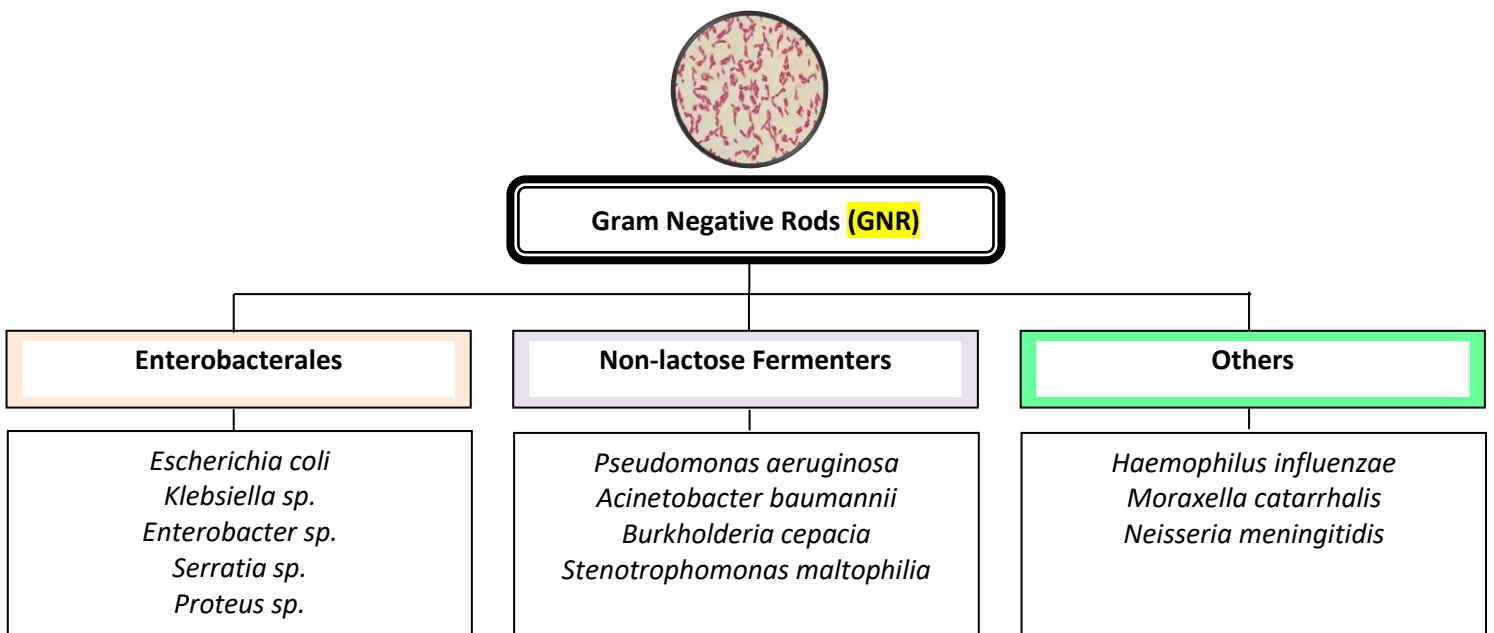
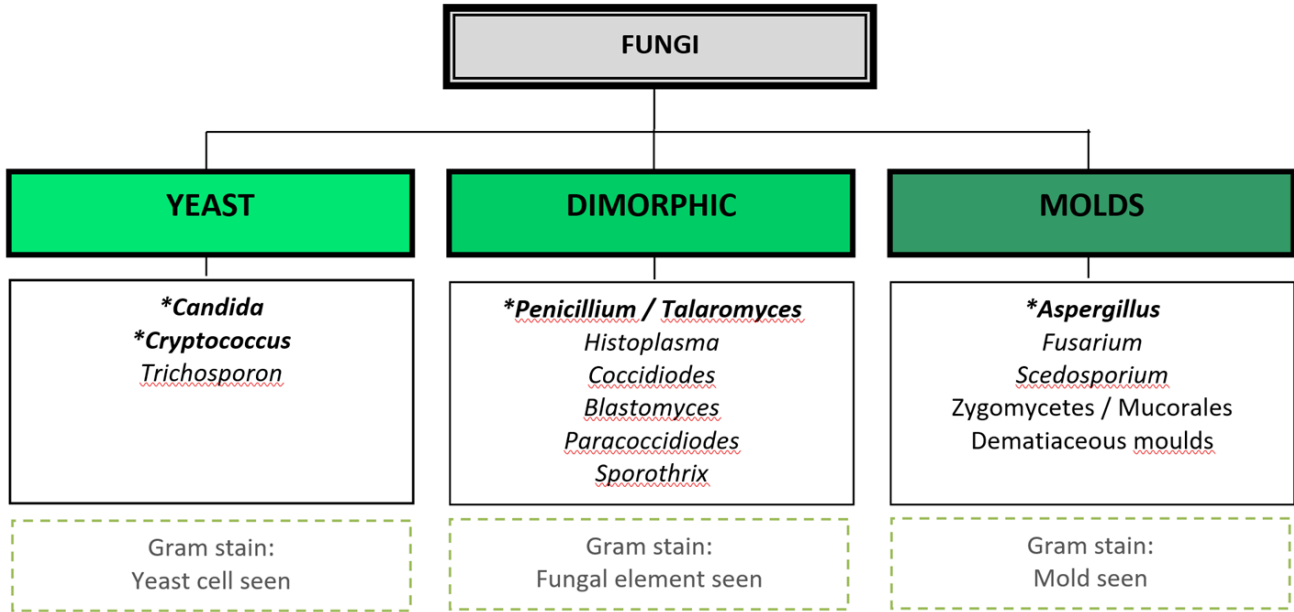


