

- 1. Proposal GTFCC Laboratory WG:
Revise classification of *Vibrio cholerae***
- 2. Country Support - Quality Assurance**
- 3. Do we need to compose document on
Accessing Sequence Data?**

1. Proposal GTFCC Laboratory WG: Revise classification of *Vibrio cholerae*

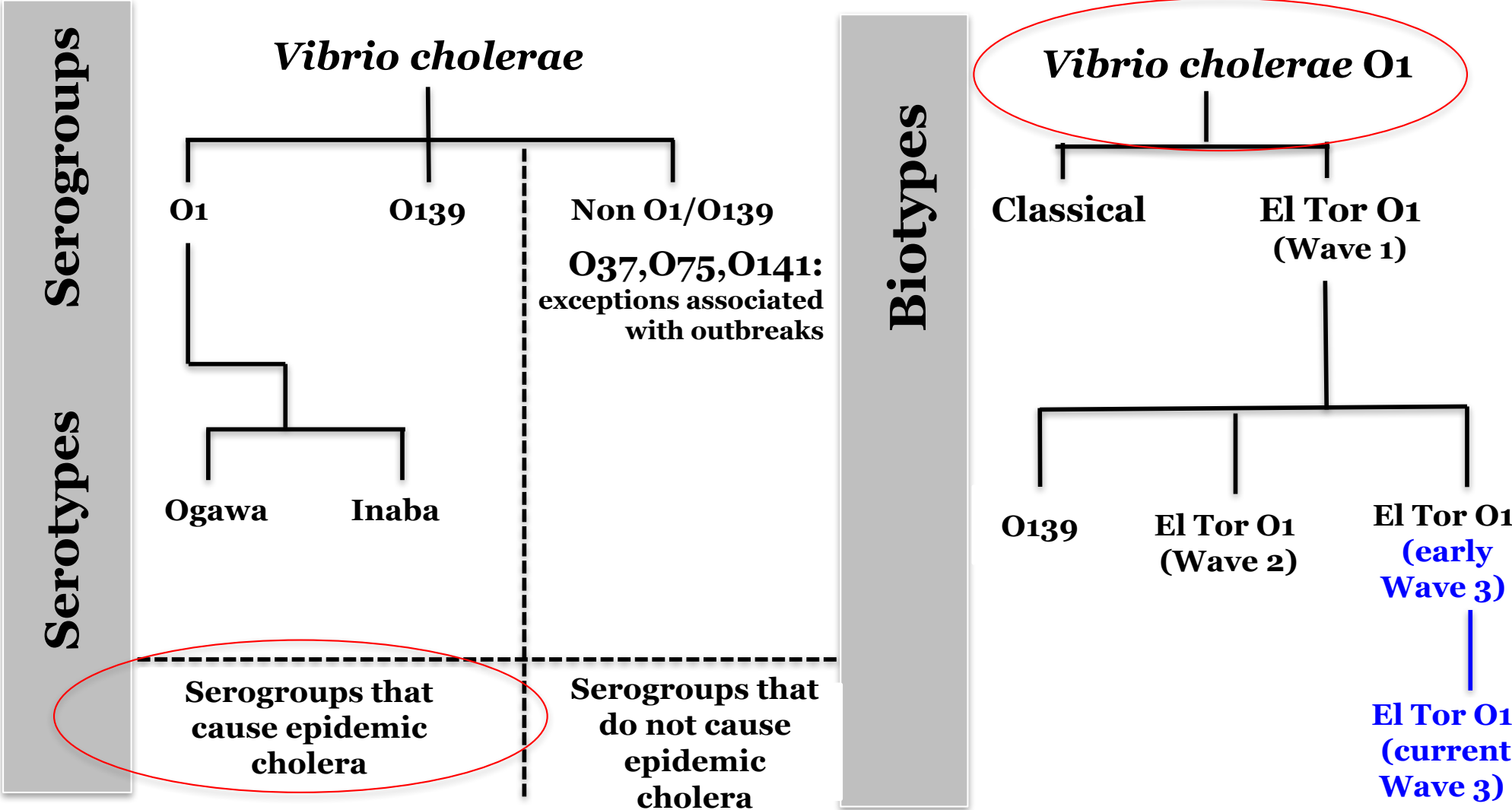
For discussion:

