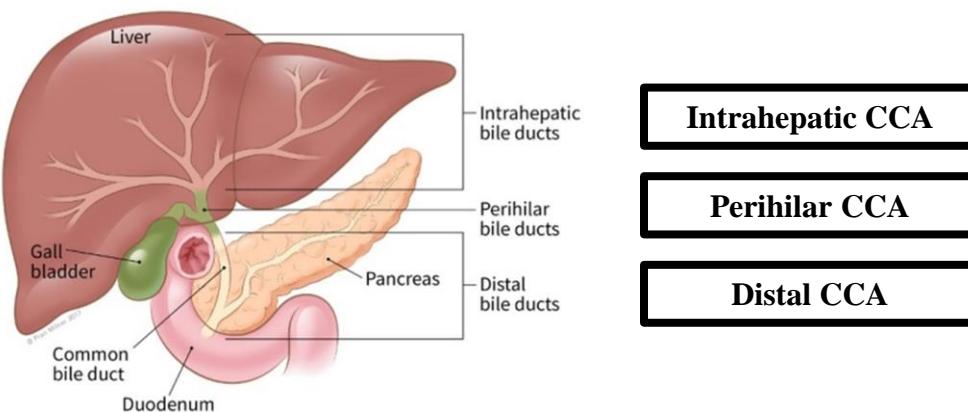


Introduction

- Cholangiocarcinoma (CCA) is the second commonest form of liver cancer worldwide and arises from a malignant transformation of cholangiocytes (epithelial lining of the bile ducts/ductules within the liver).
- Rising incidence and mortality rates, combined with a dismal prognosis (5-year survival < 10%) make this an important clinical problem to be addressed.
- Recognized risk factors for CCA, including chronic Hepatitis B or C infection, liver fluke infection and autoimmune-related diseases such as Primary Sclerosing Cholangitis, all point to an immune component in the pathogenesis of this disease. However, immune-related mechanisms of disease pathogenesis and progression in CCA remain largely understudied.
- A better understanding of the immune tumour microenvironment (TME) in this disease is necessary to enhance development of new treatment and prognostication strategies for patients with CCA.
- We therefore aimed to investigate the potential contribution of immune cells to CCA pathology.



Courtesy American Cancer Society

Figure 1: Anatomic classification of CCA

Methods

- Using immunohistochemistry, we first analyzed formalin-fixed, paraffin-embedded CCA tumours to investigate the presence of tumour-infiltrating immune cells in human CCAs (n=20).
- We next conducted a systematic review and meta-analysis of existing literature in accordance with the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines^{1,2}, to determine the correlation between immune cell presence (assessed histologically) and clinical outcomes in patients diagnosed with CCA. Also see Systematic Review and Meta-Analysis Protocol on next slide.
- Finally, using NanoString™ Digital Spatial Profiling (DSP), we preliminarily undertook spatially-resolved analysis of immune protein signatures within CCA tumours (n=2).

References

1. Liberati A, et al., The PRISMA statement for reporting systematic reviews and meta-analyses of studies that evaluate healthcare interventions: explanation and elaboration. *BMJ*. 2009;339:b2700.
2. Egbuniwe IU, et al. 2020. PROSPERO Int. Prospect. Regist. Syst. Rev

Results

- Our results demonstrate the following:
- Preferential accumulation of CD68⁺ macrophages and CD3⁺ T lymphocyte subsets within CCA tumours compared with non-tumour bile duct tissue.
 - Presence of intra-tumoural CD68⁺ macrophages and CD3⁺ CD8⁺ (Cytotoxic) T lymphocytes are most significantly associated with overall patient survival in patients diagnosed with CCA.
 - Prognostic significance of intra-tumoural CD68⁺ macrophages appears dependent upon spatial location within CCA tumours.
 - Corroboration of differential presence of CD68⁺ macrophages within distinct tumour regions by quantification analysis.
 - Distinct clustering patterns between CD68⁺ macrophages and the immune checkpoint molecule B7-H3 using cluster analysis.