**Staph lugdunensis* can be a significant culture as it behaves like *Staph aureus*



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* More commonly seen in HSGb

ANTIFUNGAL SPECTRUM

Daily Integration of AMS (DIAMS) Hospital Sungai Buloh Antifungal Activity Spectra <small>Revised: October 2021</small>		YEAST							MOLD							DIMORPHIC			
		Candida species							Others	Aspergillus species				Others	Miscellaneous				
		Candida albicans	Candida parapsilosis	Candida tropicalis	Candida lusitanae	Candida glabrata	Candida auris	Candida krusei	Cryptococcus neoformans (Cryptococcus / Cryptococcal meningitis)	Aspergillus fumigatus (Most common)	Aspergillus flavus	Aspergillus niger	Aspergillus terreus	Mucorales (Mucormycosis)	Fusarium spp	Scedosporium spp	Talaromyces marneffei (Penicilliosis)	Histoplasma capsulatum (Histoplasmosis)	Sporothrix schenckii (Sporotrichosis)
Polyenes	Amphotericin B (AMB)	++	++	++	±	++	±	++	++	++	+	++	+	+	+	+	++	++	++
Azoles	Fluconazole (FLU)	++	++	++	++	C	X	X	++	X	X	X	X	X	X	X	+	±	±
	Itraconazole (ITR)	+	+	+	+	±	±	±	+	±	±	±	±	±	±	±	++	++	++
	Voriconazole (VOR)	+	+	+	+	±	±	±	+	++	++	++	++	X	++	+	++	+	+
	Posaconazole (POS)	+	+	+	+	±	±	±	+	+	+	+	+	++	++	±	X	+	+
	Isavuconazole (ISA) [N/A]	+	+	+	+	±	±	±	+	++	++	++	++	++	++	±	X	+	+
Echinocandins	Micafungin (MICA)	++	+	++	++	++	++	++	X	±	±	±	±	X	X	X	X	X	X
	Andulafugin (ANI)	++	+	++	++	++	++	++	X	±	±	±	±	X	X	X	X	X	X
	Caspofungin (CAS) [N/A]	++	+	++	++	++	++	++	X	±	±	±	±	X	X	X	X	X	X
Fluoropyrimidines	Flucytosine (5FC)	+	+	+	+	+	+	++	X	X	X	X	X	X	X	X	X	X	

Guide:

- ++ **Active (Recommended)** : Drug is recommended due to reliable in-vitro activity, clinically effective & recommended in guidelines.
- + **Active (Alternative)** : Drug may be used as alternative therapy. It has in-vitro activity & likely to be clinically effective but not used as 1st line due to overly broad spectrum/toxicity profile / limited clinical evidence or experience).
- ± **Variable coverage** : Variable sensitivity.
- C **Conditional Use** : Can only be used in specific condition (i.e; only as part of combination therapy /sensitivity-dose-dependent (SDD) (May be sensitive in higher doses).
- X **Not Recommended** : Agent is not recommended (likely to be resistant due poor penetration/unfavourable toxicity profile/insufficient data/inherent resistance).
- [N/A] **Not Available** : Registered in Malaysia but not available in HSGb