Simplify to

***7th Pandemic El Tor lineage *Vibrio cholerae*
(subsumes O1, O139, CTX variants)***

Requires greater socialisation and specific consultation with GB Nair and DW Kim

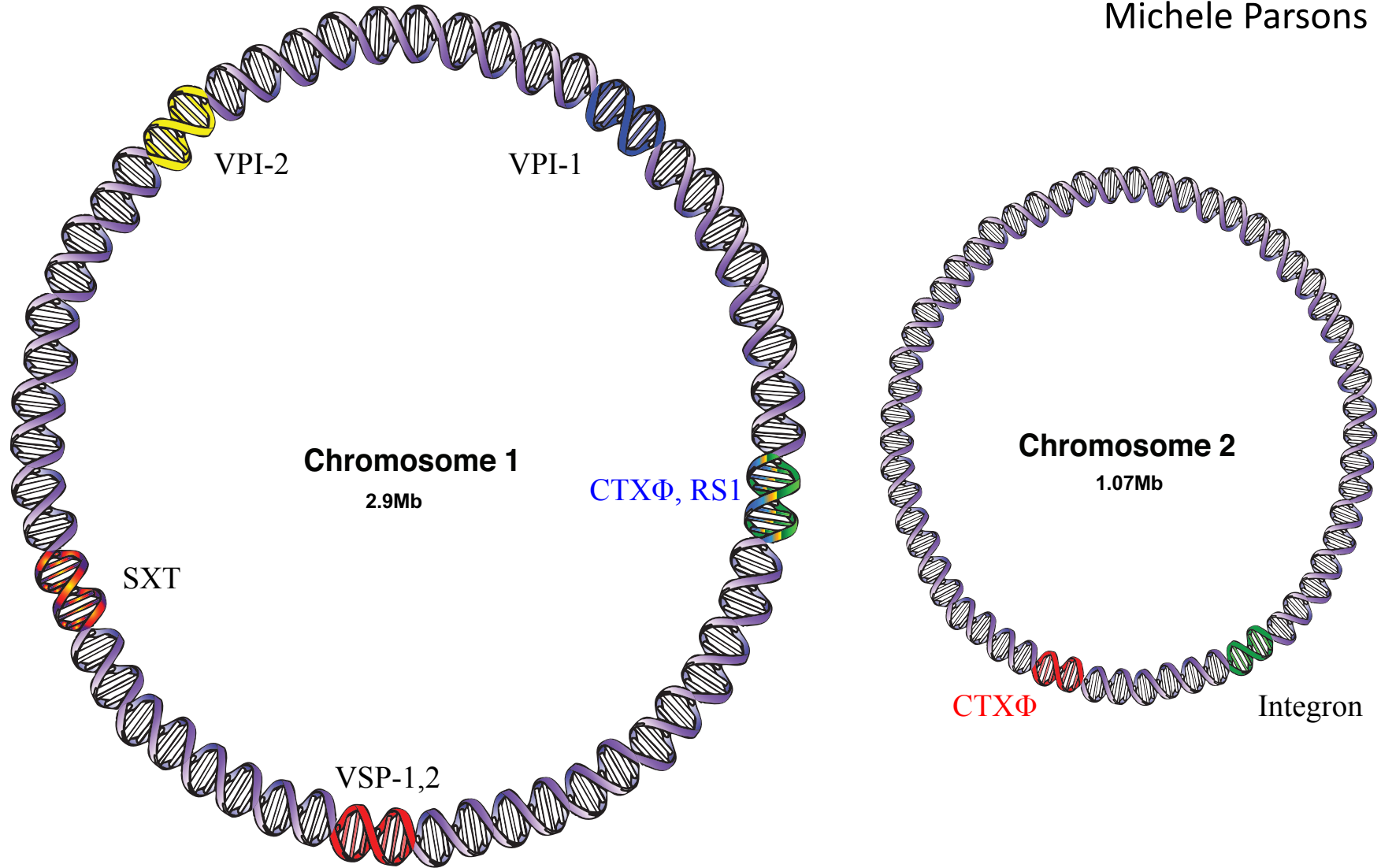
Previously 2015: Revised classification of *Vibrio cholerae*