Macrophages preferentially accumulate within CCA tumour tissue compared to non-tumour bile duct tissue

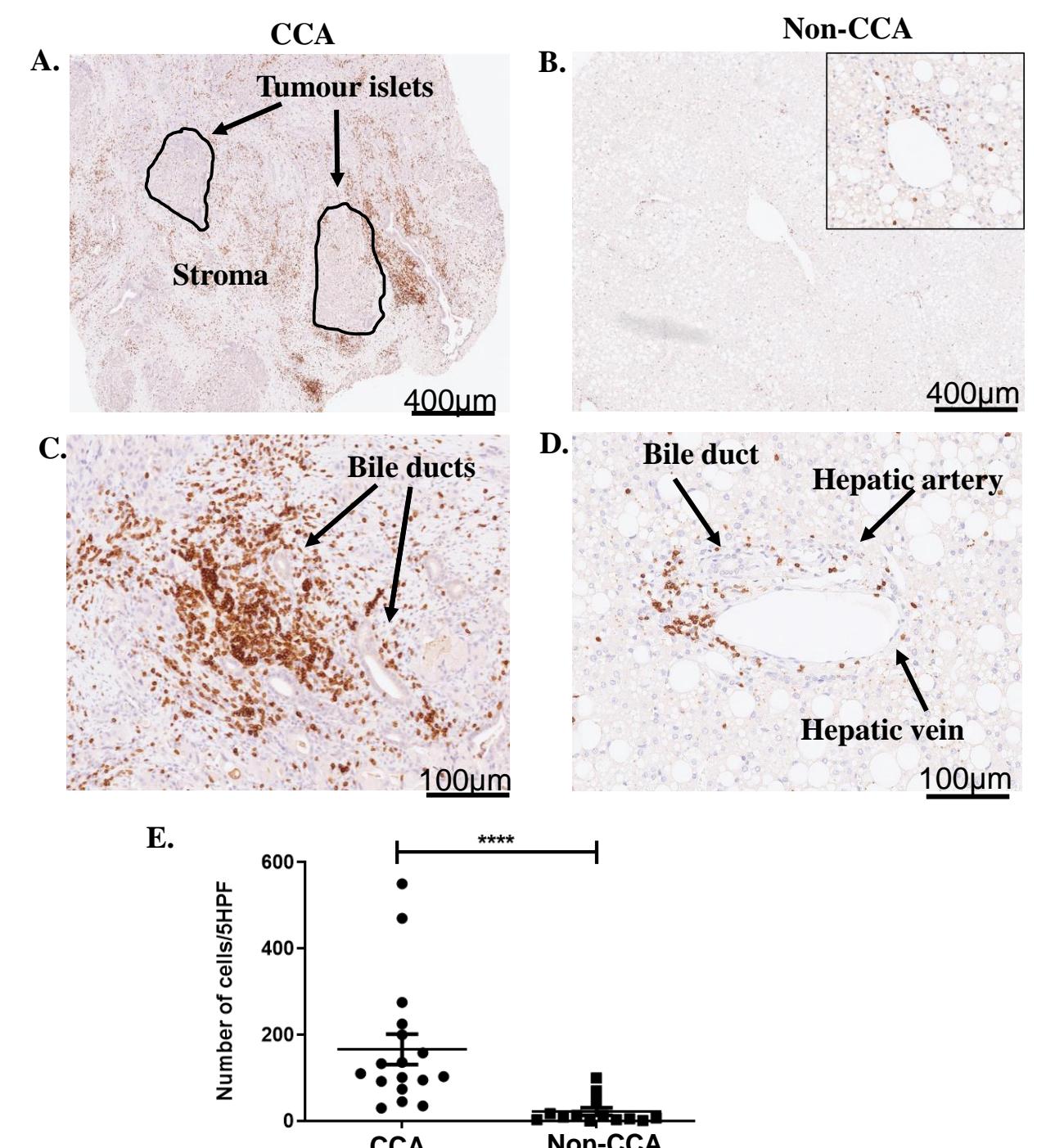


Figure 2: Immunohistochemical staining of CCA tumours (A and C), and matched non-tumour bile duct tissue (B and D) shows tumour-infiltrating CD68⁺ macrophages (brown dots) within tumour islets and stroma. Comparatively fewer CD68⁺ macrophages are present within non-tumour tissue obtained from the same patient sample. (E) Quantification of CD68⁺ macrophages within CCA tumour and non-tumour tissue. P<0.0001; Mann-Whitney U Test; Error bars = Mean ± SEM.

Results

T lymphocyte subsets are also enriched within CCA tumour tissue compared to non-tumour bile duct tissue

