

Mapping Maternal Group B Streptococcus Colonization in Nepal: First Steps Toward Surveillance to Inform Vaccine Policy

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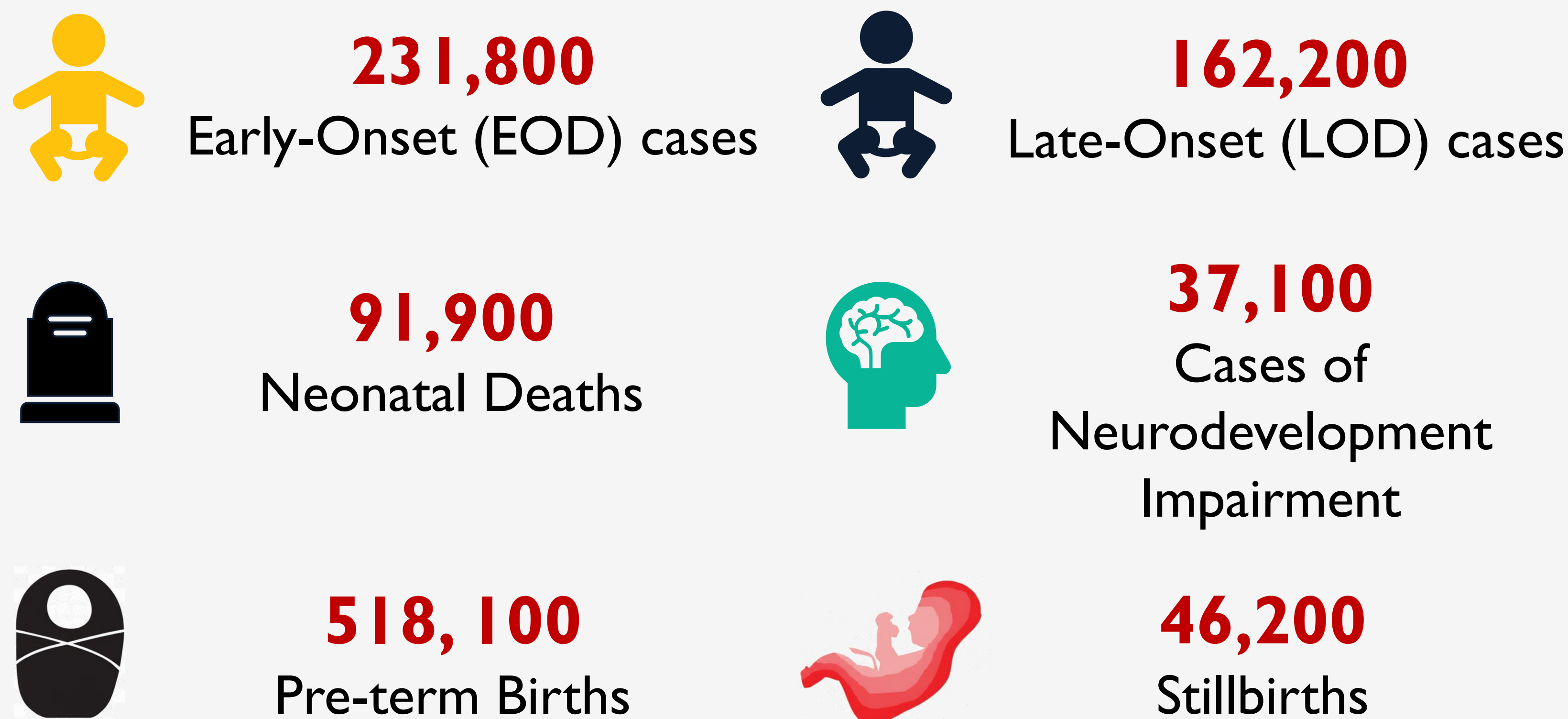
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BACKGROUND: SIGNIFICANCE OF GROUP B STREPTOCOCCUS (GBS)

ANNUAL BURDEN OF GBS¹



CURRENT PREVENTION STRATEGY²

Intrapartum Antimicrobial Prophylaxis (IAP)

Administration of intravenous antimicrobials to pregnant women before delivery based on either **universal GBS screening** or **risk-based approach**

Limitations of IAP

- △ No protection against LOD, stillbirths, pre-term births, and maternal invasive GBS disease (iGBS)
- △ Risk of disruption of neonatal microbiome
- △ High antimicrobial use » acceleration of AMR
- △ Difficult to implement in LMICs

