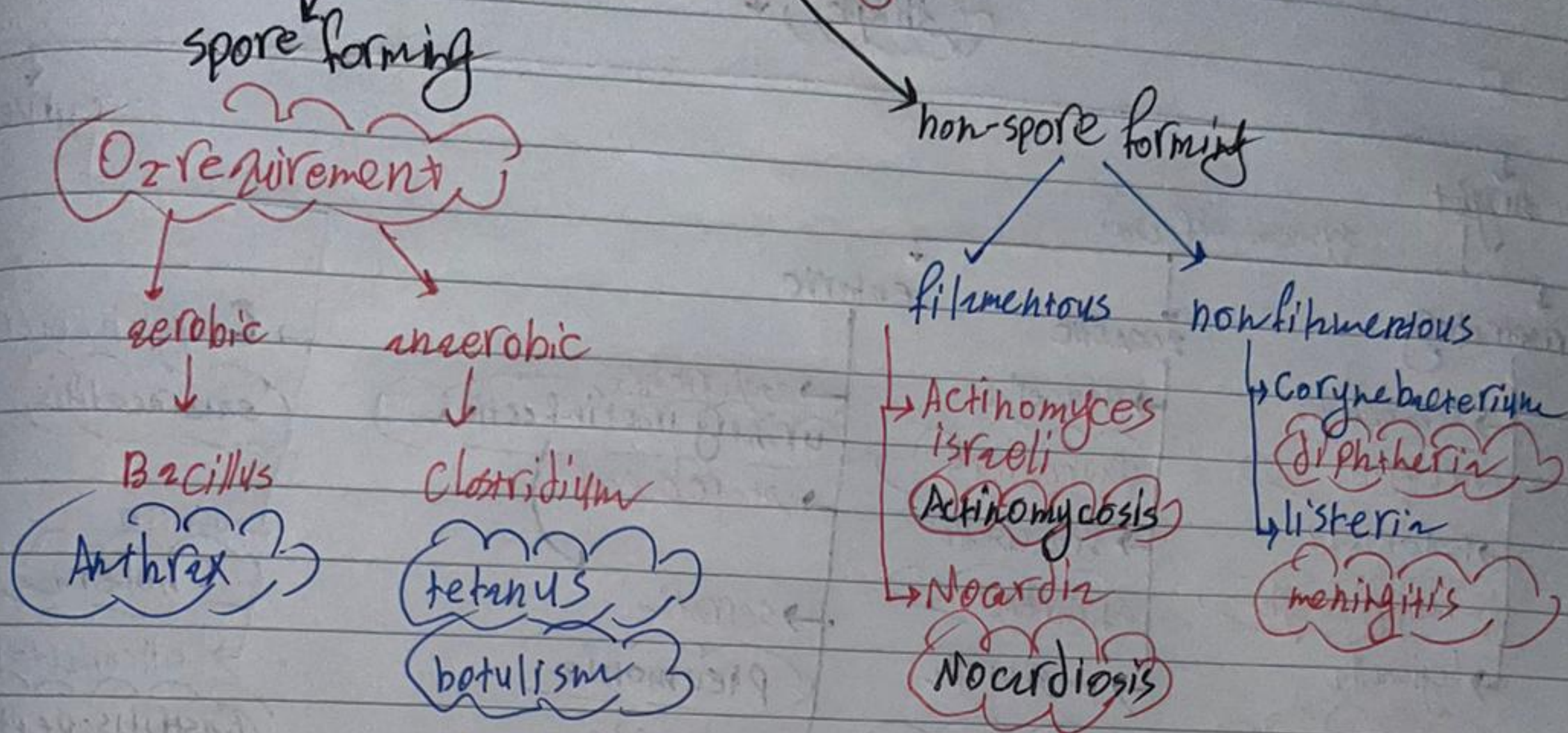
→ staphylococci

Catalase^{-ve}

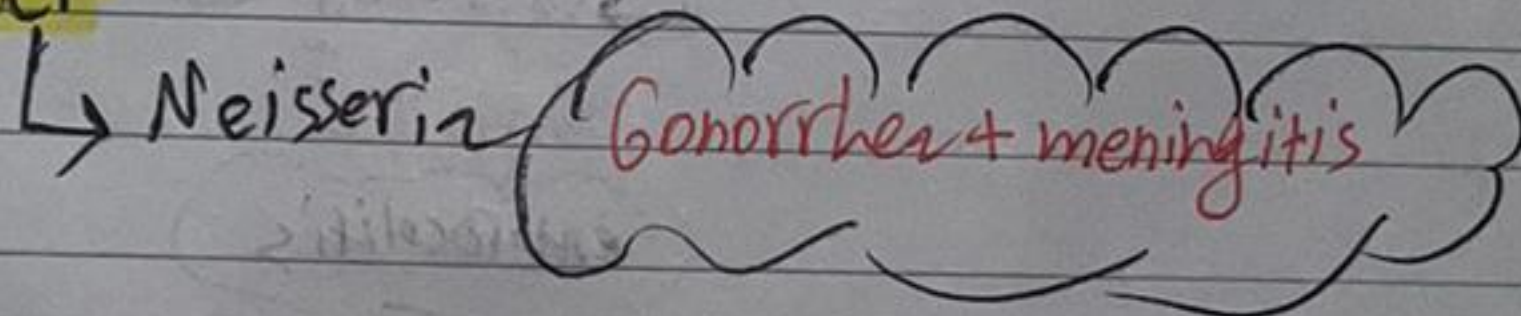
Catalase^{ve}

→ Abscess of skin

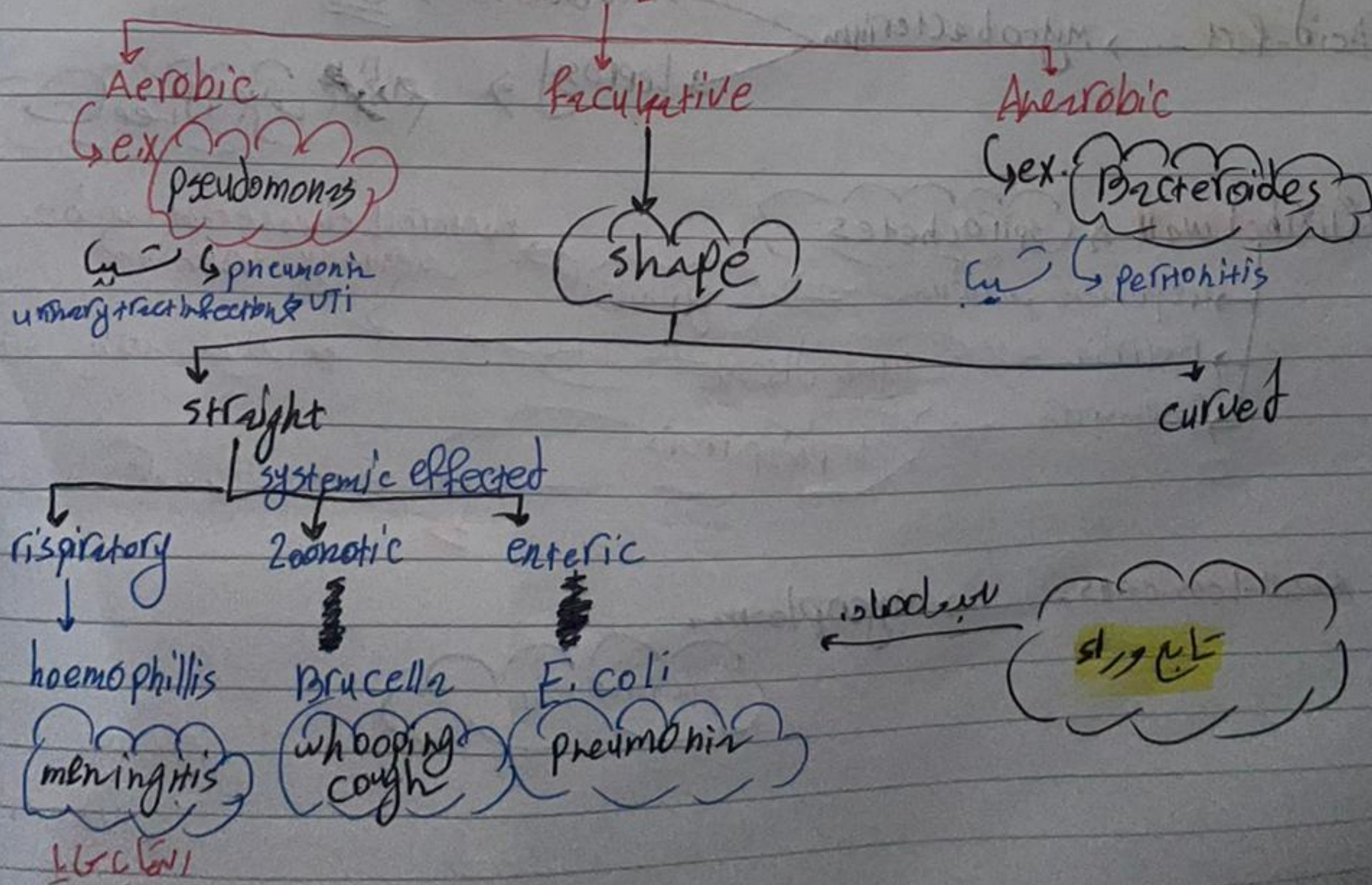
* G⁺ bacilli → spore-forming ability

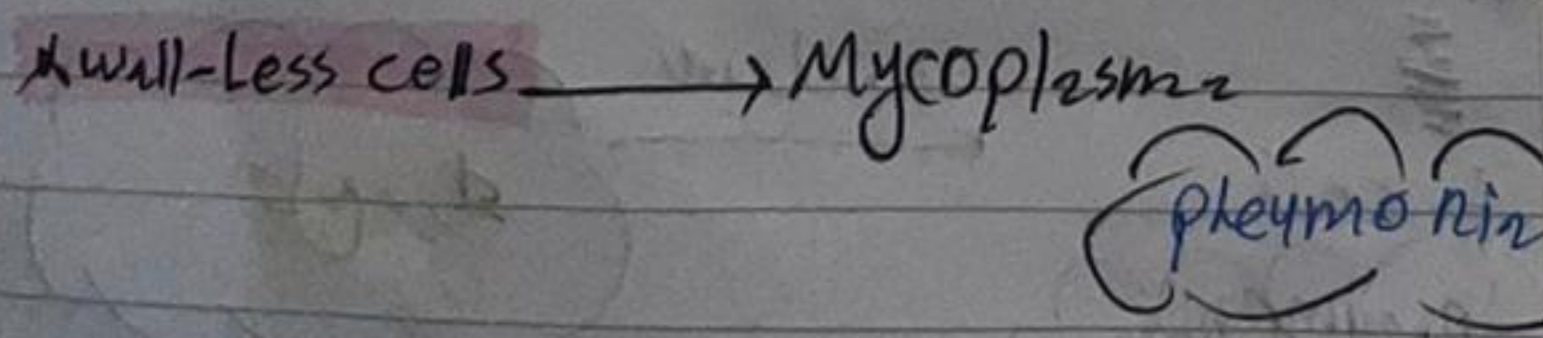
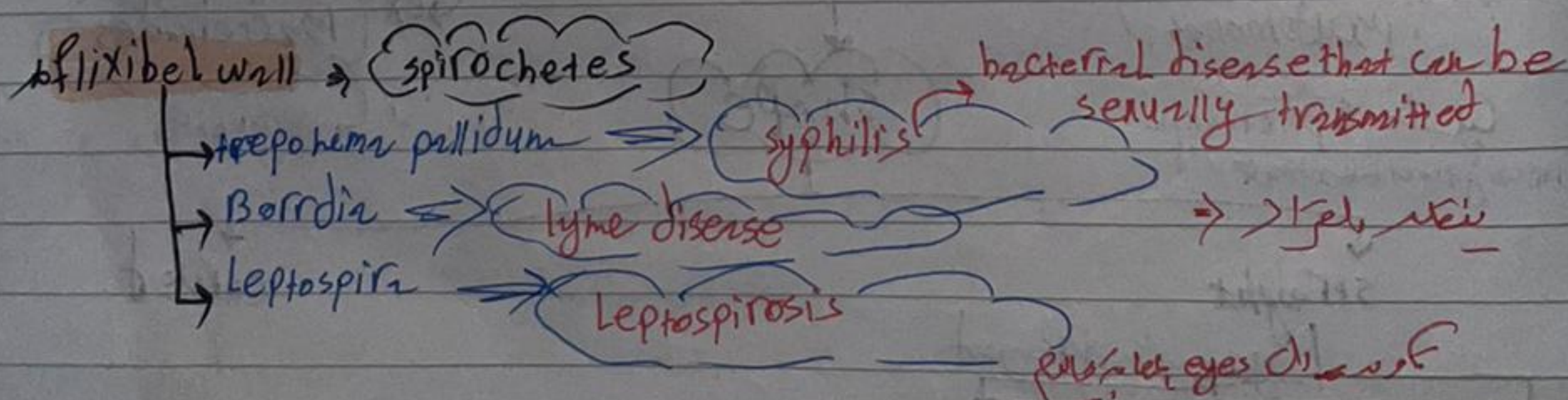