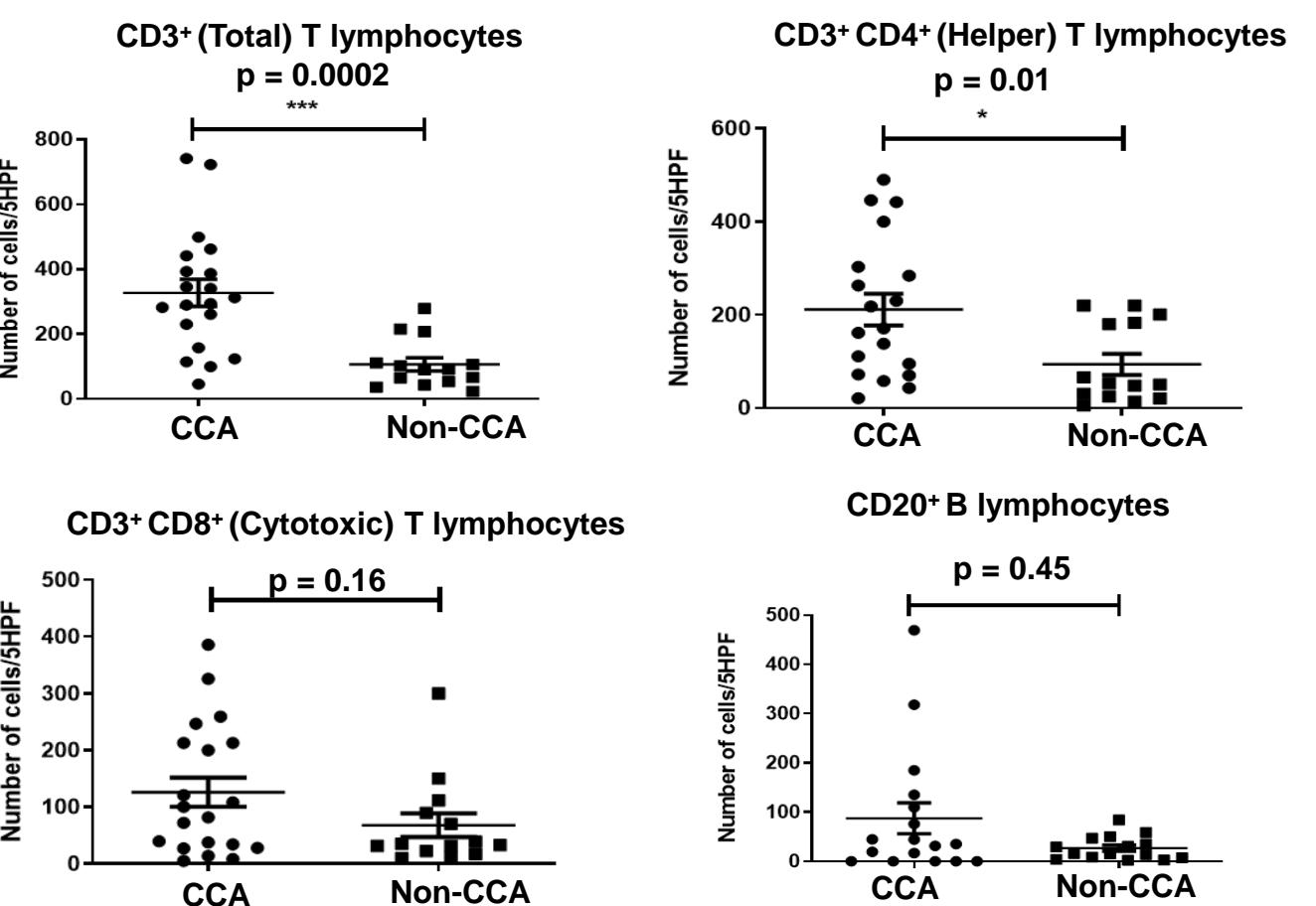


Figure 3: Increased numbers of total (CD3⁺), helper (CD3⁺CD4⁺) and cytotoxic (CD3⁺CD8⁺) T lymphocytes are present within CCA tumours compared to non-tumour bile duct tissue.

Prognostic significance of intra-tumoural macrophages is dependent upon location within CCA tumours