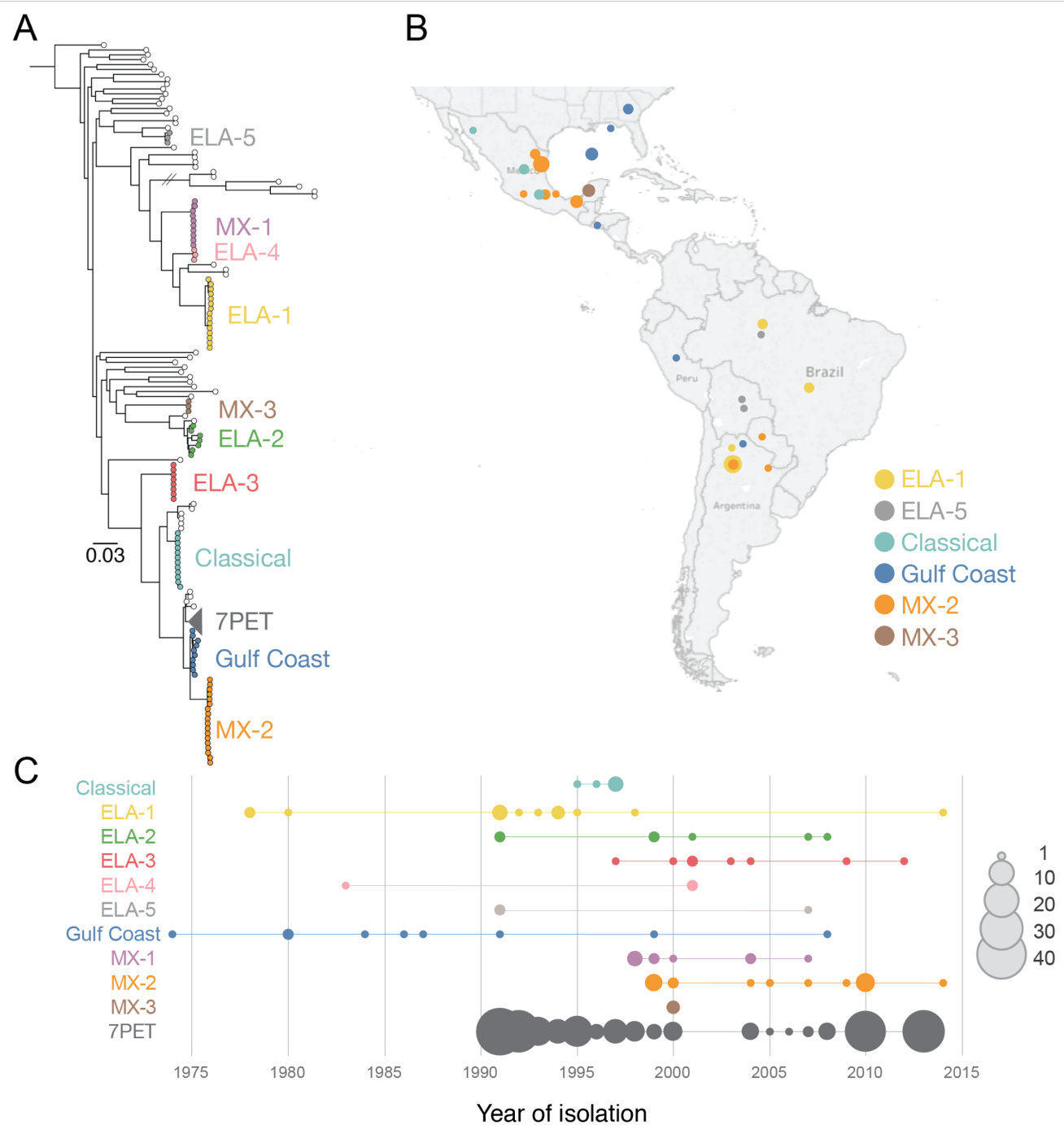
Dong Wook Kim & G. Balakrish Nair

Vibrio cholerae genome

