

International Nurse Researcher Hall of Fame and Conversation with the Honorees - Session 4

Title:

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Keywords:

award recipient, nurse researcher hall of fame and research journey

Summary:

Attendees will experience the opportunity to hear International Nurse Researcher Hall of Fame honorees discuss their work and their research journeys. Session moderated by 2019=2021 President, Richard Ricciardi, PhD, CRNP, FAANP, FAAN.

References:

- No references were used in the production of this session.

Organizer

Connie White Delaney, PhD, RN, FAAN, FACMI, FNAP, RN, PhD
University of Minnesota
School of Nursing
Dean and Professor
Minneapolis, Minneapolis
USA

Author Summary: Dr. Delaney is recognized as a leader and visionary with unique organizational ability to re-create/co-create integrative communities, partnerships, programs, systems and collectives that produce sustainable relationship empowered impact, wholeness, conscious wellbeing, planetary health, and responsible stewardship. She is committed to transforming challenges into opportunities that advance discovery that transforms health, health systems, education, and integrative Informatics. She cultivates leadership transcendence in academia, health care, and informatics.

Organizer

Carol M. Musil, PhD, RN, FAAN, FGSA, PhD
Case Western Reserve University
Bolton School of Nursing
Associate Professor
Cleveland, Ohio
USA

Author Summary: Dean and Edward J. and Louise Mellen Professor of Nursing, Case Western Reserve University Frances Payne Bolton School of Nursing, Cleveland, Ohio, USA Research area: Older adult physical and psychological health, family well-being during strain and transition

Organizer

Jacquelyn Taylor, PhD, PNP-BC, RN, FAHA, FAAN

New York University

Rory Meyers College of Nursing

Vernice D Ferguson Endowed Professor in Health Equity

New York, New York

USA

Author Summary: Dr. Taylor is a nationally renowned genomics nurse scientist. Her utilizes many laboratory techniques such as targeted candidate gene, genome wide association, epigenome wide association and genome sequencing to answer research questions related to gene-environment interactions on chronic health conditions among minority populations.