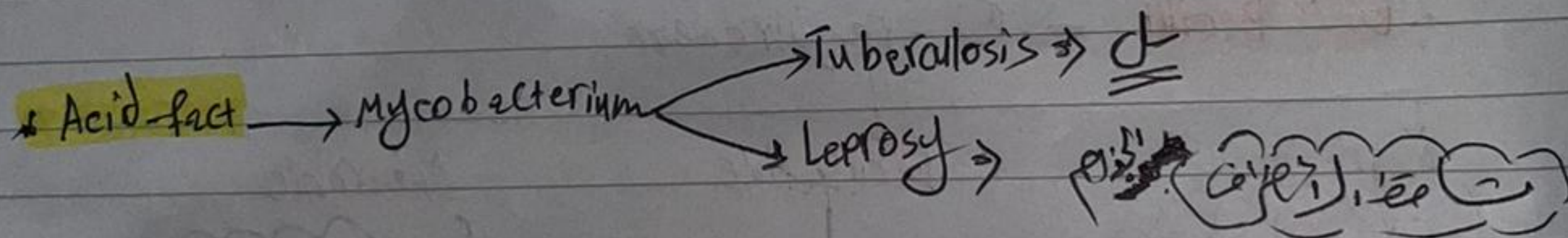
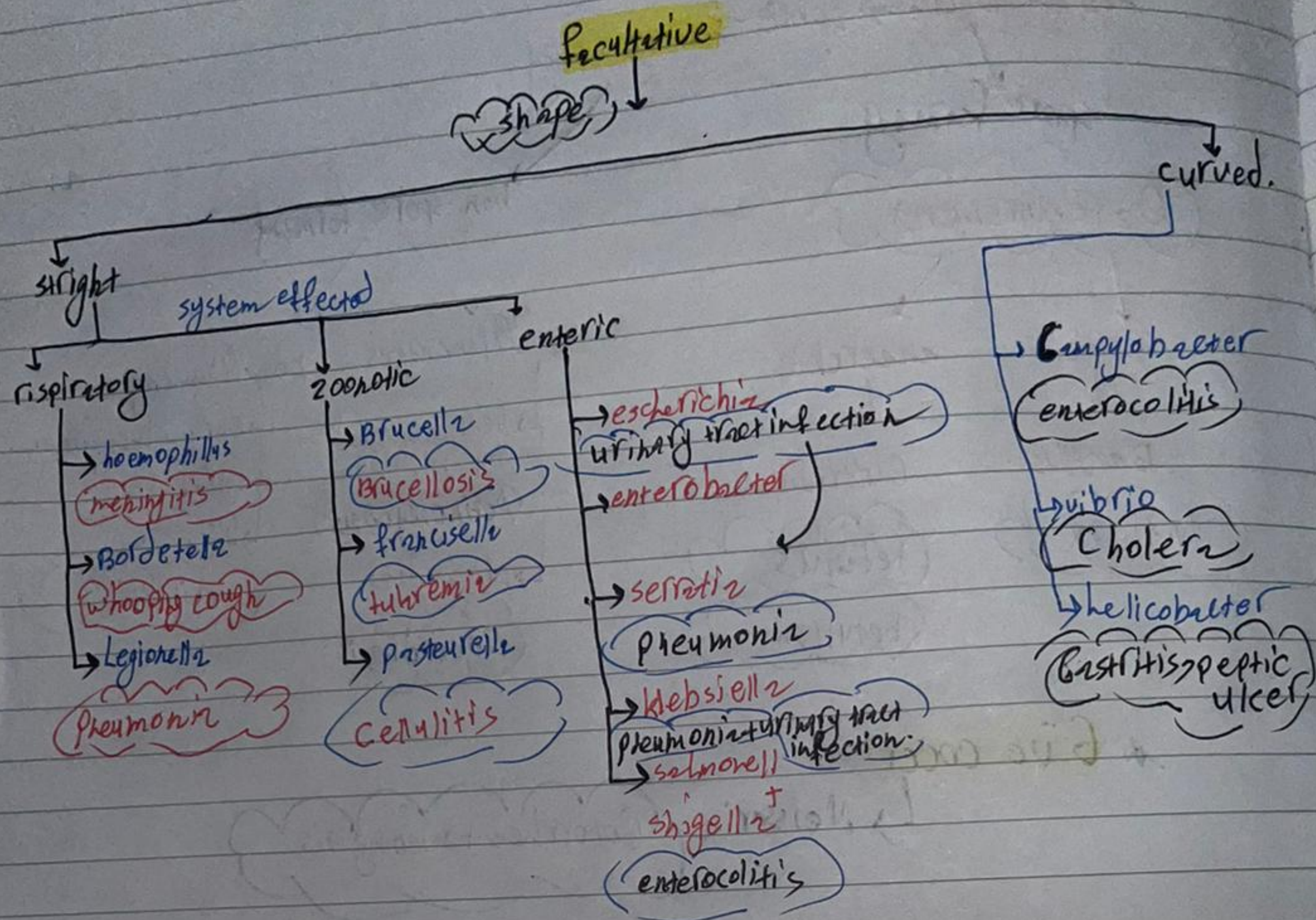


* G⁻ cocci



* G⁻ Bacilli → O₂ - requirement





Genetic test

↓

isnt

Pseudomonas from the Burkholderia family because the difference between 2 DNA

↓

DNA fingerprinting molecular test.

