Agile review of the literature using a structured, non-systematic approach: example application for a genomic diversity question Field P,^a Couto CA-M,^a Schofield H,^a Eichinger C^a and Mackintosh M^b ^aOxford PharmaGenesis, Oxford, UK; ^bGenomics England, London, UK

Introduction

 When asking broad research questions that cover a large evidence base, with differences in keywords and terminology between studies, a systematic literature

Methods

 We followed an iterative process (Figure 2), with semi-reproducible methodology (Figure 3).

Results

 Our initial search strings identified references with low specificity; agile reviewing led to approximately 200 references being reviewed in detail (Figure 4).

- review (SLR) can take more time and resources than are available.
- There is little guidance on semi-reproducible, non-systematic alternatives and no guidance, to our knowledge, on supplementing targeted Population, Intervention, Comparison, Outcomes and Study-based searches with artificial intelligence (AI)-driven searching.

Objectives

 We wanted a methodology that would help us to identify key studies without having to screen a large number of citations and without already knowing the areas of focus in the research area (Figure 1).

Figure 1. Aims of our agile methodology.

We wanted to identify key studies

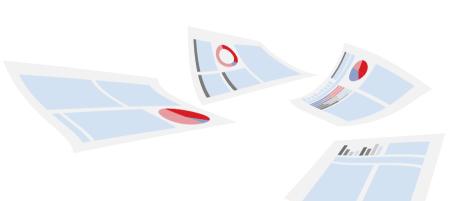
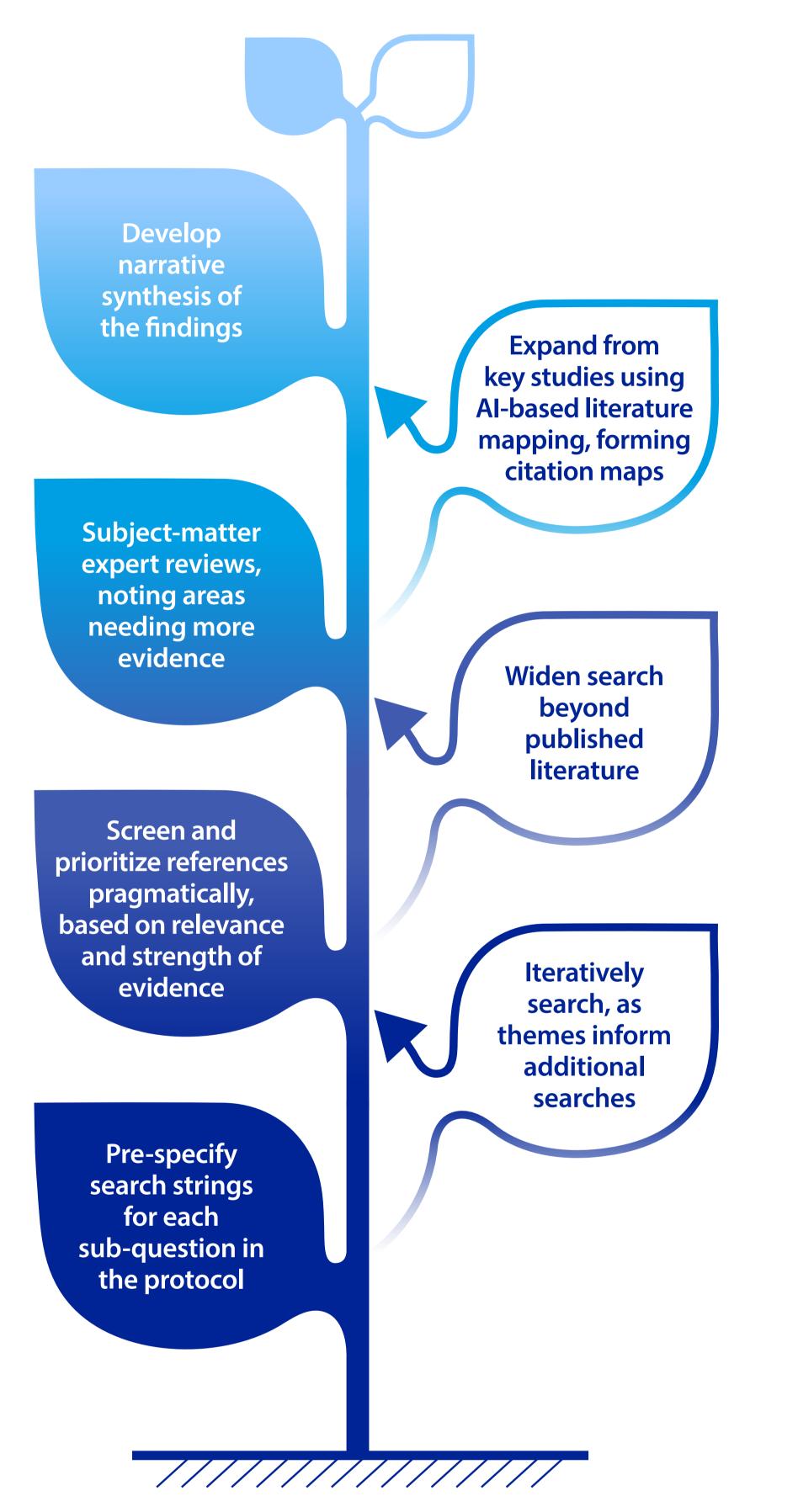