MATERNAL GBS VACCINATION³

A maternal GBS vaccination (with 80% efficacy & 50-90% coverage) can prevent **127,000 - 231,000** cases in infants and pregnant women & **60,000 - 108,000** stillbirths/neonatal deaths annually

EVIDENCE GAPS

Important data including **prevalence** of GBS colonization in mothers, **serotype distribution**, **incidence of iGBS** are lacking in LMICs and none available for Nepal

METHODS

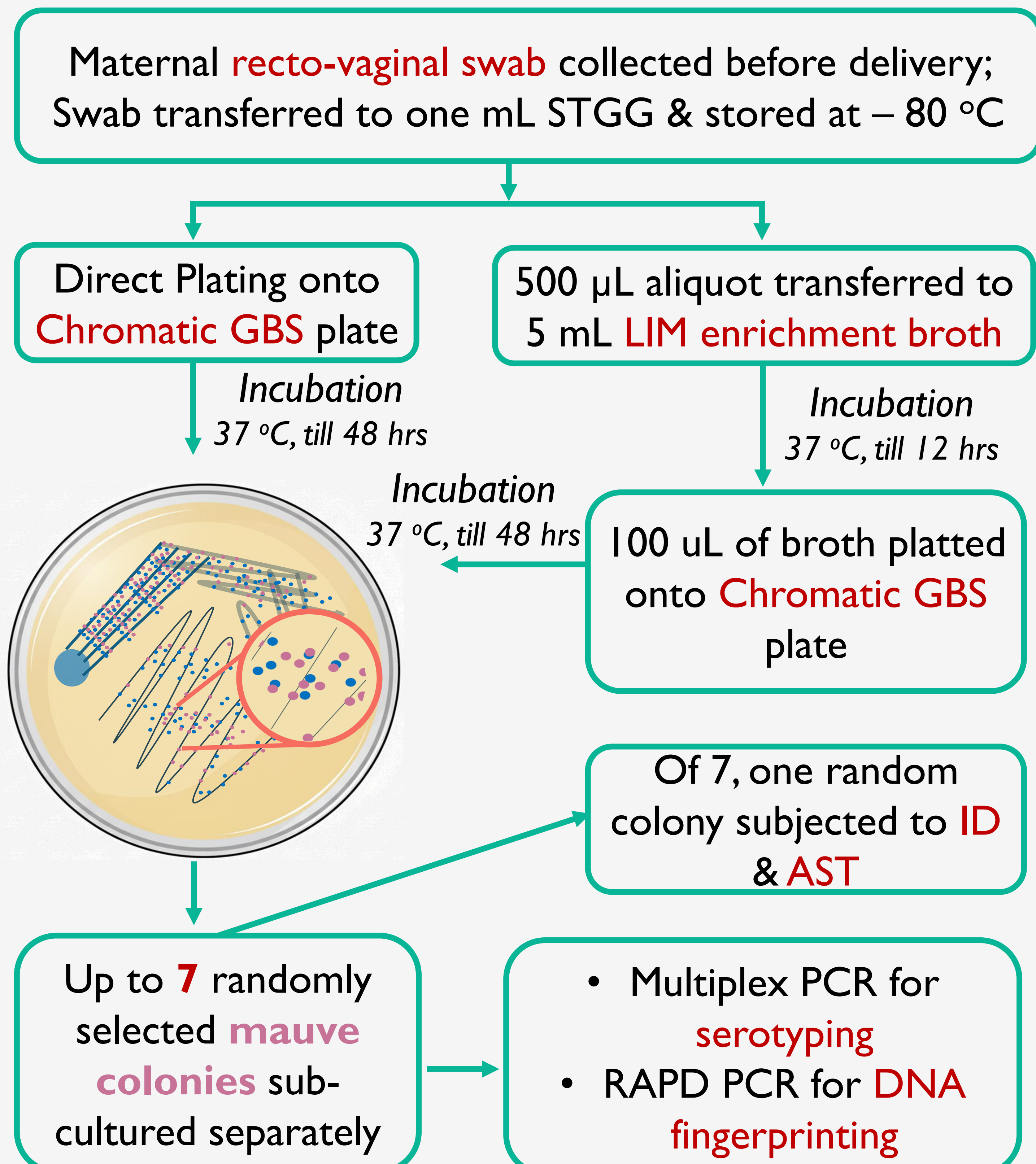
Study Design: Prospective Cohort

Study Sites: Siddhi Memorial Hospital & Bhaktapur Hospital

Study Population: 387 pregnant women & 390 neonates

Data Collection: Detail Socio-demographic & Clinical Data

Follow-up duration: One month from the day of discharge



CONCLUSION

- Low maternal GBS colonization & no iGBS in mothers or neonates; Universal GBS screening may not be feasible
- Within sample genotypic diversity could have IPC implications, warranting further investigations