Chapter 6

Normal Flora

AP: Dr. Said S. Algora



: Concept of normal Flora

مفهوم الفلورا الطبيعية

Normal flora is the term used to describe the various bacteria and fungi that are permanent resident of certain body sites, especially the skin, oropharynx, colon, and vagina

→ not considered (normal flora)

The viruses and parasites, which are the two other major groups of microorganisms, are usually not considered members of the normal flora, although they can be present in asymptomatic individuals

The normal flora organisms are often referred to as commensals, which are organisms that derive benefit from another host but do not damage that host

normal flora = commensals

مستعمرات غير ضارة

: Contents

- concepts of normal Flora-
- normal Flora of the skin-
- normal flora of the respiratory tract-
- normal flora of the intestinal tract-
- normal flora of the Genitourinary tract-

TABLE 6-1 Summary of the Members of Normal Flora and Their Anatomic Locations

Members of the Normal Flora	Anatomic Location
Bacteroides species	Colon, throat, vagina
Candida albicans	Mouth, colon, vagina
Clostridium species	Colon
Corynebacterium species (diphtheroids)	Nasopharynx, skin, vagina
Enterococcus faecalis	Colon
Escherichia coli and other coliforms	Colon, vagina, outer urethra
Gardnerella vaginalis	Vagina
Haemophilus species	Nasopharynx, conjunctiva
Lactobacillus species	Mouth, colon, vagina
Neisseria species	Mouth, nasopharynx
Propionibacterium acnes	Skin
Pseudomonas aeruginosa	Colon, skin
Staphylococcus aureus	Nose, skin
Staphylococcus epidermidis	Skin, nose, mouth, vagina, urethra
Viridans streptococci	Mouth, nasopharynx

UT 1
أكثر شيوعاً في مجرى البول

The members of the normal flora vary in both
 Although number and kind from one site to another.
 of the normal flora extensively populates many areas
 the body, **the internal organs usually are sterile**.
 Areas such as the central nervous system, blood,
 lower bronchi and alveoli, liver, spleen, kidneys,
 and bladder are free of all but the occasional
 . transient organism

*note → normal flora
 skin ③ urogenital tract ② GIT ① ↓
 note → sterile system divided normal flora (e.g. as
 Leix → CNS, blood.

There is also a distinction to be made between
 members of the normal flora, which are the
 permanent residents, and the colonization of the
 individual with a new organism. In a sense, we are
 all colonized by the normal flora organisms, but the
 term colonization typically refers to the acquisition
 of a new organism. After the new organism colonize
 (i.e., attaches and grow, usually on a mucosal
 membrane), it may cause an infectious disease or it
 may be eliminated by our host defenses.

Furthermore, the person colonized by a new
 organism can transmit that organism to others (i.e.,
 . act as a reservoir of infection for others)
Colonization → Acquisition of new organism that may cause disease or be
 eliminated or transmitted to others.
 pathogenesis. **dissemination** → the carrying of new organism

There is a distinction between the presence of these
 organisms and the carrier state. In a sense, we all
 are carriers of microorganisms, but that is not the
 normal use of the term in the medical context. The
 term carrier implies that an individual harbors a
 potential pathogen and therefore can be a source of
 . infection of others
 + Can transmit infections.
 infection → potential pathogen
 ↳ PD → chronic carrier.

It is most frequently used in reference to a person
 with an asymptomatic infection or to someone who
 has recovered from a disease but continues to carry
 . the organism and may shed it for a long period

*notes → carrier **asymptomatic** infection
 *note → carrier state → asymptomatic infection
 ↳ example: hepatitis B

The members of the normal flora play a role both in
the maintenance of health and in the causation of
disease in three significant ways
 ↳ E. coli in urinary tract → UTI
 ↳ immune system weakness
They can cause disease, especially in (1) → change their location
 immunocompromised and debilitated individuals.
 Although these organisms are nonpathogens in their
 usual anatomic location, they can be pathogens in
 . other parts of the body

*note → colonization as normal flora
 ↳ example: E. coli

The nonpathogenic resident bacteria occupy attachment sites on the skin and mucosa that can interfere with colonization by pathogenic bacteria. The ability of members of the normal flora to limit the growth of pathogens is called colonization resistance.

If the normal flora is suppressed, pathogens may grow and cause disease. For example, antibiotics can reduce the normal colonic flora that allows *Clostridium difficile*, which is resistant to the antibiotics, to overgrow and

[illegible]

Location	Important Organisms ¹	Less Important Organisms ²
Skin	<i>Staphylococcus epidermidis</i>	<i>Staphylococcus aureus</i> , <i>Corynebacterium diptheroides</i> , various streptococci, <i>Pseudomonas aeruginosa</i> , anaerobes (e.g., <i>Propionibacterium</i>), yeasts (e.g., <i>Candida albicans</i>)
Nose	<i>Staphylococcus aureus</i> ¹	<i>S. epidermidis</i> , <i>Corynebacterium diptheroides</i> , various streptococci
Mouth	Various streptococci	Various streptococci, <i>Elizabethkingella</i>
Dental plaque	<i>Streptococcus mutans</i>	<i>Prevotella intermedia</i> , <i>Porphyromonas gingivitis</i>
Gingival crevices	Various anaerobes (e.g., <i>Bacteroides</i> , <i>Fusobacterium</i> , streptococci, <i>Actinomyces</i>)	
Throat	<i>Viridans streptococci</i>	Various streptococci (including <i>Streptococcus pyogenes</i> and <i>Streptococcus pneumoniae</i>), <i>Moraxella</i> species, <i>Neisseria meningitidis</i> , <i>S. epidermidis</i> , <i>Staphylococcus</i> , <i>Escherichia coli</i>
Colon	<i>Bacteroides fragilis</i> , <i>Escherichia coli</i>	<i>Streptococcus</i> , <i>Escherichia coli</i> , <i>Enterococcus faecalis</i> and other streptococci, <i>Clostridium</i>
Vagina	<i>Lactobacillus</i> , <i>E. coli</i> , group B streptococci ¹	Various streptococci, various gram-negative rods, <i>B. fragilis</i> , <i>Corynebacterium diptheroides</i> , <i>C. albicans</i>
Urethra		<i>S. epidermidis</i> , <i>Corynebacterium diptheroides</i> , various streptococci, various gram-negative rods (e.g., <i>E. coli</i>) ³

Diagrams that are less medically significant or present in smaller numbers.

These organisms are not part of the normal flora in that location but are important colonizers

They may serve a nutritional function. The(3)

intestinal bacteria produce several B vitamins and vitamin K . Poorly nourished people who are treated with oral antibiotics can have vitamins deficiencies as a result of the reduction in the normal flora .

However, since germ-free animals are

well-nourished, the normal flora is not essential for proper nutrition

flora of the skin

(all surface of body like chest hair, genital area)

predominates in skin
↑
↑ endocoritis in w/rif values.

The predominant organism is **staphylococcus**

epidermidis, which is a nonpathogen on the skin but such can cause disease when it reaches certain sites, as artificial heart valves and prosthetic joints. It is found on the skin much more frequently than its

- pathogenic relative staphylococcus aureus - *Staph. aureus*

There are about 10^3 - 10^4 organisms/cm² of skin. Most of them are located superficially in the stratum corneum, but some are found in the hair follicles and act as a reservoir to replenish the superficial flora after hand washing. Anaerobic organisms, such as Propionibacterium and Peptococcus, are situated in the deeper follicles in the dermis, where oxygen tension is low. Propionibacterium acnes is a common skin anaerobe that is implicated in the pathogenesis of acne.

Location	Important Organisms	Less Important Organisms
Skin	<i>Staphylococcus epidermidis</i>	<i>Staphylococcus aureus</i> , <i>Corynebacterium</i> (diphtheroid), various streptococci, <i>Pseudomonas aeruginosa</i> , anaerobes (e.g. <i>Propionibacterium</i>), yeasts (e.g. <i>Candida albicans</i>)

The yeast *Candida Albicans* is also a member of the normal flora of the skin. It can enter a person's bloodstream when needles pierce the skin (e.g., in patients with intravenous catheters or in those who use intravenous drugs). It is an important cause of systemic infections in patients with reduced cell-mediated immunity.

* *Yeast candida albicans* in skin can enter blood stream 'needles' leading to systemic infection. cell-mediated immunity is very low.