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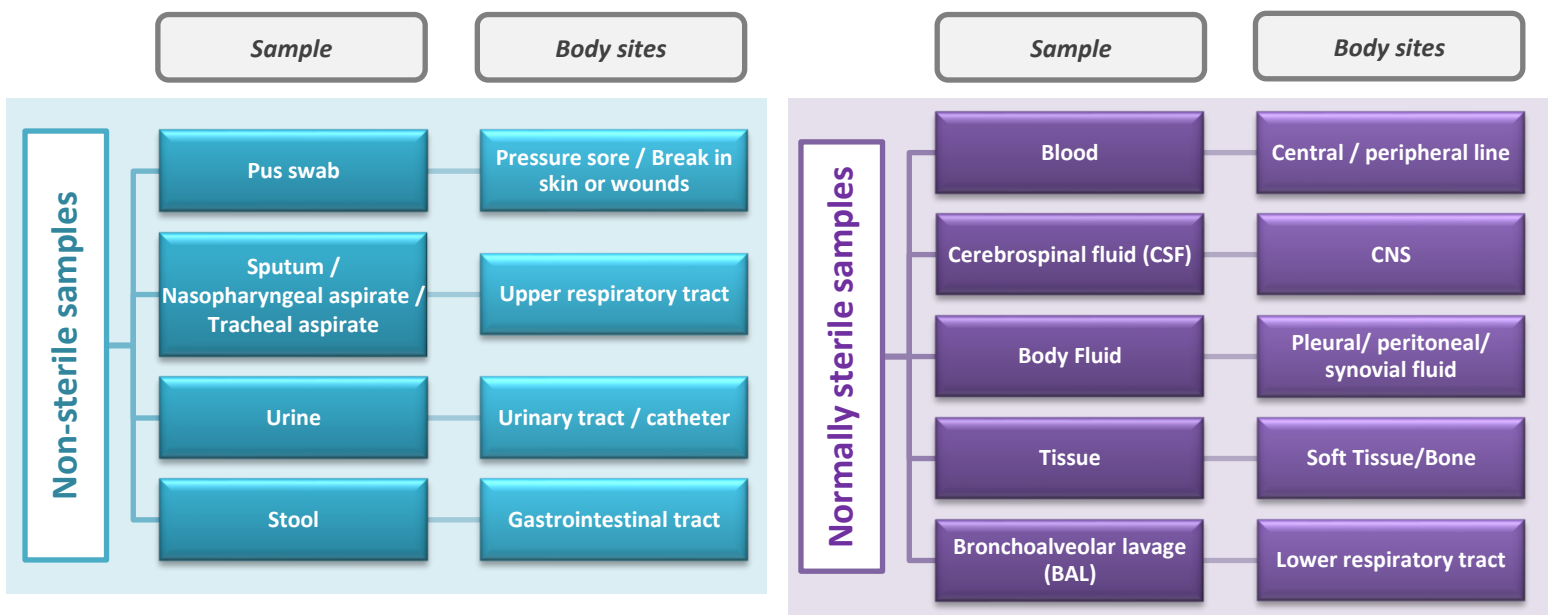
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CONTAMINANT VS COLONISER VS PATHOGEN

