incidence and risk factors of oesophageal cancer by histological subtypes using data from 178 countries.

Methods The data on the incidence of oesophageal cancer by histological types in 2018 were estimated from GLOBALCAN and Cancer Incidence in Five Continents (CI5). Age-standardized rates (ASRs) for oesophageal cancer incidence by histological subtypes were evaluated by Segi–Doll population. The prevalence of tobacco use, alcohol drinking, physical inactivity, obesity, diabetes, and lipid disorders for each country were retrieved from the Global Health Observatory. The association between the ratio of histological subtypes and risk factors was examined by multivariable linear regression.

Results We estimated a total of 63,470 (12.6%) and 502,669 new cases of oesophageal adenocarcinoma (AC) and squamous cell carcinoma (SCC) in 2018, respectively. The incidence among males was 3.6-fold and 2.2-fold of that among females for AC and SCC, respectively. The highest AC:SCC ratio was found in the UK (ratio 2.880, ASR 7.5), New Zealand (2.667, 4.2), the Netherlands (2.536, 7.7), Bahrain (2.143, 0.9), and Canada (2.000, 3.7) among males (figure 1). As for females, the highest AC:SCC ratio was observed in Moldova (1.000, 0.2), the Netherlands (0.800, 1.2), Iceland (0.750; 0.5), the UK (0.700; 1.4), and Cyprus (0.667; 0.3). A higher AC:SCC ratio was associated with a higher prevalence of obesity (male: β 0.039, 95% CI 0.023 to 0.055; female: 0.009, 0.004 to 0.146) and high cholesterol (male: 0.028, 0.010 to 0.047; female: 0.011, 0.004 to 0.019); but a lower prevalence of tobacco use (male: -0.007, -0.014 to -0.001) and diabetes (male: 0.009, 0.004 to 0.146; female: -0.021, -0.038 to -0.003).

Conclusions While SCC is the predominant subtype of oesophageal cancer, the incidence of AC has surpassed SCC in a substantial proportion of countries, probably due to the increasing prevalence of obesity and metabolic disorders. Future research should investigate the reasons behind these epidemiological changes.

IDDF2020-ABS-0141 THE GUT MICROBIOME AND SERUM METABOLOME ORCHESTRATE HEALTHY AGING AND LONGEVITY WITH NOVEL IMPLICATIONS FOR RENAL FUNCTION

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10.1136/gutjnl-2020-IDDF.87

Background Aging is closely associated with the occurrence and development of many complex diseases such as cardiovascular conditions, type 2 diabetes, gastrointestinal (GI) and renal diseases. Detecting changes as early as possible in the human body due to aging is thus crucial significant to potentially mitigate its impacts on physical health.

Methods Here, we present a comprehensive metagenome association study and serum metabolomics profiling in a registry Guangxi-Longevity cohort aged from 20–111 years ($n=151$) and Kunming cohort ages ranging from 20–80 years ($n=80$).

Results We identified uremic toxins as key factors in serum metabolomics highly associated with aging, and this finding has been validated in an independent Kunming-Aging cohort aged from 20–80 years ($n=80$). We also observed that aging-associated systemic inflammation levels were positively associated with uremic toxins. Moreover, the increased *Escherichia coli*, *Odoribacter splanchnicus*, *Bilophila wadsworthia* and *Parabacteroides* spp. abundances were related to serum levels of uremic toxins, and the accumulating rate of uremic toxins and specific microbial species was robustly much slower in centenarians than in nonagenarians. We further investigated the frailty and health status in long-living individuals aged above 90 years, and found that the frailty status might be a putative extreme aging phenomenon characterized by novel uremic toxin accumulation patterns.

Conclusions Our findings reveal novel potential links between gut microbiota alterations, uremic toxins and aging, and highlight the preponderance of gut microbiota and serum metabolism in aging.

IDDF2020-ABS-0142 PERFORMANCE INDICATORS OF ORGANISED COLORECTAL CANCER SCREENING PROGRAMMES USING FAECAL IMMUNOCHEMICAL TESTS AND COLONOSCOPY: A SYSTEMATIC REVIEW AND META-ANALYSIS

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10.1136/gutjnl-2020-IDDF.88

Background The success of population-based colorectal cancer (CRC) screening is dependent on the optimal achievement of various targets. We estimated the pooled performance indicators of CRC screening programmes that used faecal immunochemical tests (FIT) as a primary screening modality and colonoscopy as a subsequent confirmatory test in various countries.

Methods We searched PubMed, Ovid MEDLINE, Embase, and Cochrane from inception to 1st Jan 2020. We included original articles published in the English language describing population-based CRC screening programme that used FIT and colonoscopy, and relevant national CRC screening reports by hand searching. We extracted data to pool early performance indicators, including participation rate, invalid FIT rate, FIT positive rate, adenoma/CRC detection rate of FIT, colonoscopy compliance rate, rate of adequate bowel preparation, colonoscopy completion rate, colonoscopy complication rate, and positive predictive values (PPV) of FIT for adenoma/CRC. We used Metaprop to conduct a meta-analysis via R software (version 3.6.3). The Freeman-Tukey double arcsine transformation was used to stabilise the variances, and a random-effects model was used to pool the rates with proportions.

Results A total of 85 studies (64 articles and 21 reports) were included in this meta-analysis. The pooled participation rate was 52.2%, and the pooled proportion of invalid FIT was