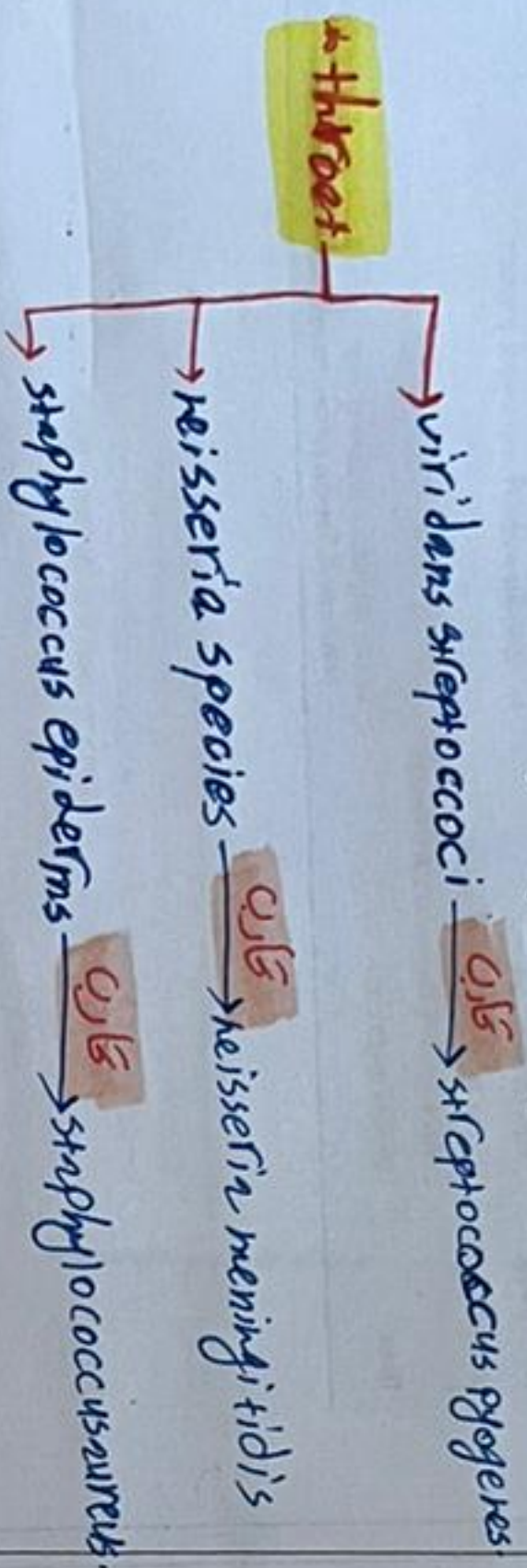


: Normal flora of the respiratory tract

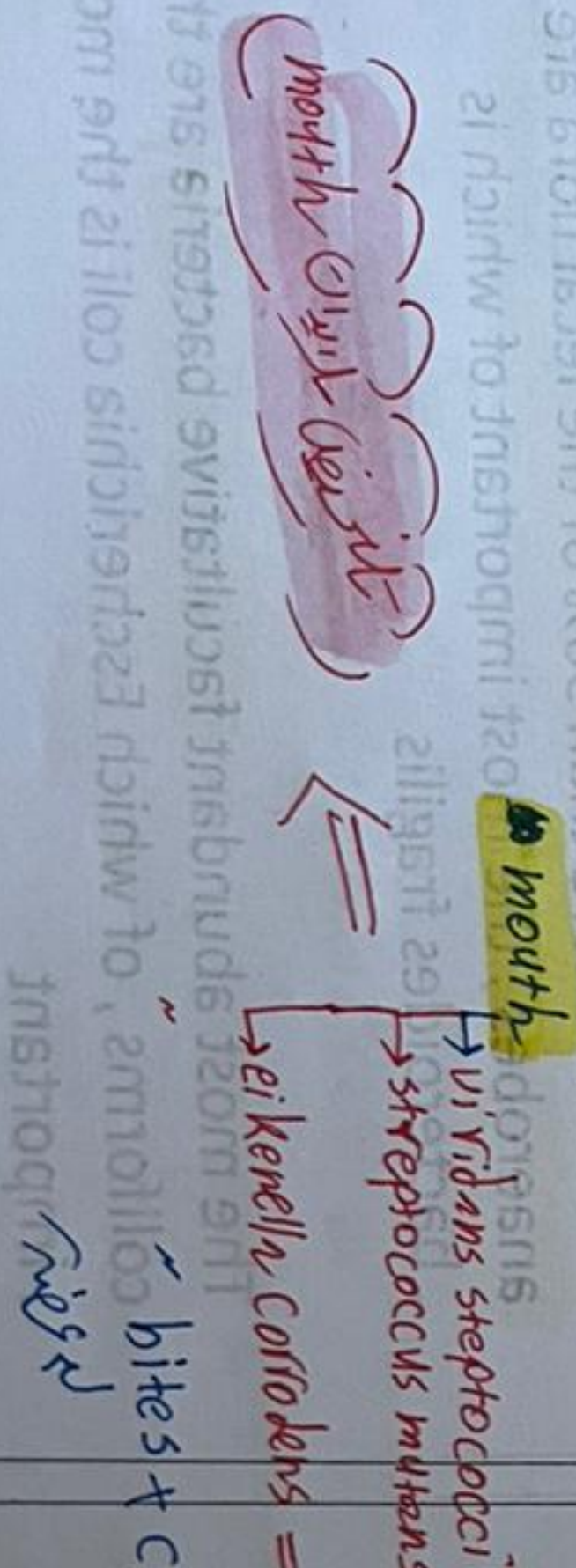
A wide spectrum of organisms colonize the nose, throat, and mouth, but the lower bronchi and alveoli typically contain few, if any, organisms. The nose is colonized by a variety of streptococcal and which staphylococcal species, the most significant of is the pathogen *S. aureus*. Occasional outbreaks of disease due to this organism, particularly in the newborn nursery. Can be traced to nasal, skin, or perianal carriage by health care personnel.

note nose → streptococci, staphylococci & aureus
alveoli & bronchi → staphylococcus aureus

The throat contains a mixture of viridans streptococci, *Neisseria* species, and *S. epidermidis*. These nonpathogens occupy attachment sites on the pharyngeal mucosa and inhibit the growth of the pathogens *Streptococcus pyogenes*, *Neisseria meningitidis*, and *S. aureus*, respectively.



The plaque on the enamel surface is composed of gelatinous, high-molecular-weight glucans secreted by the bacteria. The entrapped bacteria produce a large amount of acid, which demineralizes the enamel and initiates caries.

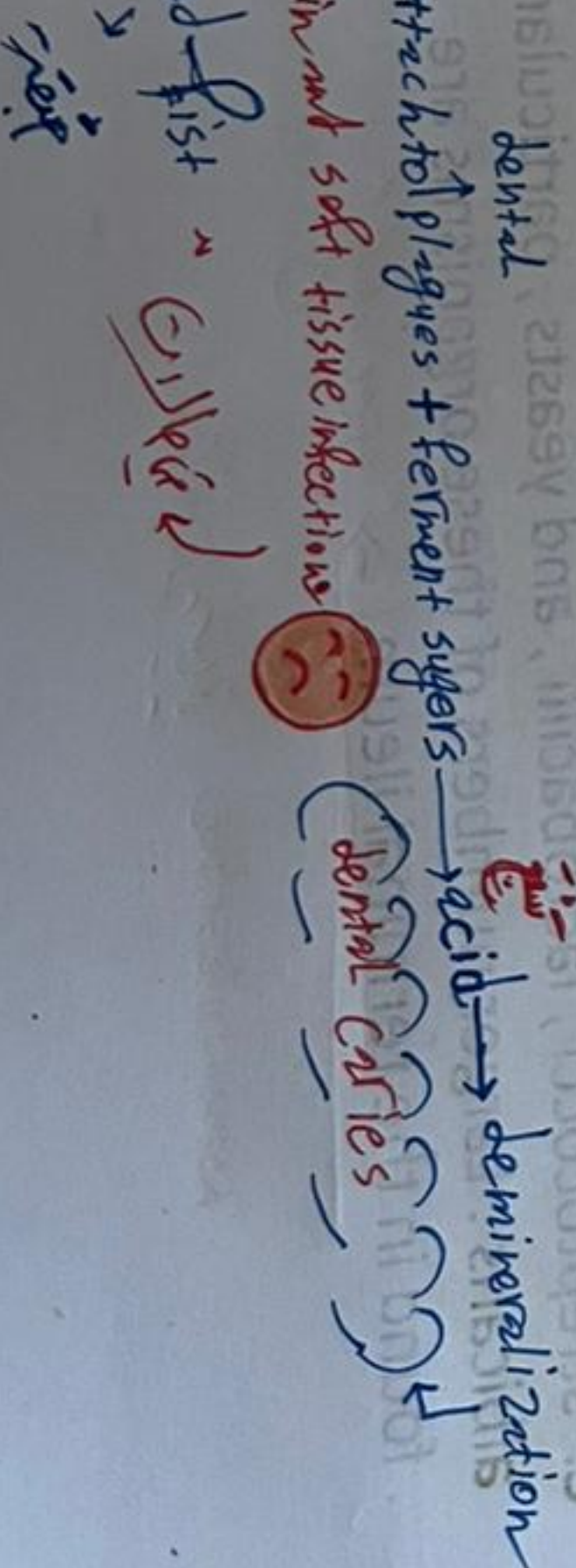


In the mouth, viridans streptococci make up about half of the bacteria. *Streptococcus mutans*, a member of the viridans group, is of special interest since it is found in large numbers in dental plaque, the precursor of caries.

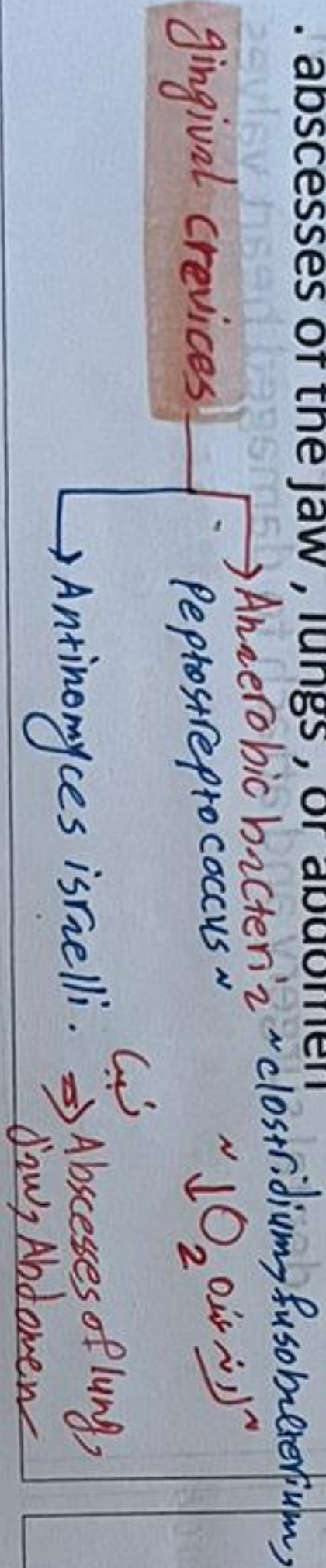
The viridans streptococci are also the leading cause of subacute bacterial (infective) endocarditis. These organisms can enter the bloodstream at the time of dental surgery and attach to damaged heart valves.

where reaching blood in dental surgery.

Eikenella corrodens, also part of the normal oral flora, causes skin and soft tissue infections associated with human bites and "clenched-fist" injuries (i.e., injuries to the hand that occur during fist fights).