CONTAMINANT	
<ul style="list-style-type: none"> ✚ Introduced into sample due to poor sampling technique. ✚ Do not require treatment, unless repeated C&S isolate the same microorganism(s) despite proper technique & patient demonstrates clinical infection. 	<p>Common contaminants:</p> <ul style="list-style-type: none"> ▪ Coagulase-negative species (CONS) ▪ <i>Bacillus</i> spp. ▪ <i>Corynebacterium</i> spp. ▪ <i>Propionibacterium acnes</i> ▪ <i>Micrococcus</i> spp.
COLONISER	
<ul style="list-style-type: none"> ✚ Grow in or on <u>non-sterile body sites/prosthetic devices</u> without causing infection. ✚ Does not normally harm the patient nor require antibiotic treatment. ✚ Colonisation may become infectious when there is: <ul style="list-style-type: none"> ○ Weakened immune systems (opportunistic pathogen) ○ Access to a normally sterile site through invasive procedures 	<p>Common colonisers:</p> <ul style="list-style-type: none"> ▪ Skin flora (eg: <i>Staphylococci</i>) ▪ Enteric flora (eg: <i>E. coli</i>, <i>Enterococci</i>, anaerobes) ▪ Oral/urogenital flora (eg: <i>Candida albicans</i>)
PATHOGEN	
<ul style="list-style-type: none"> ✚ Infectious agents that cause disease or illness to its host. ✚ Microorganisms isolated from a <u>normally sterile site</u> that may cause infection. ✚ Signs & symptoms of true infection: <ul style="list-style-type: none"> ○ Generalised: Fever, malaise ○ Localised: Swelling due to inflammation, heat, pain, erythema 	<p>Common pathogens:</p> <ul style="list-style-type: none"> ▪ <i>Staphylococcus aureus</i> ▪ <i>Streptococcus pneumoniae/pyogenes/agalactiae</i> ▪ <i>Enterobacteriaceae</i> ▪ <i>Pseudomonas aeruginosa</i> ▪ Fungus (yeast i.e. <i>Candida albicans</i>)

NON-STERILE & STERILE CULTURE SAMPLES FROM BODY SITES/ DEVICES



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COMMON RESISTANCE IN HOSPITAL ACQUIRED INFECTIONS

Microorganism	Description						
Methicillin-resistant <i>S. aureus</i> (MRSA)	Resistance conferred by <i>MecA</i> gene, leading to resistance to Methicillin, Oxacillin & Cephalosporins.						
Vancomycin-intermediate <i>S. aureus</i> (VISA) Vancomycin-resistant <i>S. aureus</i> (VRSA)	Rare resistance, seen mostly in patients with long-term Vancomycin therapy (MIC creep).						
Vancomycin-resistant Enterococcus (VRE)	Resistance conferred by plasmid-mediated VanA and VanB gene complexes.						
AmpC β -Lactamase-Producing Enterobacteriales** <i>**Enterobacteriales previously known as Enterobacteriaceae</i>	<ul style="list-style-type: none"> Mutations which confers resistance to penicillins & 3rd generation cephalosporins. Exists in 3 forms: <ul style="list-style-type: none"> On chromosome + inducible On chromosome + de-repressed On plasmid + constitutive May be inducible by certain antibiotics. Refer table below [click here]. 						
Extended Spectrum Beta-Lactamase-producing Enterobacteriales** (ESBL-E) <i>**Enterobacteriales previously known as Enterobacteriaceae</i>	<ul style="list-style-type: none"> TEM-/SHV-/CTX-M enzyme-producing organisms under Ambler Class A. Carbapenem is the first line treatment for ESBL-E infection outside urinary tract based on data from a large clinical trial (MERINO trial) which showed 30-d mortality difference of 12% vs 4% when meropenem was used as compared to piperacillin/tazobactam. 						
Carbapenem-resistant Enterobacteriales** (CRE) <i>**Enterobacteriales previously known as Enterobacteriaceae</i>	<p>Heterogenous group of pathogens with multiple potential mechanism of resistance. It can be broadly divided into 2 groups:</p> <p>a) Carbapenemase-producing (CP-CRE)</p> <p>b) Non-carbapenemase producing (Non CP-CRE)</p> <table border="1" style="margin-left: auto; margin-right: auto;"> <thead> <tr> <th colspan="2">CRE Mechanism of Resistance</th> </tr> <tr> <th>CP-CRE</th> <th>Non CP-CRE</th> </tr> </thead> <tbody> <tr> <td> <ol style="list-style-type: none"> Serine (Ambler Class A) Eg: KPC Metallo-β-lactamases (Ambler Class B) Eg: NDM, VIM, IMP Serine (Ambler Class D) Eg: OXA-48 </td> <td> <ol style="list-style-type: none"> Efflux pump Porin Mutations Other β-lactam hydrolyzing enzymes Eg: ESBLs, AmpC </td> </tr> </tbody> </table>	CRE Mechanism of Resistance		CP-CRE	Non CP-CRE	<ol style="list-style-type: none"> Serine (Ambler Class A) Eg: KPC Metallo-β-lactamases (Ambler Class B) Eg: NDM, VIM, IMP Serine (Ambler Class D) Eg: OXA-48 	<ol style="list-style-type: none"> Efflux pump Porin Mutations Other β-lactam hydrolyzing enzymes Eg: ESBLs, AmpC
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<p>Carbapenem-resistant <i>Acinetobacter baumannii</i> (CRAB)</p> <p>#Pan-sensitive <i>A. baumannii</i> are sensitive to sulbactams, meropenem, pip/tazo, cefepime, AMG & FQs.</p>	<p>Once <i>Acinetobacter baumannii</i> exhibits carbapenem resistance, it generally acquired resistance against most other antibiotics expected to be active against pan-sensitive[#] (wild-type) <i>A. baumannii</i>.</p> <ul style="list-style-type: none"> • Presence of carbapenemases such as OXA-24/40-like, OXA-23-like, metallo-β-lactamases and additional serine carbapenemases causes carbapenem resistance. • Mutation to the PBP & β-lactamases production confers additional resistance towards sulbactam. • Presence of Aminoglycoside modifying enzyme (AME) causes additional resistance towards aminoglycosides. • Upregulation of efflux pumps mediates resistance to FQs.
<p>Difficult-to-treat resistant (DTR) <i>Pseudomonas aeruginosa</i></p>	<ul style="list-style-type: none"> • Exhibit resistance against antibiotic that commonly used to treat <i>Pseudomonas</i> (pip/tazo, ceftazidime, cefepime, aztreonam, meropenem, imipenem, ciprofloxacin & levofloxacin). • Multiple complex resistance mechanisms can exist simultaneously including reduced porin expression (OprD), efflux pump upregulation (MexAB-OprM), PBP mutations, AmpC & carbapenemase productions.