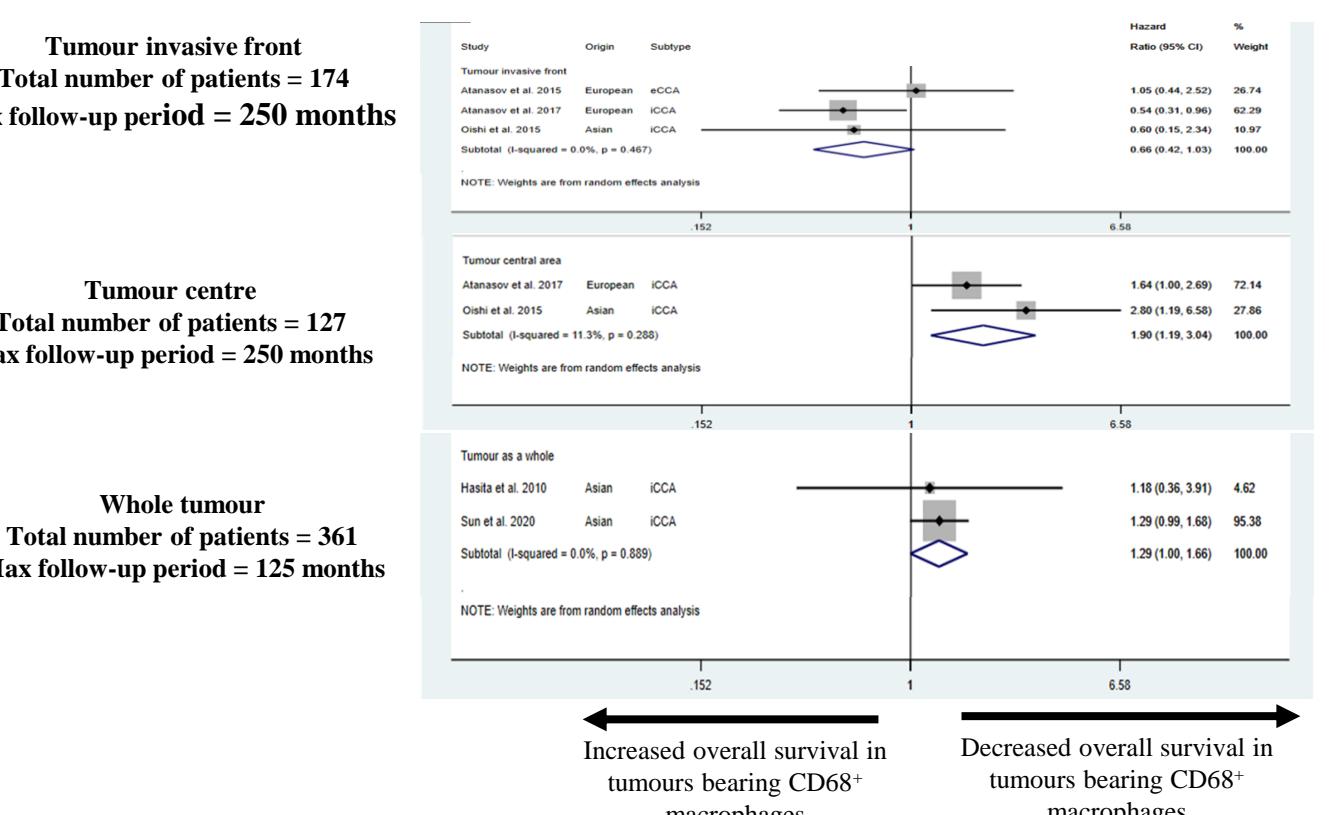


Figure 4: Forest plots show correlation between intra-tumoural location of CD68⁺ macrophages and overall survival in patients diagnosed with CCA. Measure of Effect = Pooled Hazard Ratio; Measure of Statistical Significance = 95% Confidence Interval (CI).

Results

Quantification and Cluster analyses corroborate differential presence of CD68⁺ macrophages within distinct tumour regions