Anaerobic bacteria, such as species of Bacteroids, Prevotella, Fusobacterium, Clostridium, and Peptostreptococcus, are found in the gingival crevices, where the oxygen concentration is very low. If aspirated, these organisms can cause lung abscesses, especially in debilitated patients with poor dental hygiene. In addition. The gingival crevices are the natural habitat of Actinomyces israeli - an anaerobic actinomycete that can cause abscesses of the jaw, lungs, or abdomen



: Normal flora of the intestinal tract

In normal fasting people, the stomach contains few organisms, primarily because of its low pH. The small intestine usually contains small numbers of *C. streptococci*, *Lactobacilli*, and yeasts, particularly *albicans*. Larger numbers of these organisms are found in the terminal ileum → "specimen"

small intestine → *Clostridium*

Nose	<i>Staphylococcus aureus</i>	<i>S. epidermidis</i> , <i>Corynebacterium</i> (diphtheroids), various streptococci
Mouth	<i>Vibrio streptococci</i>	Various streptococci, <i>Enterobacter</i>
Dental plaque	<i>Streptococcus mutans</i>	<i>Prevotella intermedia</i> , <i>Porphyromonas gingivalis</i>
Gingival crevices	Various anaerobes (e.g., <i>Bacteroides</i> , <i>Fusobacterium</i> , streptococci, <i>Actinomyces</i>)	Various streptococci (including <i>Streptococcus pyogenes</i> and <i>Streptococcus pneumoniae</i>), <i>Neisseria</i> species, <i>Haemophilus influenzae</i> , <i>S. epidermidis</i>
Throat	<i>Vibrio streptococci</i>	

The colon is the major location of bacteria in the body. Roughly 20% of the feces consists of bacteria, approximately 10^{11} organisms/g

→ e.g. *Bacteroides fragilis*

Note that more than 90% of the fecal flora are anaerobes, the most important of which is *bacteroides fragilis*

→ e.g. *E. coli*

The most abundant facultative bacteria are the coliforms, of which *Escherichia coli* is the most important

TABLE 6-3 Major Bacteria Found in the Colon

Bacterium ¹	Number/g of Feces	Important Pathogen
<i>Bacteroides</i> , especially <i>B. fragilis</i>	10^{10}	Yes
<i>Bifidobacterium</i>	10^{10}	No
<i>Eubacterium</i>	10^{10}	No
<i>Coliforms</i>	10^7 – 10^8	Yes
<i>Enterococcus</i> , especially <i>E. faecalis</i>	10^7 – 10^8	Yes
<i>Lactobacillus</i>	10^7	No
<i>Clostridium</i> , especially <i>C. perfringens</i>	10^6	Yes

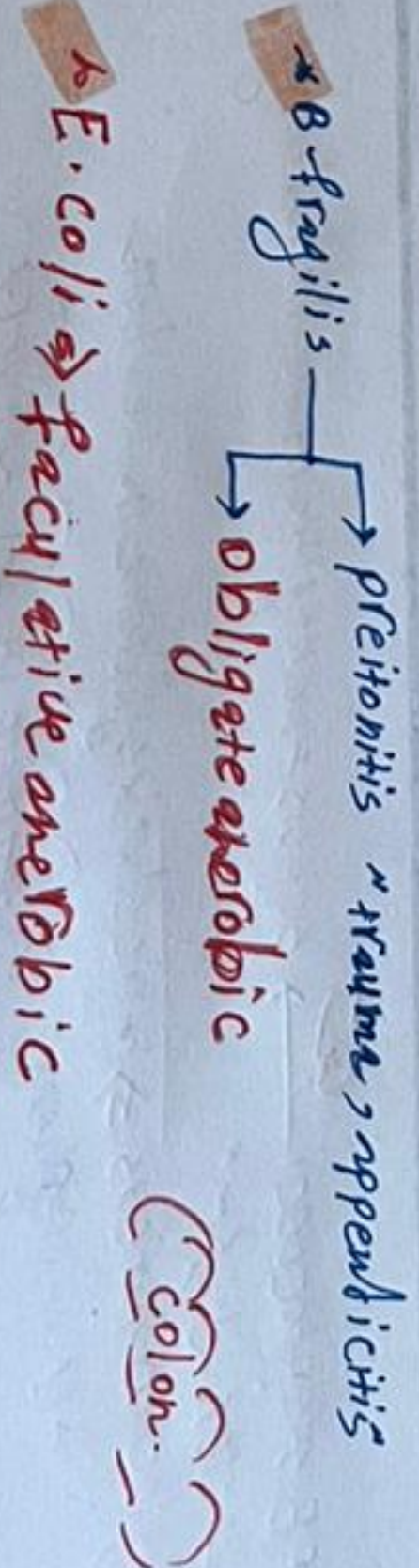
¹*Bacteroides*, *Bifidobacterium*, and *Eubacterium* (which make up more than 90% of the fecal flora) are anaerobes. *Coliforms* (*Escherichia coli*, *Enterobacter* species, and other gram-negative organisms) are the predominant facultative anaerobes.

Other important anaerobic pathogens include *Fusobacterium* and *Peptostreptococcus*, and other important facultative bacteria include *Enterococcus faecalis*, which causes urinary tract infections, particularly in hospitalized patients with decreased host defenses.

Pseudomonas
P. Aeruginosa is present in 10% of normal stools, as well as in soil and water.

The normal flora of the intestinal tract plays a significant role in extraintestinal disease.

For example, *E. coli* is the leading cause of urinary tract infections and *B. fragilis* is an important cause of peritonitis associated with perforation of the intestinal wall following trauma, appendicitis, or diverticulitis.



Antibiotic therapy (e.g., with clindamycin) can suppress the predominant normal flora, thereby allowing a rare organism such as the toxin-producing *Clostridium difficile* to overgrow and cause severe colitis. Administration of certain antibiotics, such as neomycin orally, prior to gastrointestinal surgery to "sterilize" the gut leads to a significant reduction of the normal flora for several days. Followed by a gradual return to normal levels.

6. sepsis + meningitis in newborn during delivery

20-15% of newborns

staphylococci aureus → toxic shock syndrome in newborn

5% of pregnant → antibiotic before delivery

Colon	Bacteroides fragilis, Escherichia coli	Bifidobacterium, Eubacterium, Fusobacterium, Lactobacillus, various aerobic gram-negative rods, Enterococcus faecalis and other streptococci, Clostridium
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group B streptococci → sepsis + meningitis in newborn during delivery

20-15% of newborns

staphylococci aureus → toxic shock syndrome in newborn

5% of pregnant → antibiotic before delivery

The vagina is located close to the anus and can be colonized by members of the fecal flora. For example, women who are prone to recurrent urinary tract infections harbor organisms such as E. coli and Enterobacter in the introitus. About 15% to 20% of women of childbearing age carry group B streptococci in the vagina. This organism is an important cause of sepsis and meningitis in the newborn and is acquired during passage through the birth canal. The vagina is colonized by S. aureus in approximately 5% of women, which predisposes them to toxic shock syndrome

: Normal flora of the genitourinary tract

The vaginal flora of adult women consists primarily of lactobacillus species. Lactobacilli are responsible for producing the acid that keeps the pH of the adult woman's vagina low

need estrogen

Before puberty and after menopause, when estrogen levels are low, lactobacilli are rare and the vaginal pH is high. Lactobacilli appear to prevent the growth of potential pathogens, since their suppression by antibiotics can lead to overgrowth by C. albicans.

Overgrowth of this yeast can result in Candida vaginitis

Urine in the bladder is sterile in the healthy person, but during passage through the outermost portions of the urethra, it often becomes contaminated with S. epidermidis, coliforms, diptheroids, and nonhemolytic streptococci. The area around the urethra of women and uncircumcised men contains secretions that carry Mycobacterium smegmatis, an acid-fast organism. The skin surrounding the genitourinary tract is the site of Staphylococcus saprophyticus, a cause of urinary tract infections in women

women's UTI

Ch-7 $\Rightarrow$ pathogenesis

terms

pathogen $\Rightarrow$ microorganism that can cause disease.

opportunistic pathogen $\Rightarrow$ pathogen that cause the disease in immunocompromised patient.
ميكروب (ممرض)، مسبب المرض

parasite

- $\rightarrow$ any organism that can cause harm
- $\rightarrow$ usually for protozoa + helminths
- $\rightarrow$ intracellular parasites " rickettsia, chlamydia "
- $\rightarrow$ there are facultative parasites

infection

- $\rightarrow$ literally $\Rightarrow$ an organism has entered the body.
- $\rightarrow$ in medical context $\Rightarrow$ an organism has entered the body + caused symptoms "disease"
"normal flora" (ميكروبيوتة طبيعية)

virulence

- $\rightarrow$ The No. of organisms required to cause disease.
- $\rightarrow$ LD₅₀ $\rightarrow$ Lethal dose of 50% of population.
- $\rightarrow$ ID₅₀ $\rightarrow$ infective dose of 50% of population.
- $\rightarrow$ ID $\propto \frac{1}{\text{virulence}}$
 - $\rightarrow$ e.g. $\rightarrow$ ID₅₀ in salmonella is 10^5
 - $\rightarrow$ ID₅₀ shigella is 10^2
 - $\rightarrow$ ID₅₀ in yersinia 10

notes

$\Rightarrow$ Bacteria may have more than 1 virulence factor.

$\rightarrow$ the main equation of pathogenesis is

(pathogen vs immune system)

number, site, virulence

humoral, cellular, disease

virulence factor

- $\rightarrow$ agent used by pathogen to sneak disease into the body
- $\rightarrow$ it helps the pathogen for:

① colonization $\rightarrow$ needs attachment by pili, Glycocalyx

"N. gonorrhoea"

"streptococcus"

② immunoevasion

$\rightarrow$ by Genetic variation

$\rightarrow$ by Ricket transmission

$\downarrow$
"meningitis"

Agg (مجموع)

بعض الميكروبات

③ Immunosuppression → inhibit the immune response.

④ entry and exit of cells → like intracellular bacteria

⑤ Obtain nutrition from the host

*Types of virulence factor

① Adherence component → pili, Glycocalyx

② capsule → for evasion ~ antiphagocytic

③ invasion enzymes → hyaluronidase, collagenase, fibrinolytic enzyme.
↓
degrade collagen
+ hyaluronic acid, important cellulitis by streptococcus
↓
streptokinase

→ coagulase ⇒ staphylococcus aureus + yersinia

④ toxins ⇒ exotoxins → enterotoxin of E. coli.