RESULTS

- Prevalence of maternal GBS recto-vaginal colonization **4.4% (95% Confidence Interval: 2.8% to 6.9%)**

- **No cases** of maternal or neonatal **GBS infection**

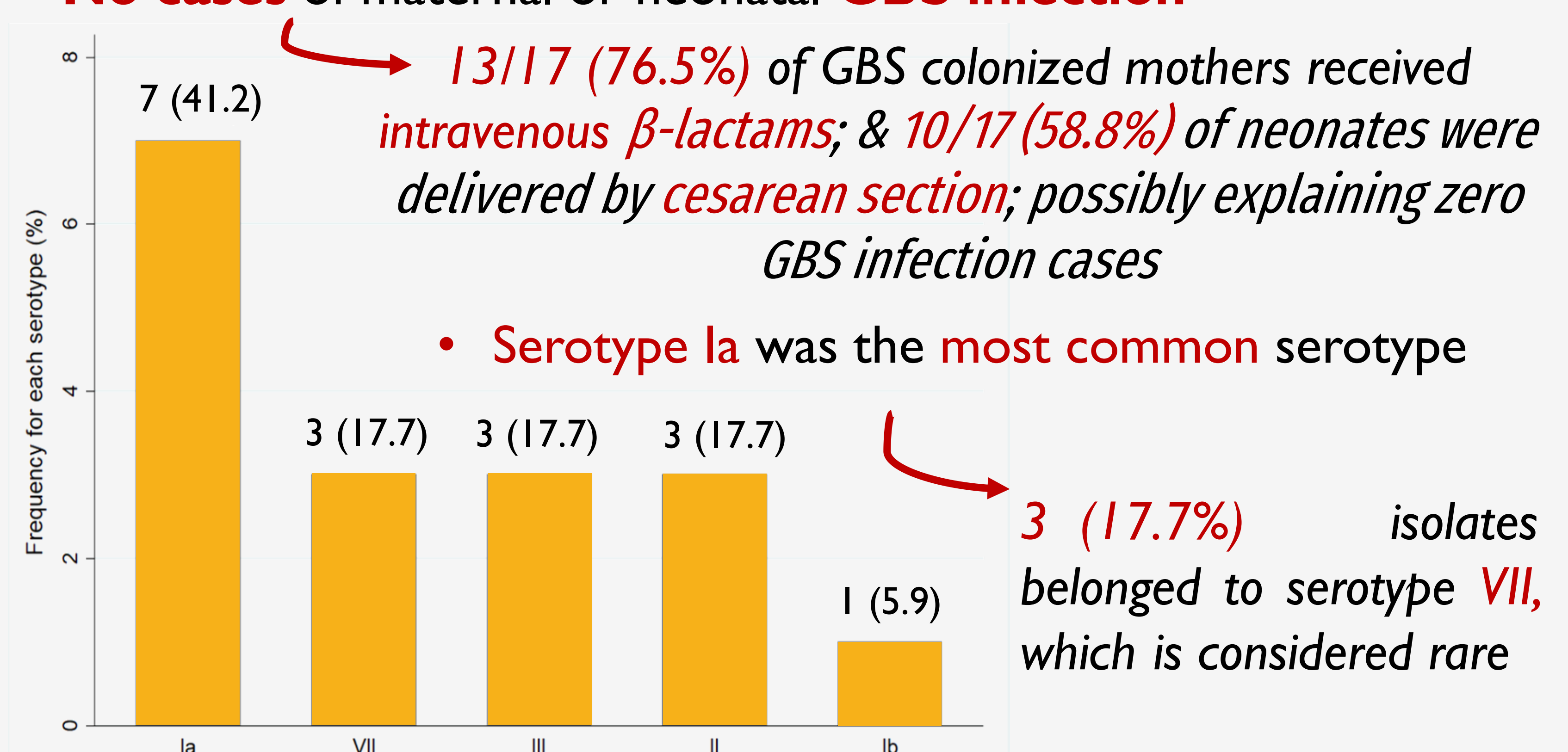


Figure 1: Distribution of serotypes of the 17 GBS isolates

- Of 17 isolates, proportion of non-susceptibility to **tetracycline**, **ofloxacin**, **erythromycin**, & **clindamycin** were **64.7%**, **23.5%**, **11.8%** and **11.8%**, respectively
- Full susceptibility to penicillin, ceftriaxone, chloramphenicol, linezolid
- Median MIC in µg/mL (IQR) of ampicillin, flomoxef, and vancomycin were 0.125 (0.125), 0.25 (0.25), & 0.38 (0.13), respectively