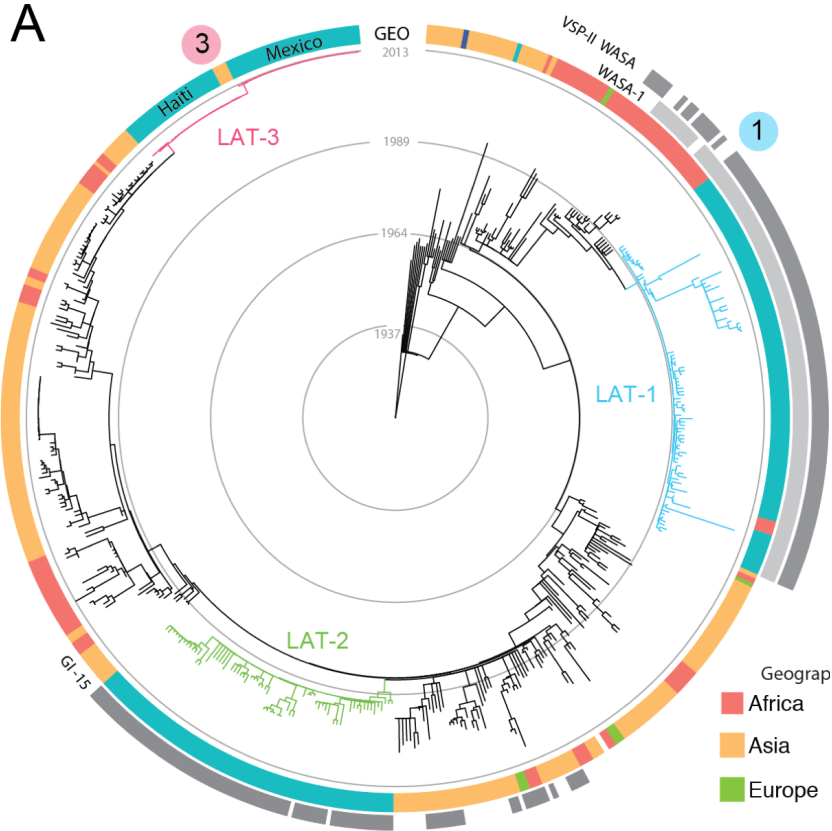
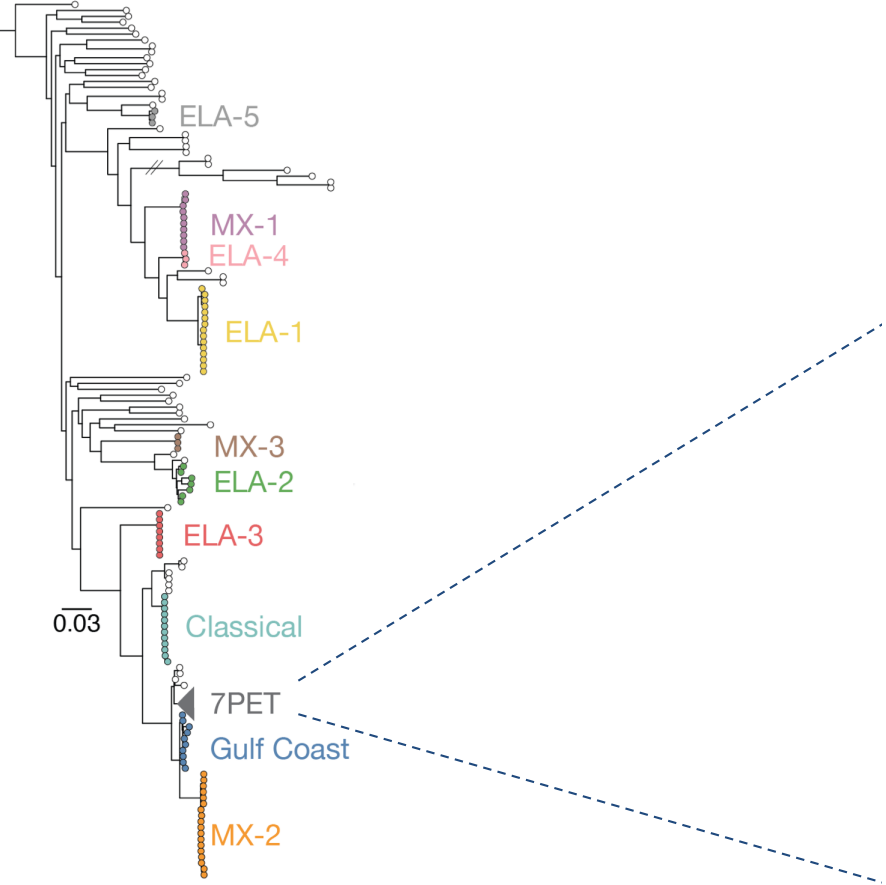
Michele Parsons