↓
endotoxins ⇒ lipid A, Teichoic Acid
↓
G⁻ ↓ ↓
G⁺

*note ⇒ Bacteria differs in their virulence factors

→ strain of E. coli ⇒ exotoxins 1 ⇒ watery diarrhea

→ other strain of E. coli ⇒ exotoxins 2 ⇒ Bloody diarrhea.

Why dose people get infectious diseases?

① from organism's perspective ⇒ ① infection & No ② infection & virulence.

② from host's perspective ⇒ weak immunity ~ innate ↓, humoral ↓, cellular ↓

• why the immune system becomes weak?

- ① → Genetic immunodeficiency → agammaglobulinemia
- ② → Acquired " " → AIDS, diabetes
- ③ → drug-induced immunosuppression →
organ transplant, autoimmune, cancer

• خزل مرضى كل الاصابه بالعدوى

L → serum dig Ag dig
 (e.g. → hepatitis B surface Ag)

L → serum dig Ab dig
 (e.g. → hepatitis C Ab, hepatitis A Ab)

① communicable → can be transmitted "TB via air"

② non communicable $\rightarrow$ not transmitted " botulism ✓

- **epidemic** → outbreak of the disease
- **endemic** → present at low level in specific population.
- **pandemic** → world wide distribution.

① subclinical infection → there is no or mild symptoms to escape diagnosis

الحياة ماري

(2) Asymptomatic $\rightarrow$ No symptoms
(3) chronic carrier $\rightarrow$ the organism continues grow with or without symptoms. $\rightarrow$ typhoid Mary
in this life organism and reactivation of symptoms.

(4) latent state $\rightarrow$ there is reactivation of the organism and reactivation of symptoms.

من 1 و 2 و 3 / عامل مشترك في اسمنا عليه العدوى
من 3 و 4 / عدوى من 3 و 4 / عدوى من 3 و 4
العدوى من 3 و 4 / عدوى من 3 و 4
reactivation / عدوى من 3 و 4

• How to detect asymptomatic people? colonization, normal flora of nasal cavities

وللأزيم فكره من مريد عند الرقة، العينة

② Ab titer $\rightarrow$

* stages of bacterial pathogenesis:

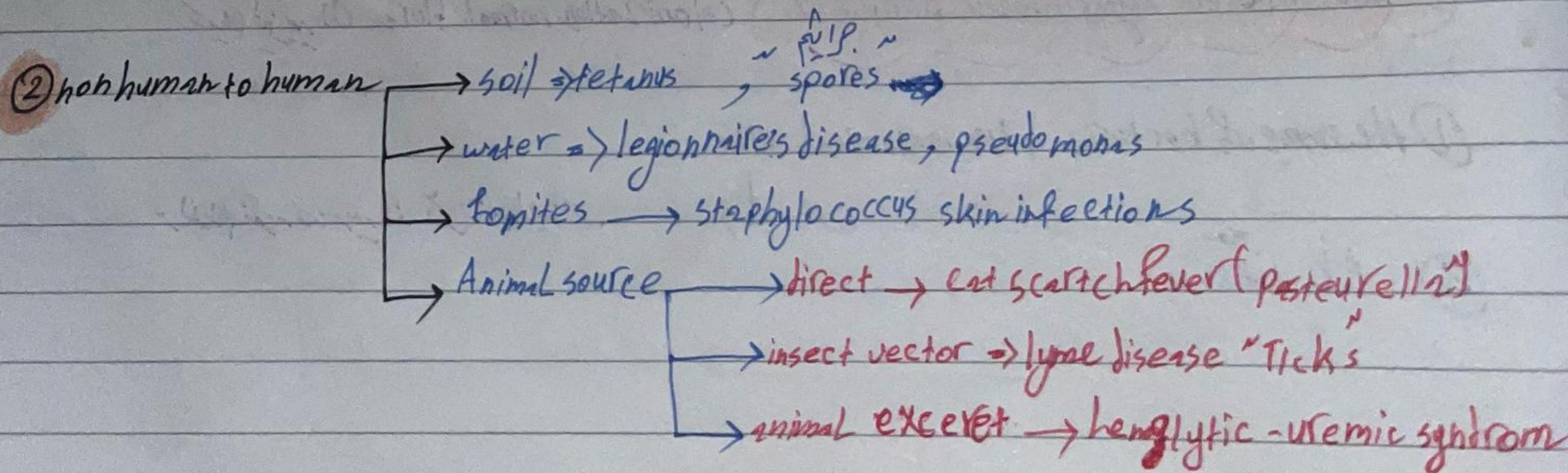
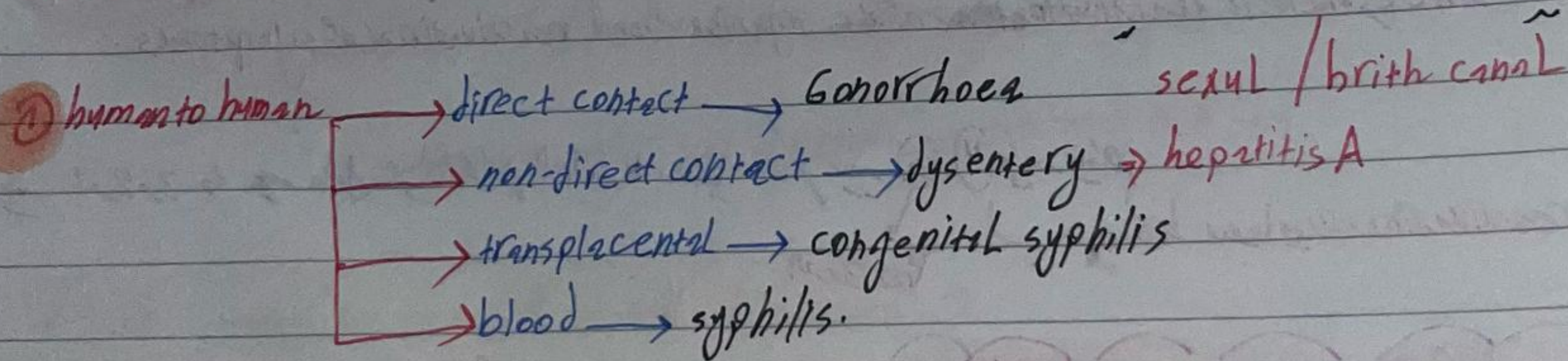
- ① transmission $\rightarrow$ to the portal of entry. $\leftarrow$ دخول
- ② evasion $\rightarrow$ host defense دفاع
- ③ Adherence $\rightarrow$ التعلق $\rightarrow$ mucous membranes by pili.
- ④ Colonization $\rightarrow$ $\uparrow$ No of bacteria $\rightarrow$ Adherence التعلق
- ⑤ disease production $\rightarrow$ exotoxin, invasion + inflammation.
- ⑥ Host response $\rightarrow$ استجابة $\rightarrow$ Adherence $\checkmark$, colonization $\checkmark$, disease production $\checkmark$
- ⑦ progression or resolution.

determination of bacterial pathogenesis

- ① transmission
- ② Adherence to cell surface
- ③ invasion + inflammation
- ④ toxin production
- ⑤ immunopathogenesis.

① transmission

- modes of transmission



N.F
↳ normal flora can cause disease, then infection via airborne droplets ~ respiratory N.F.
or fecal oral ~ GIT

hepatitis B, C / HN ~ ← org ~ / (فحص) screening في الدول المتقدمة

horizontal transmission ~ person to person

vertical transmission $\rightarrow$ mother $\rightarrow$ foetus

- ① placenta $\rightarrow$ syphilis, toxoplasma, cytomegalovirus
- ② Birth canal $\rightarrow$ Gonorrhoea, streptococcal.
- ③ milk $\rightarrow$ viruses $\rightarrow$ bacteria $\leftarrow$ staphylococcus aureus.

① Respiratory

② GIT

③ skin

~ GUT ~
④ Genital tract

① *Pili* → *Neisseria* + *E. coli*

② Glycocalyx → streptococcus + staphylococcus epidermidis
③ capsule (antiphagocytic + Adherent) → G +ve in general.

the molecules of attachment are called **adhesins**

→ the molecules of attachment are called ~~zone~~ zinc finger proteins.
E. coli, proteus ⇒ have flagella that allow them to move against urine flow.

Biofilms

- ① structure → polysaccharides + proteins surround microbes.
- ② function → protection from Antibiotics + cellular immunity
- ③ predisposing factor → Foreign bodies / synthetic valves or joints
- ④ risk → ~~retard~~ wound healing in diabetics
pseudomonas persistence in cystic fibrosis patients
- ⑤ mech → cell entry → colonization → Quorum sensing
attack ← biofilm + virulence factors → microbial density

notes → E. coli + salmonella produce curli protein
binding on endothelium + fibrinetic
active factor XII → Thrombosis → DIC
disseminated intravascular coagulation
كلية الغيرة دلفي أوسه درية

Invasion + inflammation

invasion ⇒ entry to host cell "need enzyme"

enzymes

① (collagenase + hyaluronidase) → degrade collagen + hyaluronic acid → spreading bacteria
e.g. strept. pyogenes

Cellulitis

② coagulase → coagulation "fibrinogen" → fibrin → walling off → protection
e.g. S. aureus, yersinia.

③ IgA protease → degrade IgA in mucus → Adherence + colonization.
"immunoglobulin" site of attachment
e.g. H. influenzae, strept. pneumoniae.

④ leukocidin → Kill neutrophils + macrophages.

other virulence factors for invasion

① Anti phagocytic factors

capsule ~ ^{G+ve} N. meningitidis, ^{G+ve} streptococcus pneumoniae, ^{G+ve rods} H. influenzae

② cell wall proteins of ^{G+ve} cocci

M protein ~ group A streptococcus → anti phagocytic
A protein (S. aureus) → ⊖ complement.

