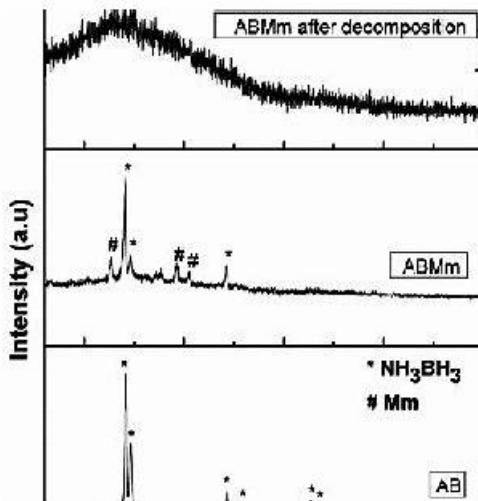


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American Society of Microbiology ABMM Sample Questions (Q111-Q116):

NEW QUESTION # 111

A microbiology laboratory receives a blood culture from a patient with suspected bacterial endocarditis. After 24 hours of incubation, Gram-positive cocci in clusters are observed on the Gram stain. Catalase test is positive, and coagulase test is negative. Further testing reveals susceptibility to novobiocin. The MOST likely organism is:

- A. *Staphylococcus epidermidis*
- B. *Staphylococcus lugdunensis*
- C. *Staphylococcus aureus*
- D. *Staphylococcus saprophyticus*

Answer: A

NEW QUESTION # 112

The gut microbiome has been implicated in the pathogenesis of autoimmune diseases. Dysbiosis, a disruption in the composition and function of the gut microbiota, can lead to increased intestinal permeability and the translocation of microbial products that trigger inflammatory responses. Which of the following bacterial components is a potent activator of the innate immune system and has been linked to the development of autoimmunity?

- A. Peptidoglycan
- B. Flagellin
- C. Lipoteichoic acid (LTA)
- D. **Lipopolysaccharide (LPS)**

Answer: D

NEW QUESTION # 113

A foodborne outbreak is suspected after several individuals who attended a potluck dinner develop nausea, vomiting, and abdominal cramps within 1-6 hours of eating. No fever or diarrhea is reported. The MOST likely bacterial toxin involved is:

- A. Shiga toxin
- B. Botulinum toxin
- C. *Clostridium perfringens* enterotoxin
- D. **Staphylococcal enterotoxin**

Answer: D

NEW QUESTION # 114

A research laboratory is investigating the mechanisms of antigenic variation in *Trypanosoma brucei*, the causative agent of African trypanosomiasis. The parasite evades the host immune response by periodically changing its surface coat protein. This process primarily involves:

- A. Segmental gene conversion events involving a large repertoire of silent VSG genes.
- B. RNA interference-mediated silencing of expressed VSG genes followed by activation of a new VSG gene.
- C. Recombination between different VSG gene loci on minichromosomes.
- D. High-frequency point mutations in the gene encoding the variant surface glycoprotein (VSG).

Answer: A

NEW QUESTION # 115

A research team is investigating the mechanism of bacterial resistance to a novel fluoroquinolone antibiotic.

They isolate a resistant strain of *Escherichia coli* from a patient with a urinary tract infection. Genetic analysis of this strain reveals a point mutation in the *gyrA* gene. This mutation MOST likely results in:

- A. Modification of the ribosomal target site, preventing antibiotic binding.
- B. Increased expression of efflux pumps that actively transport the antibiotic out of the cell.
- C. **Alteration of the DNA gyrase enzyme, reducing its affinity for the fluoroquinolone.**
- D. Production of an enzyme that inactivates the fluoroquinolone molecule.

Answer: C

NEW QUESTION # 116

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