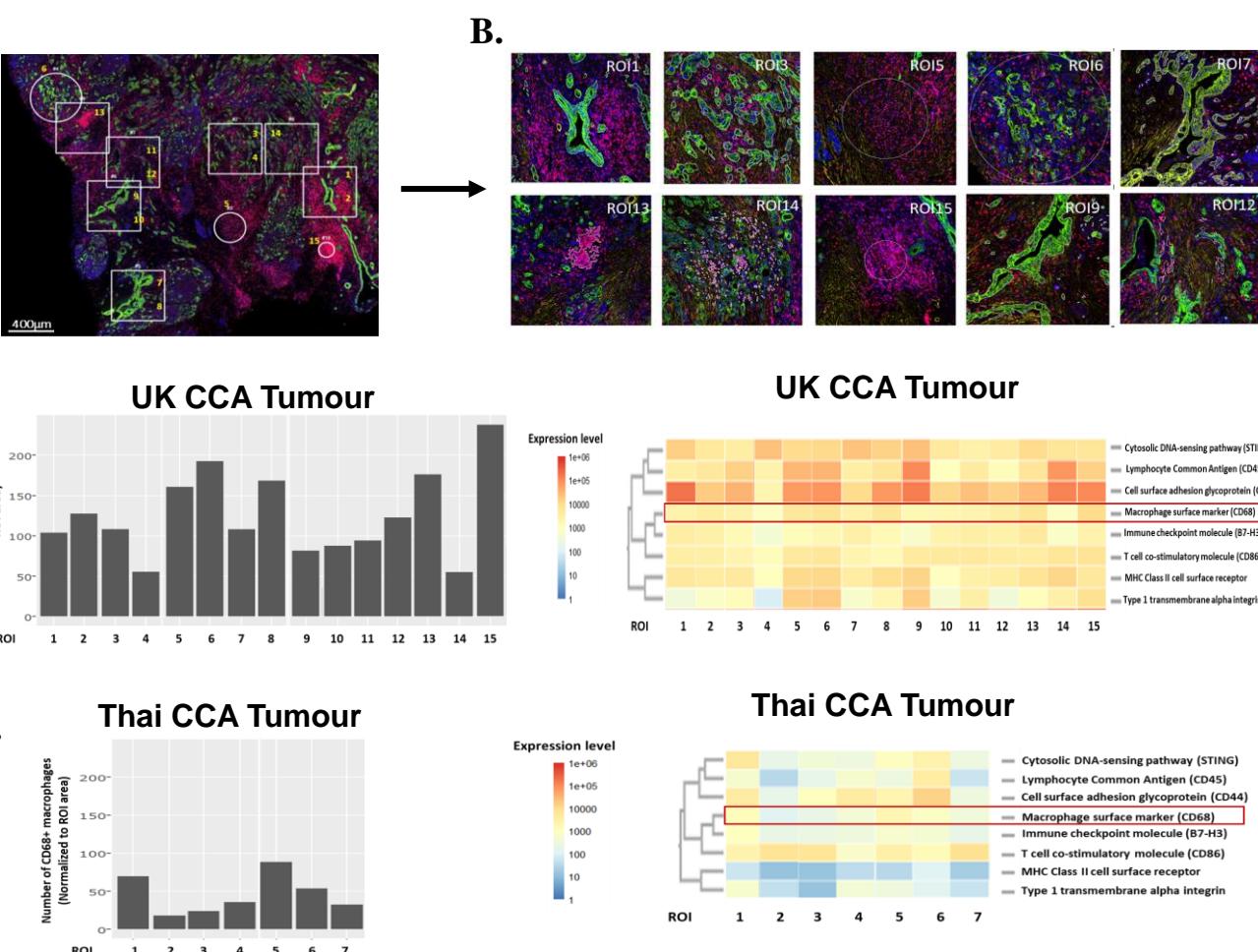


Figure 5: Patient CCA tumours were stained to identify tumour cells (green = pan-cytokeratin), T cells (red = CD3), stromal cells (yellow = smooth muscle actin) and cell nuclei (blue = DAPI stain). A and B = images of tumour obtained from a UK patient. Regions of interest (ROIs - white boxes/circles) representing tumour-rich, immune-rich or stroma-rich areas were further probed to generate quantification graphs and heatmaps (C and D; left and right panels respectively) of stromal and immune signatures. Differential presence of CD68⁺ macrophages is seen within different regions in CCA tumours from one UK (C) and one Thai (D) patient. Heatmaps also show clustering of CD68⁺ macrophages with immune checkpoint molecule (B7-H3) within both tumours.

Conclusions

- CD68⁺ macrophages and CD3⁺(Total) T lymphocytes are enriched within CCA tumours compared to non-tumour tissue.
- CD68⁺ tumour-infiltrating macrophages correlate with overall survival in patients with CCA.
- Prognostic significance of intra-tumoural CD68⁺ macrophages appears dependent upon location within CCA tumours – likely related to underlying intra and inter-tumour heterogeneity
- Findings highlight the need to consider spatial context in the evaluation of tumour specimens in research and diagnostic/prognostic endeavours.