Figure 2. Process.



- The combination of PubMed and Google searches and AI-based mapping had greater sensitivity and specificity than any of these techniques alone.
- Citation mapping was able to deepen evidence on particular subjects by providing example case studies; however, it was less useful for uncovering new topics and, if used alone, it would not have covered the breadth of studies.

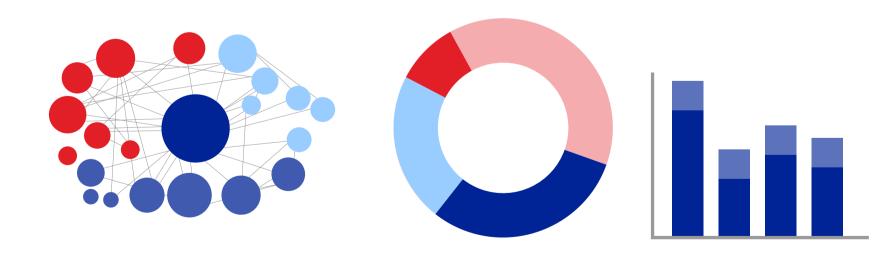
Figure 4. Results summary showing the number of references identified or reviewed.





... without having to screen a very large number of citations ...

... so we could understand and summarize leading research themes.



• We were willing to lose some sensitivity – we did not have to identify all relevant studies.

AI, artificial intelligence.

Figure 3. Key features.



500



Identified through AI

strings



Reviewed in detail

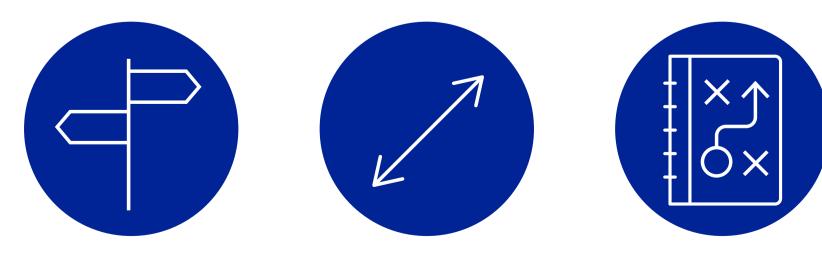
AI, artificial intelligence.

Conclusions

- Agile literature reviewing efficiently orientated us around a complex evidence landscape.
- We prioritized understanding the breadth of information but maintained an element of reproducibility from the initial search terms.
- Findings from our agile approach could help to design SLRs into specific themes.

We wanted a methodology somewhere between one that was fully reproducible and one that was fully pragmatic.

 We formalized an agile literature review methodology as an alternative to an SLR and applied this to the complex evidence landscape for how diversity is defined in genomics research.



Inclusion/exclusion Expanded to criteria selection similar publications

PICOS, Population, Intervention, Comparison, Outcomes and Study.

Disclosures

Polly Field, Anne-Marie C Couto, Helen Schofield and Christian Eichinger are paid employees of Oxford PharmaGenesis Ltd. Polly Field owns shares in Oxford PharmaGenesis Ltd. Maxine Mackintosh is a paid employee of Genomics England.

Funding

Narrative

synthesis

The development of the methodology and the development of this poster were funded by Oxford PharmaGenesis. The literature review was funded by Genomics England.



Poster presented at the International Society for Pharmacoeconomics and Outcomes Research (ISPOR) Europe 2022 conference, 6–9 November 2022, Vienna, Austria and virtual