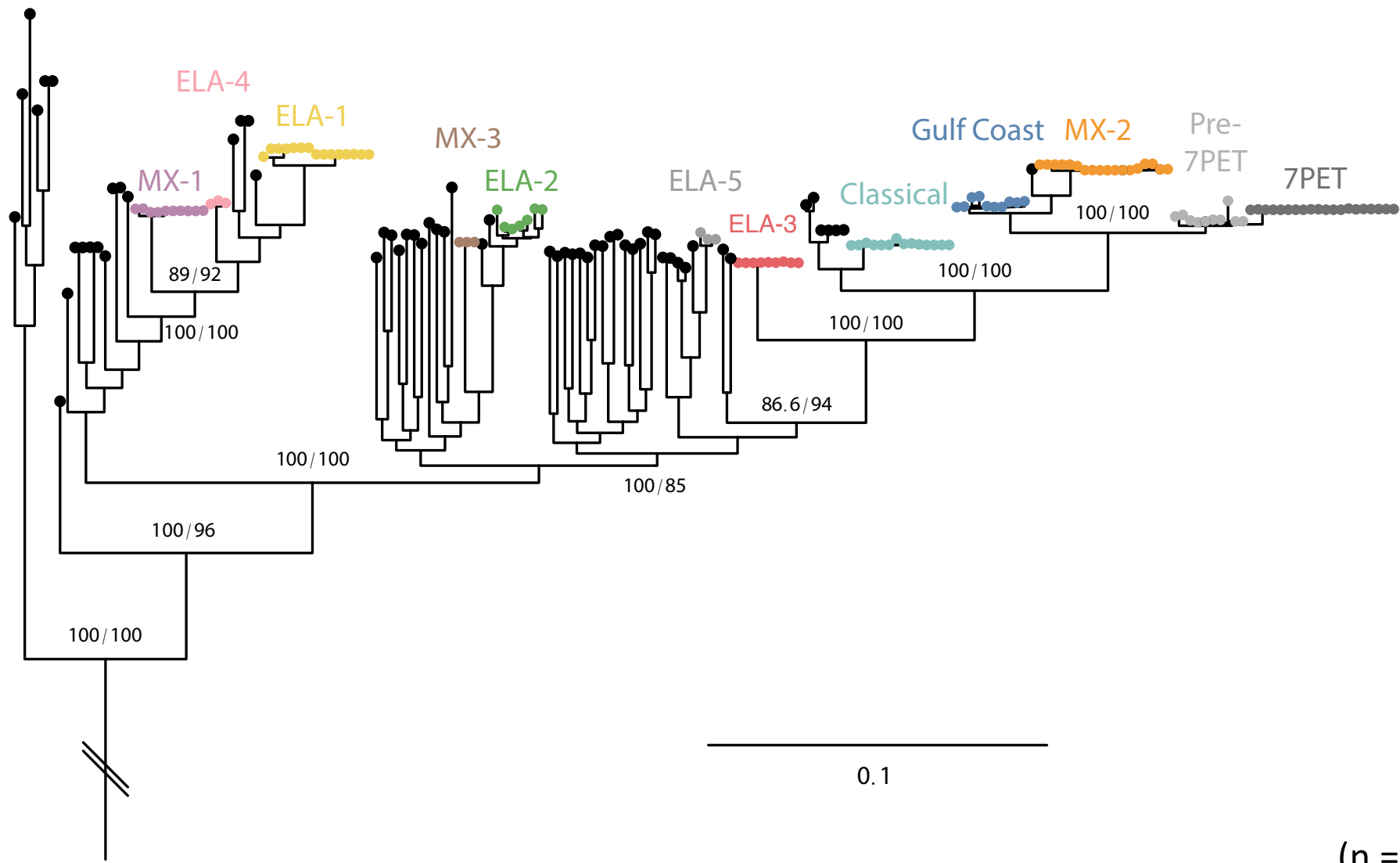
Qu: What information is actually needed and by Whom?