Significant AmpC β -Lactamase-Producing Enterobacterales (CEK)

C	<i>Citrobacter freundii</i>
E	<i>Enterobacter cloacae</i> complex
K	<i>Klebsiella aerogenes</i> (previously known as <i>Enterobacter aerogenes</i>)

- **CEK:** Treat with Cefepime or Meropenem for initial treatment even when C&S shows susceptibility to Ceftazidime/Tazocin.
- Antibiotic(s) capacity for inducing AmpC production and their efficacy against AmpC-producing organism as listed below:

	Weak AmpC Inducer Antibiotics	Strong AmpC Inducer Antibiotics
Effective against AmpC producing pathogens	<p>Cefepime (subjected to inoculum effect)</p>	<p>Carbapenems</p>
Not effective against AmpC producing pathogens	<p>Piperacillin\pmTazobactam 2nd & 3rd gen Cephalosporins Aztreonam</p>	<p>Ampicillin\pmSulbactam Amoxycillin\pmClavulanate 1st gen Cephalosporins</p>

* **Inoculum effect:** Efficacy of antibiotic reduced due to significant increase in Abx MIC because of high bacterial load.

Lower risk for significant AmpC β -Lactamase-Producing Enterobacterales (SPM)

S	<i>Serratia marcescens</i>
P	<i>Providencia</i> spp.
M	<i>Morganella morganii</i>

- **SPM:** Treat as per susceptibility testing. In severe infection & high bacterial burden* conditions, there is a risk of treatment failure.
- → refer to ID if indicated. (*High bacterial burden eg: undrained abscesses, infective endocarditis, ventriculitis, etc)

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ESKAPE

- ESKAPE pathogens are leading causes of nosocomial infections with rapidly growing multidrug resistance (MDR) and virulence

E	<i>Enterococcus faecium</i>
S	<i>Staphylococcus aureus</i>
K	<i>Klebsiella pneumoniae</i>
A	<i>Acinetobacter baumannii</i>
P	<i>Pseudomonas aeruginosa</i>
E	<i>Enterobacter spp.</i>

HACEK

- Fastidious GNR organisms commonly colonize the human oropharynx as normal flora with propensity to cause infective endocarditis (IE)
- May appear as “culture negative endocarditis” as HACEK grows slowly in blood culture media

H	<i>Haemophilus parainfluenzae/ aphrophilus</i>
A	<i>Actinomyces comitans (Aggregatibacter spp.)</i>
C	<i>Cardiobacterium hominis</i>
E	<i>Eikenella corrodens</i>
K	<i>Kingella kingae</i>
*Culture negative endocarditis organisms:	<i>Brucella spp.</i>
	<i>Coxiella burnetii (agent of Q fever)</i>
	<i>Bartonella spp.</i>

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