**Publicaciones VITEK MS PRIME - ECCMID 2024 - POSTERS**

P0624 - Optimizing the preparation of non-tuberculous mycobacterial (NTM) isolates for MALDI-TOF (VITEK) MS identification and speciation

P0674 - Improved identification of Mycobacterium avium complex by MALDI-TOF coupled with machine learning

P0747 - Direct identification of microorganisms by MALDI-TOF MS from positive blood cultures and rapid antibiotic susceptibility testing

P0874 - The impact of rapid microbial identification report on the clinical and health economics of hospitalised patients with bloodstream infection

P1746 - High accuracy and prediction values for mecA+ methicillin-resistant (MRSA) and phenotypic oxacillin-resistant Staphylococcus aureus (ORSA) on the basis of MALD

P1753 - Detection of lipid A by MALDI-TOF for colistin heteroresistant strains

P1989 - Is the correct identification directly from blood culture bottles just a dream

P1991 - Evaluation of a rapid MALDI-TOF protocol for the detection of pathogenic microorganisms on positive blood cultures in an MDRO-endemic hospital environment

P1992 - Performance and workflow comparison between two MALDI TOF MS systems in microbiology routine laboratory settings

P1993 - Performance evaluation of Vitek MS PRIME compared to Vitek MS

P1994 - MALDI-TOF MS database expansion for robust and improved diagnostics in microbiology

P1996 - Performance of the VITEK MS PRIME was equivalent to the VITEK MS in microbial identification with improved time to result

P2001 - Bacterial identification by direct MALDI-TOF MS from blood culture bottles.pdf

P2002 - Evaluation of fast direct MALDI-TOF identification of organisms from positive blood-culture bottles (BCB) the saponin method vs full extraction

P2005 - FASTinov A simple protocol for rapid bacterial identification by MALDI-TOF MS directly from positive blood cultures

P2008 - Colibri Evaluation of VITEK MS prime bacterial identification performance in conjunction with a fully automated slide preparation system in urine cultur

P2009 - Detection of cfiA gene in Bacteroides fragilis using MALDI-TOF