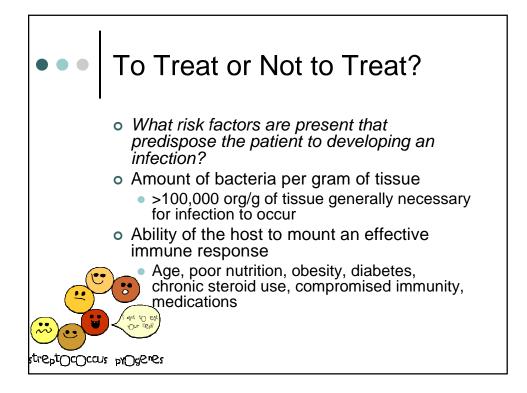
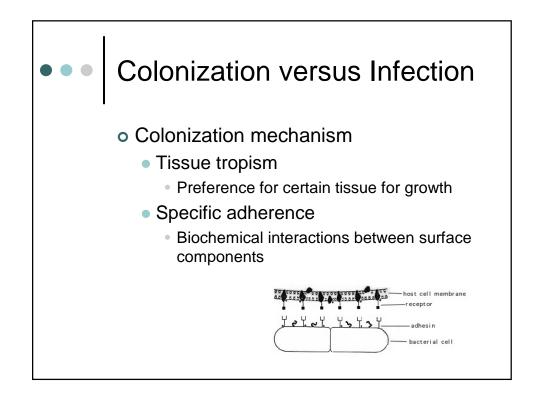
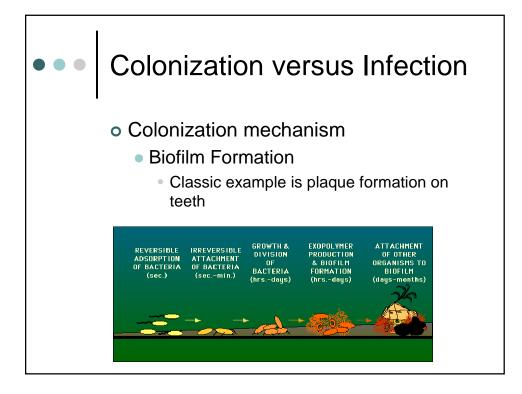


•••	Normal Flora		
	Body Site	Indigenous Microbiota	
	Skin	Diphtheroids (e.g. <i>Corynebacterium spp.</i>), <i>Propionobacteria</i> , Staphylococci (especially <i>Staph. epi</i>)	
	Gastrointestinal tract	Bacteroides spp., Clostridium spp. (some species), Diphtheroids, Enterobacteriaceae (e.g., <i>E.coli, Klebsiella spp.</i>), Enterococcus, <i>Candida spp.</i> (usually <i>C. albicans</i>)	
	Upper respiratory tract	Haemophilus spp., Neiserria spp., Viridans strep., Strep. pneumo., Staphylococci, Diphtheroids	
	Genital tract	<i>Corynebacterium spp.</i> , Enterobacteriaceae, <i>Lactobacillus, spp., Mycoplasma spp.,</i> Staphylococci, Streptococi, <i>Candida spp.</i>	







•••	Colonization versus Infection			
	Bacteria	Site of colonization	Pathogenic Impact	
	MRSA	Nostrils, belly button, under arms, groin, etc	Bacteremia, Pneumonia, Endocarditis, Meningitis, Osteomyelitis	
	<i>E. Faecalis</i> (VRE)	Gastrointestinal tract	Bacteremia, Endocarditis, Pyelonephritis	
	Strep. pneumo	Upper respiratory tract	Pneumonia (LRT), Meningitis	
	S. pyogenes	Mouth/Pharynx	Pneumonia, Endocarditis, Nephritis, Rheumatic Fever, CSSI	
	Neisseria	Upper respiratory tract/Pharynx	Meningitis	
	E. Coli	Small intestine	Bacteremia, Intestinal infections, UTI	
	Pseudomonas aeruginosa	Small intestine	Quintessential opportunistic pathogen of humans	