(pyogens)

spores

some capsules can be used in vaccination

no capsule → non pathogenic

anticapsule Ab enhance phagocytosis.

opsonization

→ enhancing pathogen phagocytosis by labeling it with opsonins IgG, C3b

inflammation

host defense mechanism, against pathogens

signs → redness, pain, loss of function, swelling, heat

mediators → histamine, LTs, PGs → prostaglandins
Leukotrienes

types

pyogenic → neutrophils are ^{ant} predominant cells.

granulomatous → lymphocytes + macrophages.

↑ permeability

intracellular survival

Def → the ability of pathogens to grow within cells.

They can be grown on media

These pathogens are not obligate intracellular

e.x → bacteria → TB, legionella, listeria.

→ fungi → histoplasma

benefits → protection from Ab + neutrophils.

mechanism

① inhibition of fusion of phagosome with lysosome

② inhibition of acidification of phagosome

③ escape from phagosome into cytoplasm

TB + legionella

e.x → listeria

avoid degradation

↓ lysosomal enzyme activity.

invasins → bacterial surface proteins interact with cellular receptor to allow invasion

phagosome
cytoplasm
listeria → move from one cell to another by actin rockets to evade the immune system

yops → yersinia outer-membrane proteins

→ inhibit phagocytosis by neutrophils

→ inhibit cytokine production by macrophages → inhibit immune system

notes → yersinia also use to coagulate to inhibit phagocytosis
→ yop J → cleaves the proteins in the cascade for TNF induction (hemorrhagic fever)
→ type of yersinia → GI → pneumonic

pathogen/city islands

cluster of genes that encode many virulence factors
→ invasins
→ exotoxins
→ adhesins

transfer together via conjugation/transduction

note cannot replicate independently

e.x → Gram negative bacilli → E. coli, salmonella, pseudomonas

→ Gram positive cocci → streptococcus pneumoniae

receptor → target → (note)

pseudomembranes → Thick adherent exudate on mucosal surface

e.x → Diphtheria → Corynebacterium diphtheriae

→ pseudomembranes colitis → Clostridium

④ toxin production.

properties of exotoxin

- ① excreted by living cell
- ② In G⁺, G⁺
- ③ polypeptide
- ④ highly antigenic
- ⑤ toxoid can be made
- ⑥ highly toxic
- ⑦ heat labile
- ⑧ bind to specific receptor.
- ⑨ not-pyrogenic
- ⑩ No enzymatic activity.
- ⑪ extra-chromosomal gene.

(endotoxin in the opposite)

- integral part, after cell death.
- G⁻ only
- lipopolysaccharide
- weak
- cannot
- moderately
- stable
- don't
- pyrogenic $\Rightarrow$ interleukin 1
- Yes
- chromosomal

examples

S. aureus | food poisoning |
Bacillus cereus | |
Bacillus anthracis (α -toxin) |
Streptococcus pyogenes.

E. coli, *Salmonella*
Shigella, *Cholera* "cholera gene" "

Disease

tetanus, diphtheria, botulism

G⁺ bacilli.
Meningococcus, sepsis

notes

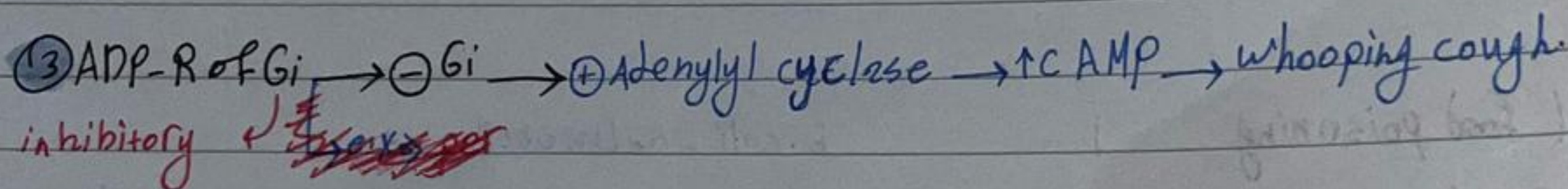
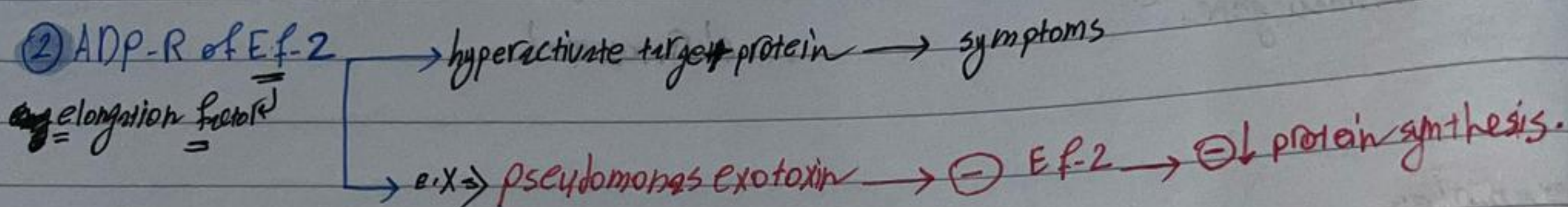
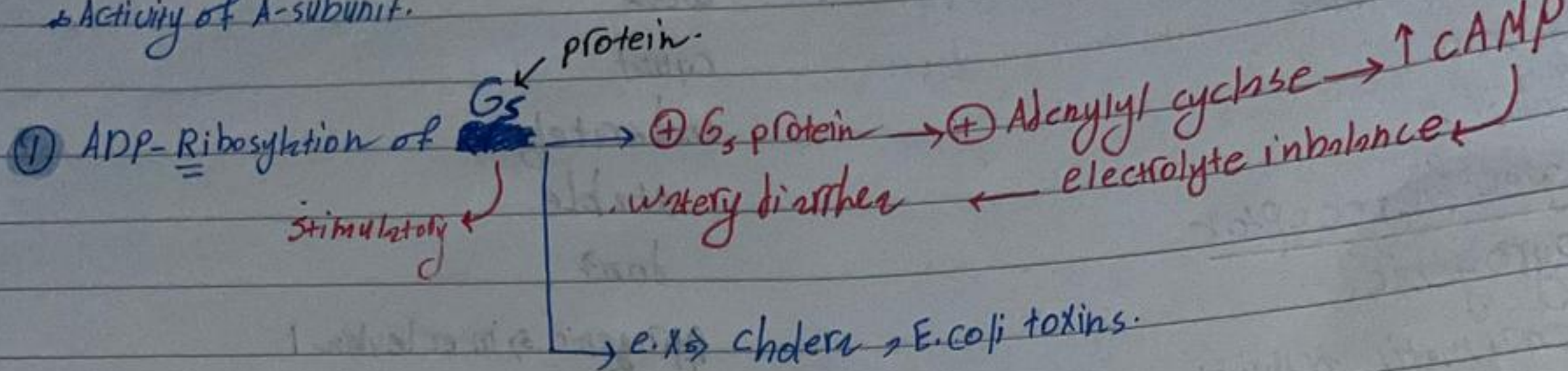
- $\rightarrow$ toxoid $\Rightarrow$ toxin + formalin processing or heat processing.
 - $\rightarrow$ specific receptors "ACh receptors"
 - $\rightarrow$ exotoxin found in plasmid $\Rightarrow$ can be passed to other bacteria.
 - $\rightarrow$ toxoid is used to induce antitoxins $\rightarrow$ treatment/prevention
- antitoxin $\leftarrow$ specific $\leftarrow$ botulism/tetanus diphtheria
- toxoid $\leftarrow$ " " " " " "

A-B subunit structure.

- exotoxin has
 - A subunit → toxic activity.
 - B subunit → Binding to receptors

e.g. → toxins of (diphtheria, tetanus, botulism, cholera, E. coli)

Activity of A-subunit.



way of secretion → by secretion system → secret to toxin → to extracellular → cell directly

Ab (antibody) → more effective

(injectosome) Type 3 secretion system → secretion system

→ in pseudomonas, shigella, salmonella, E. coli

toxin → transport pump → molecular junction

* exotoxin of ① Gram⁺ bacteria (mechanism + examples)

① ADP-R $\Rightarrow$ exⁿ *Corynebacterium diphtheriae*
Diphtheria toxin $\Rightarrow$ ADP-R of EF-2 $\rightarrow$ $\downarrow$ protein synthesis $\rightarrow$ cell death

$\rightarrow$ pseudomembrane
 $\rightarrow$ myocarditis

tox gene expression $\Rightarrow$ exotoxin production.

regulated by lysogenic bacteriophage

$\rightarrow$ regulated by Iron (Iron $\rightarrow$ expression)

- 2+3 neurotoxin.

② Blocking inhibitory neurotransmitters (glycine)

Clostridium tetani $\rightarrow$ tetanus toxin $\rightarrow$ $\ominus$ glycine $\rightarrow$ $\oplus$ excitatory $\rightarrow$ neurons
"spinal arch lock jaw" $\leftarrow$ spastic paralysis

tetanus toxin is composed of

- heavy chain (B-subtype) $\Rightarrow$ binds gangliosides of neural membrane

- light chain (A-subtype) $\Rightarrow$ protease degrades proteins needed for release of inhibitory neurotransmitter.

③ Blocking stimulatory NT

Clostridium botulinum $\rightarrow$ botulinum toxin $\rightarrow$ $\ominus$ ACh $\rightarrow$ flaccid paralysis.

2 subunits $\rightarrow$ 1 for binding

1 protease $\rightarrow$ degrade proteins for ACh release.

there are 6 subtypes $\rightarrow$ encoded on plasmid

$\rightarrow$ // ~ chromosomal DNA

$\rightarrow$ // ~ temperate bacteriophage.

④ enterotoxin + cytotoxins

* *Clostridium difficile* → enterotoxin A → enterotoxin → watery diarrhea
 → enterotoxin B → cytotoxins → damage to colonic mucosa.

