GLOBAL (25872)

Number of detected features / Sequencing depth (million reads)

- GSM5942339: 46.1% detected
- GSM5942340: 55% detected
- GSM5942341: 54.3% detected
- GSM5942342: 59.8% detected
- GSM5942343: 55.8% detected
- GSM5942344: 66.4% detected
- GSM5942345: 62.9% detected
- GSM5942346: 63.5% detected
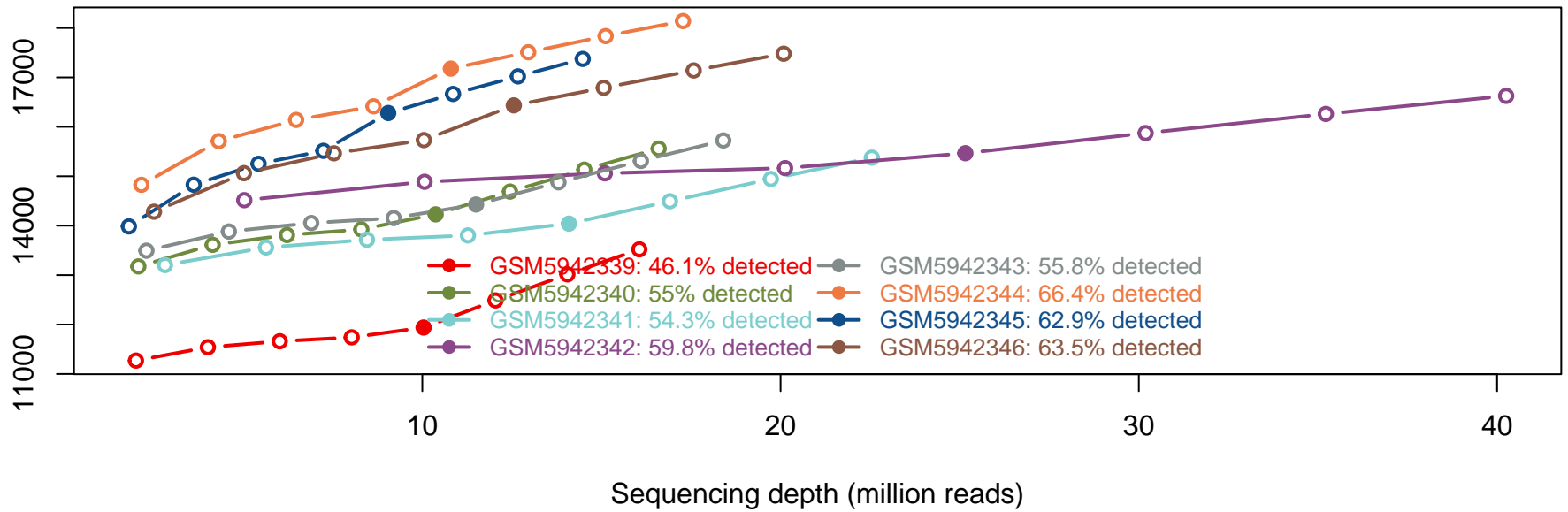
PROTEIN_CODING (19279)

Number of detected features / Sequencing depth (million reads)

- GSM5942339: 56.2% detected
- GSM5942340: 64.7% detected
- GSM5942341: 64.2% detected
- GSM5942342: 69.8% detected
- GSM5942343: 65.5% detected
- GSM5942344: 75.1% detected
- GSM5942345: 72.2% detected
- GSM5942346: 72.7% detected