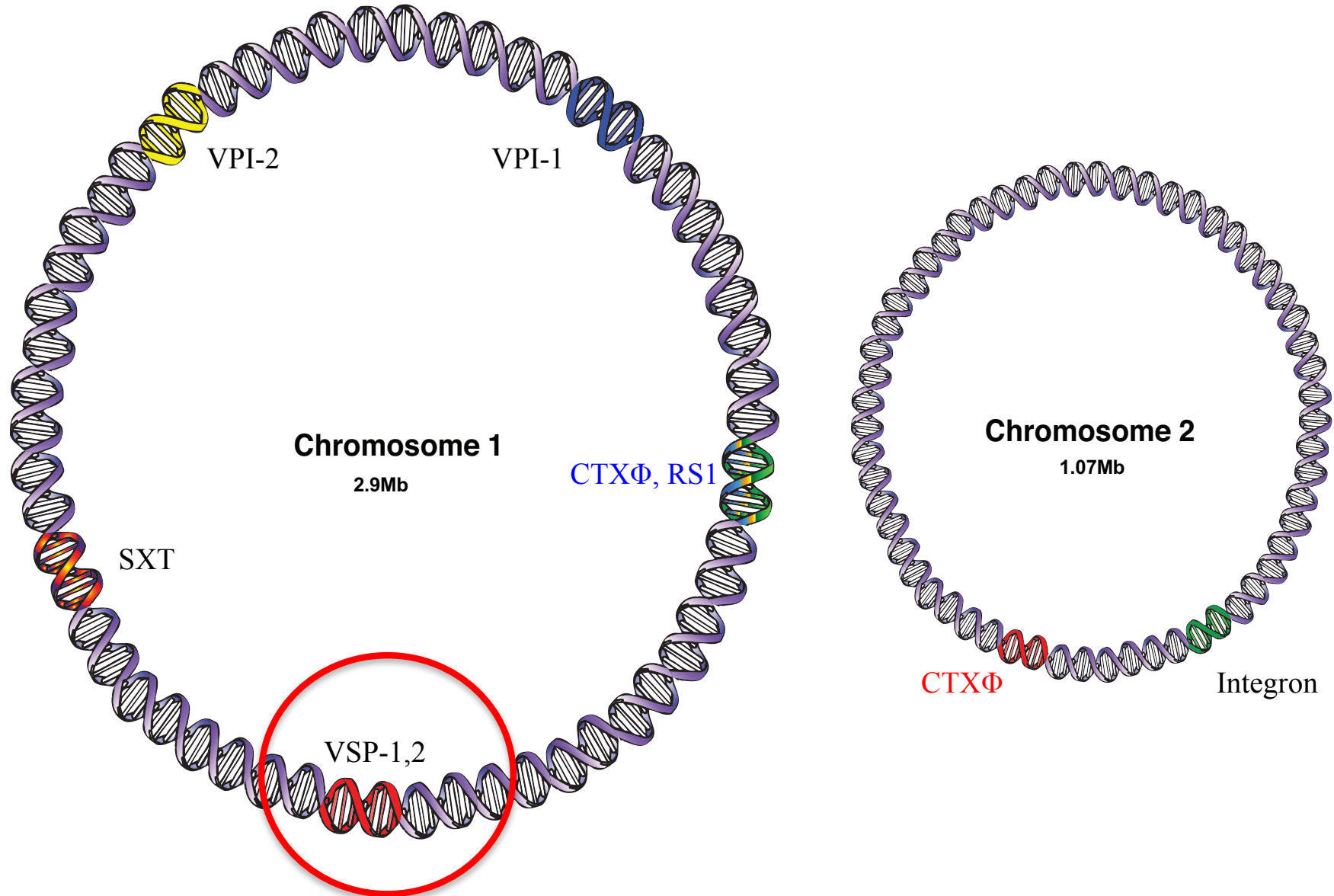
7th Pandemic El Tor lineage *Vibrio cholerae*