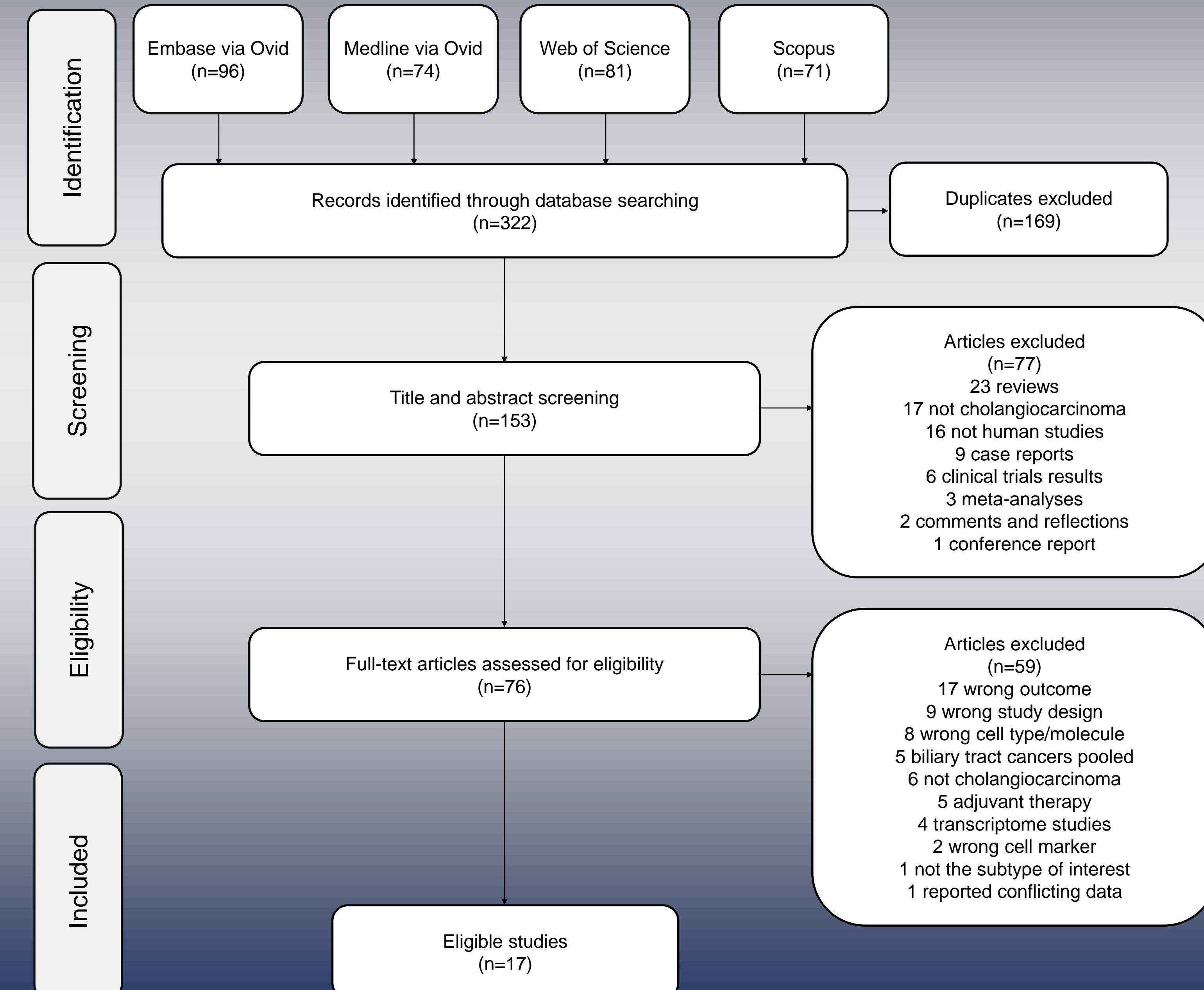
I.U Egbuniwe¹, U. Rastović¹, M. Persson², K.J Hunter³, J. Gong⁴, A.M Grabowska¹, D.O Bates¹, PS Jayaraman¹, K.Gaston¹

1. Division of Cancer and Stem Cells, Centre for Cancer Sciences, University of Nottingham, UK
2. Centre of Evidence Based Dermatology, University of Nottingham, UK
3. Cancer Research UK Birmingham Centre, University of Birmingham, UK
4. NanoString Technologies, Seattle, USA

*Corresponding author – Dr Iсиoma U. Egbuniwe, Division of Cancer and Stem Cells, Biodiscovery Institute, University of Nottingham, NG7 2RD UK; isioma.egbuniwe@nottingham.ac.uk

Systematic Review and Meta-analysis Protocol

- Study conducted following PRISMA guidelines¹
- Protocol registered on PROSPERO database²
- Study screened using Rayyan QCRI web tool (<https://rayyan.qcri.org/welcome>)
- Data extraction/Inclusion criteria:
 - Excel spreadsheet
 - Histopathological assessment
 - Meta-analysis of pooled hazard ratios for overall survival (OS)
- Risk of Bias Assessment conducted using QUIPS tool³



References

1. Liberati A. et al. 2009. *BMJ*. 339
2. Egbuniwe IU. et al. 2020. *PROSPERO Int. Prospect. Regist. Syst. Rev*
3. Hayden JA. et al. 2013. *Ann. Intern. Med.* 158:280-286