



INTERNATIONAL GENETIC EPIDEMIOLOGY SOCIETY

2022 IGES Abstract Guidelines

Abstracts should follow the same format as in previous year's published abstract booklet and should be submitted as a Word file (.doc, .docx, .rtf). (See: <http://onlinelibrary.wiley.com/doi/10.1002/gepi.22001/full>)

This year's abstract topics include:

- Biobanks and large-scale populations
- Common disease genetics (incl. cancer genetics, heart disease genetics)
- Diversity and studies of multiple ancestries
- Immunogenetics (incl. COVID-19 and SARS-CoV-2)
- Machine learning and data science
- Mendelian randomization and causality
- OMICs: methods and applications
- Polygenic risk scores and prediction (and general polygenic trait genetics)
- Statistical modeling in genetic epidemiology (GxE, GxG, Multi-trait and summary-statistic methods)
- Whole genome and exome sequencing studies

Guidelines:

- Abstracts should be named as SURNAME-FIRSTNAME.DOC (eg. Doe-John.doc).
- **Please do not submit PDF files.**
- **Abstracts must be 250 words or fewer.**
- The first line of the document should provide a short, concise title in bold font. Max title length of 150 characters including spaces.
- The next line should give the name of authors: First name, middle initial followed by a period, and last name (*example: John E. Doe*). Names are to be separated by commas, affiliations noted by numbers and with an asterisk * to specify the presenting author. The maximum number of authors is 10.
- The next line should begin a list of affiliations.
 - **US affiliations** should be spelled out and should not be abbreviated. Presenting author should be marked with an asterisk and institutions should be italicized. Example:
John E. Doe ^{1,2}, Jack B. Doe ^{2*}
¹Department of Genetics, University of North Carolina at Chapel Hill, Chapel Hill, North Carolina, United States of America; ²Lineberger Comprehensive Cancer Center, University of North Carolina at Chapel Hill, Chapel Hill, North Carolina, United States of America
 - **International affiliations** should be spelled out and should not be abbreviated. Presenting author should be marked with an asterisk and institutions should be italicized. Example:
John E. Doe ^{1,2}, Jack B. Doe ^{2*}
¹Department of Genetics, University of Liverpool, Liverpool, United Kingdom; ²Lady Davis Institute for Medical Research, Jewish General Hospital, Montreal, Canada
- Content-specific style points
 - Italics: Use for Genus/species, emphasis, journal titles in text, *P* (probability) should read *P* value and not *p*-value or *P*-value, and words or terms being defined. Do not use for Latinisms (e.g., et al., in vivo, in vitro).
 - Abbreviations: **First mention of acronyms and abbreviations in the text should be spelled out in full with acronym or abbreviation in parentheses immediately following.** In general, except for units of measure and defined acronyms and abbreviations, there should be no abbreviations unless within brackets or parentheses. See below for a list of acronyms and abbreviations that do not need to be defined.
 - Manufacturers: Insert city and state or country in parentheses.
 - Numbers: Spell out numbers 1–9 and numbers that begin a sentence.
 - % (only when used with a specific number; otherwise spell out).
 - vs. (both in text and in parentheses).
 - African American population (African American --- two words as noun and adjective).

- Change while to although.
- Change since to because.
- Change df/d^4f to df/d^4f .
- N-dashes – use only for ranges and do not use in between words in text
- References should not be included in the abstract.
- The following items are collected via the online submission form and should not be included in the abstract:
Corresponding/submitting author contact information and Topic categories (in place of keywords)
- **Permitted abbreviations:** min, hr, sec, log (one word, roman, lc), g, mg, kg, *P* (CAPITALIZED and italic).
- **Common acronyms that do not need to be spelled out:**

SD: standard deviation	RR: relative risk
SE: standard error	LOD score: logarithm of odds score
CI: confidence interval	SNP/SNP's: single nucleotide polymorphism's
OR: odds ratio	