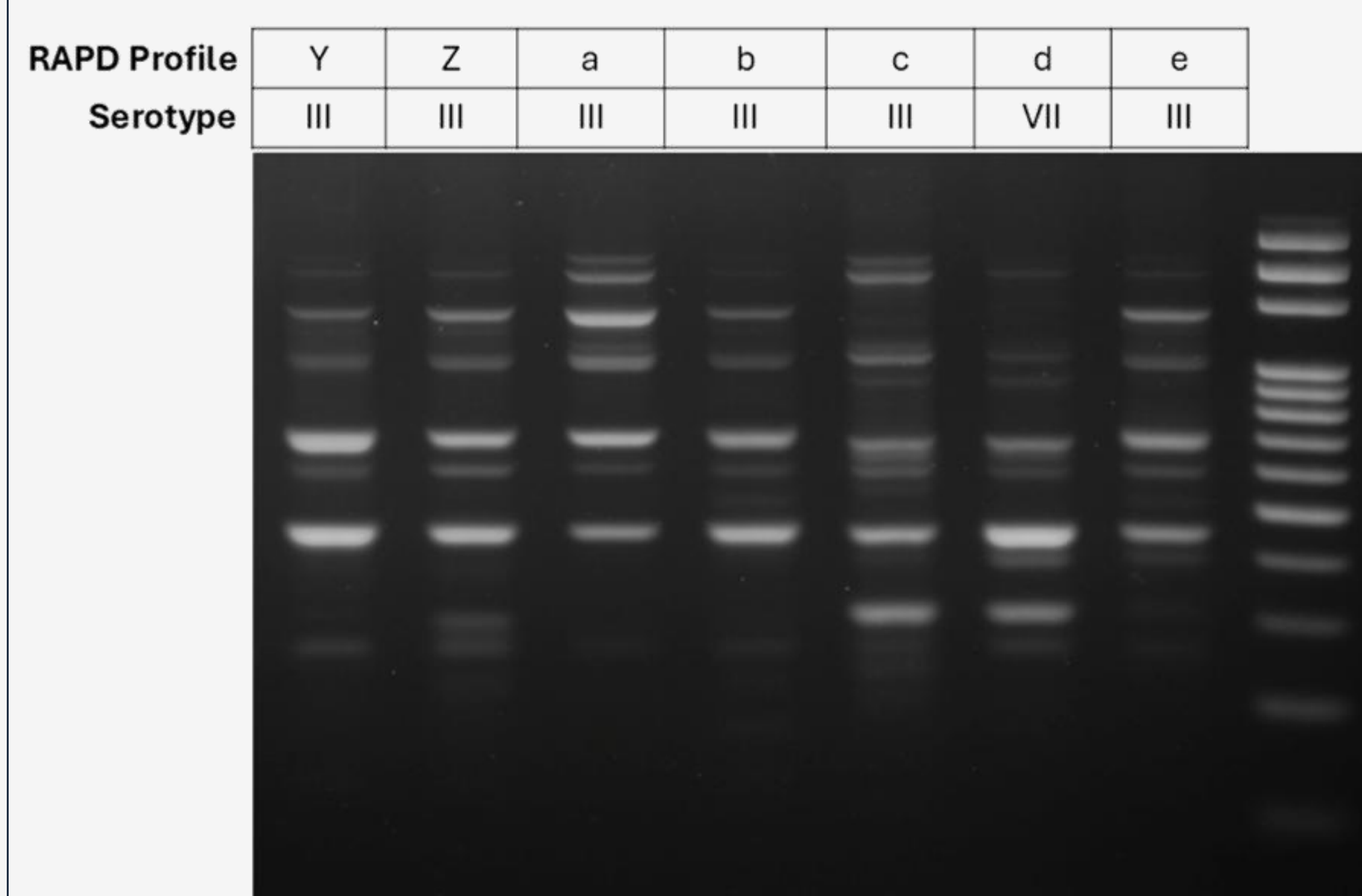


Figure 2: RAPD PCR results for one representative sample showing within-sample genetic & serotype diversity

- **Genetic diversity** within the sample = **7/17 (41.2%)**
May have implications in **outbreak investigations** in **ICU settings**
- **Serotype diversity & genetic diversity** within the sample = **1/17 (5.9%)**

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Author declare that there is no conflict of interest.



Reference:

¹Gonçalves BP et al., Lancet Glob Health., 2022

²Thorn N et al., Vaccine., 2025

³Seale AC et al., Clin Infect Dis., 2017