[A+B] → pseudomembranous colitis
 → mech → glycosyltransferase → Glucosylation of inhibition Rho GTPases → ^{inhibition} @ single transduction
 → disaggregation of actin filaments → apoptosis.

⑤ multiple toxins

* *Clostridia* have → 7 lethal factor → necrosis + hemolysis
 → 5 enzymes → collagenase, protease, DNase, hyaluronidase, α-toxin.
 ~ deoxyribonuclease. ~ ↓ lecithinase

(note) → hydrolyze Lecithinase → destruction of membrane → death
 → No single *Clostridium* species has the 12 factors
 → *Clostridium perfringens* → food poisoning
 → watery diarrhea (enterotoxin)

* *St. aureus* → superantigens ↓

exfoliation

⑥ ~~exfoliation~~

* *S. aureus* → exfoliation toxin → epidermolytic factor → cleavage of desmoglein of desmosomes
 → scalded skin syndrome → detachment of epidermis

⑦ cooperative exotoxins

* *Bacillus anthracis* → edema factor → (+) Adenyl cyclase → edema
 → Lethal factor → α protease → (-) cell growth → cell death.
 → protective Ag → bind to receptor → prevents entry of edema + Lethal factor
 (note) → Ab against protective Ag prevent the activity of 3 factor

⑧ superantigen

S. aureus → TSST ~ toxic shock syndrome toxin ~
① binds major histocompatibility II ~ MHC II on macrophages without processing → + TH cells → IL-1 ~ helper T cells ~
② T cell multiplication → cytokine overproduction

TSST → ① in it acts as superantigen → ② in it acts as T-cell mitogen.

⑨ enter superantigen

S. aureus → locally acting superantigen ~ enterotoxin ~
vomiting + watery diarrhea "food poisoning"

(note) → toxin → + macrophages → cytokines → + enteric nervous → + vomiting center → vomiting

⑩ pore-forming

S. aureus methicillin resistance *S. aureus* "MRSA" → (PU) → exotoxin → panton-valent leukocidin → destruction of WBCs, skin, subcutaneous tissue.

(note) → VRSA → vancomycin resistance *S. aureus*.

⑪ superantigen causing rash

(strawberry tongue)

Strept. pyogenes → erythrogenic toxin → scarlet fever

The DNA for the toxin is acquired by temperate bacteriophage.

Non-lysogenized → pharyngitis without scarlet fever.

② G⁺ bacteria → mechanisms & examples.

① heat labile + heat stable enterotoxins.

heat labile → inactivated by 65°C for 30 min.
→ A+B subunits ~ activate G_s
→ ex → E. coli, cholera → watery diarrhea.

heat stable → not inactivated by boiling for 30 min.
→ Acts on ~~adenylyl~~ Guanylyl cyclase → ↑ cGMP → ⊖ Na⁺ reabsorption → diarrhea.
→ ex → E. coli

(note) → Genes for heat labile + stable are on plasmid.

② shiga toxin.

E. coli O157:H7 → verotoxin → remove adenine from 28S rRNA → ⊖ protein synthesis.
bloody diarrhea ← cell death

(notes) → acts like the toxin of shigella dysentery
→ found in undercooked meat of hamburgers
→ encoded by temperate bacteriophage.

→ cell death in kidney → renal failure
→ endothelial cell death → microangiopathic hemolytic anemia.
→ hemolytic-uremic syndrome → "HUS"

→ ciprofloxacin → shiga toxin → predisposes HUS

③ enterotoxin of vibrio cholera, Bacillus cereus $\equiv$ heat-labile toxin of E. coli $\rightarrow$ watery ^{diarrhea}

④ pertussis toxin

* Bor. pertussis $\rightarrow$ pertussis toxin $\rightarrow$ ADP-R of G_i $\rightarrow$ G_i inhibition $\rightarrow$ $\oplus$ Adenylate cyclase $\rightarrow$ $\uparrow$ cAMP
edema + changes in respiratory tract "whooping cough"
chemokine receptor transduction

lymph spleen, note lymph tissue $\rightarrow$ lymphocytes $\rightarrow$ not true $\leftarrow$ lymphocytosis
اللمف في الدم

endotoxin

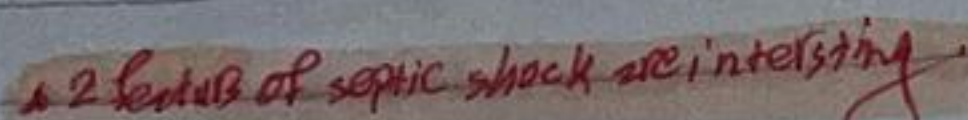
Lipopolysaccharides "LPS" $\rightarrow$ core polysaccharide
 $\rightarrow$ somatic "O" antigen (3 or 4 or 5) repeated 25 times $\rightarrow$ antigenic variation
(salmonella have > 1500 antigenic type)
 $\rightarrow$ lipid A $\rightarrow$ toxic protein

(note) $\rightarrow$ All endotoxins have the same symptoms "fever, shock", but differ in their severity.
 $\rightarrow$ endotoxin is less toxic than exotoxin.

how the LPS produce the symptoms of septic shock

LPS $\rightarrow$ $\oplus$ macrophage $\rightarrow$ IL-1, TNF $\rightarrow$ "fever, hypotension"
 $\rightarrow$ humor necrosis factor
 $\rightarrow$ $\oplus$ complement $\rightarrow$ C₃ $\rightarrow$ hypotension + edema
 $\rightarrow$ C₅ $\rightarrow$ neutrophil recruitment $\rightarrow$ $\uparrow$ $\rightarrow$ DIC
 $\rightarrow$ $\oplus$ tissue factor $\rightarrow$ $\oplus$ coagulation cascade $\rightarrow$ DIC

(notes) $\rightarrow$ other features of septic shock tachycardia, tachypnea, leukocytosis death of (30% $\rightarrow$ 60%) in ICU $\rightarrow$ intensive care units.
 $\rightarrow$ septic shock can be caused by G^+ via Teichoic acid



- ② In septic shock, bacteria killing by Antibiotics Leads to patient death

① Fever [by IL-1, IL-6, TNF on temp-regulatory center in hypothalamus]

- ③ DIC due to $\oplus$ coagulation cascade $\rightarrow$ thrombosis, tissue ischemia $\sim$ failure of vital organs
 $\hookrightarrow$ "disseminated intravascular coagulation", petechial or purpuric rash

(iv) activation of complement alternative pathway $\rightarrow C3e + C5e \rightarrow$ tissue damage + inflammation

- (1 → 5) → systemic inflammatory response syndrome ~ SIRS

→ Damage to endothelium → leakage of plasma + RBCs → hypotension
→ site for platelet aggregation → DIC

④ The evidence that the endotoxin caused the effects.

① purified LPS elicited the same effects

② antiserum against endotoxin blocked the effect.

⑤ How to detect DIC $\Rightarrow$ by assessing D-dimer (of fibrin)
lab-based, bedside

endotoxin causes the effect indirectly via IL-1, TNF

TNF $\rightarrow$ small amount $\Rightarrow$ beneficial
 $\rightarrow$ large amount $\Rightarrow$ harmful $\Rightarrow$ DIC, tumor necrosis

IV fluid sterilization

Autoclaving $\Rightarrow$ kills all organisms but release endotoxin

Filtration $\rightarrow$ physical removal of bacteria without releasing endotoxin
 $\rightarrow$ used in urine agar filtration
 $\rightarrow$ for examination of H₂O

6 true bacteria (S. aureus, strepto. pyogenes) causes of some effects of endotoxin viz lipoteichoic acid

septic shock treatment

① Antibodies against LPS, TNF release all signs of septic shock $\rightarrow$ [DIC]

② protein C / proteins $\rightarrow$ reversed DIC, but causes bleeding

immunopathogenesis

The immune system is the cause of disease not the organism

e.g. $\rightarrow$ rheumatic fever (Ab against M protein of strepto. pyogenes that cross-reacts with joint, heart, brain)
 $\rightarrow$ arthritis $\rightarrow$ carditis $\rightarrow$ chorea

Bacterial infections associated with cancers.

→ H. pylori
- helicobacter pylori associated with gastric carcinoma + gastric lymphoid tissue "MALT" lymphoma

* Campylobacter jejuni associated with intestinal "MALT" lymphoid.

how to prove it? Antibiotics in early stages causes the cancer to regress.

(notes) → h. pylori use urease "urea" → NH_3 causes ↑ pH to protect it from gastric acid
→ how to detect h. pylori → (1) IgG (2) IgM (3) stool Ag
(4) urea breath test → radioactive urea → radiation → container → urine (urine)

~~viruses that cause cancer~~
viruses that cause cancer → Epstein Barr virus, human papilloma virus (HPV), HIV

different strains of bacteria produce different diseases. Why? they have different virulence factors.

e.g. → S. aureus → pyogenic disease (endocarditis, septic arthritis, osteomyelitis)
→ non-pyogenic (scalded skin syndrome, toxic shock syndrome, food poisoning)

(notes) → The virulence factors are encoded on plasmids, transposons, pathogenicity island, lysogenic phages.
→ different genes + transmission → different virulence factor → different diseases.

typical stages of infection disease

- (1) incubation period → from acquisition of agent to symptoms.
- (2) prodrome period → Non-specific symptoms.
- (3) specific illness period → there is characteristic signs + symptoms of the disease occur.
- (4) recovery period → the patient return ~~being~~ healthy.
 - ↳ completely healthy
 - ↳ chronic carrier → transmit the disease to others.
 - ↳ latent infection → reactivation of organism (new signs)

↳ Koch's postulates:

① The microorganism found in organism suffering from disease but not in healthy

② The microorganism must be isolate + grown in pure culture.

↳ not all organisms can be isolated.

③ The cultured microorganism should cause disease when introduced into healthy organism

↳ depends to ID50 / immunity

④ The microorganism must be reisolated and be identical to the original causative agent

~~microorganism~~