Domman et al., Science 2017



Vibrio cholerae genome



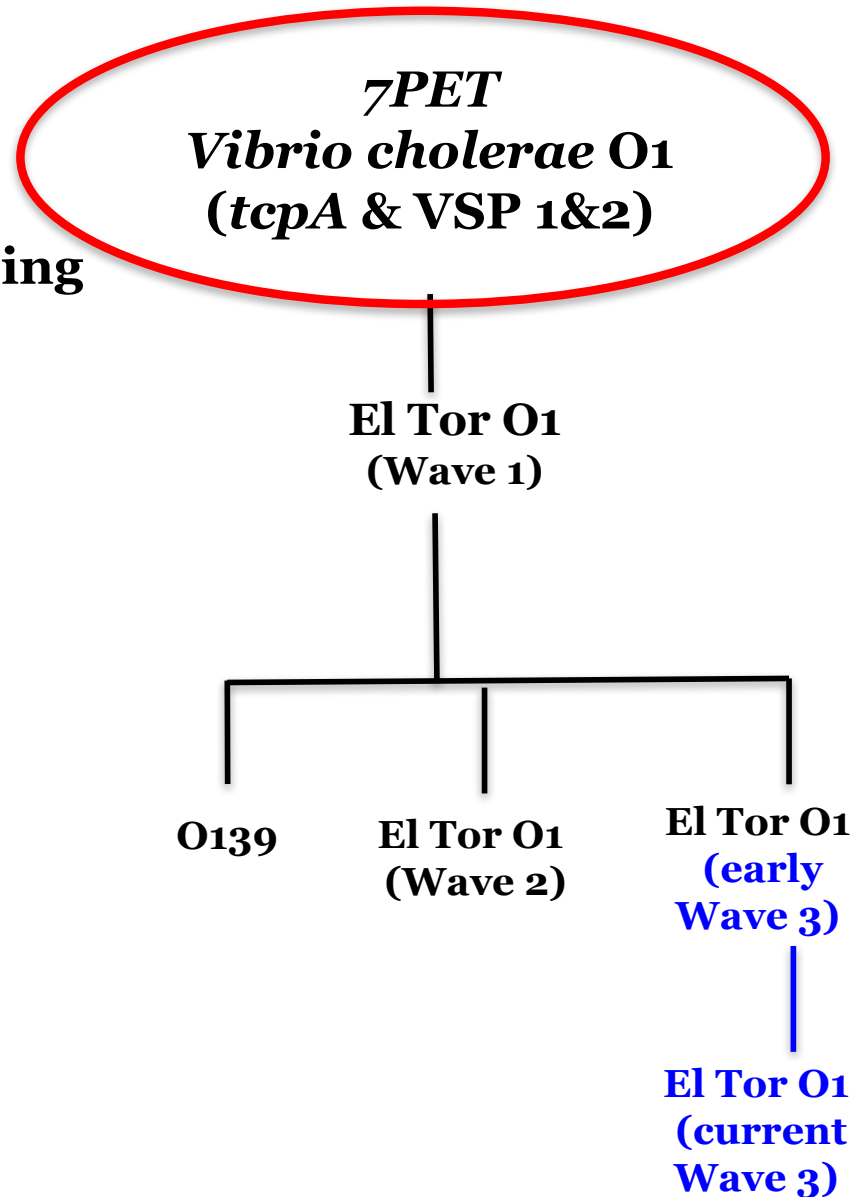
Proposal:

Genotypes

CTXB

Non epidemic genotypes

Classical *tcpA* - Typing



What is the minimum useful information required in field?