Table 1: Supplementary Table S3. Data source for final infection assignments

Method used to determine infection status in usable mice (n=169)

| Assignment Method | Number of Mice | Percentage | Description |
| --- | --- | --- | --- |
| qPCR (tissue) | 49 | 29.0% | Direct infection status from MC.Eimeria |
| Species call (qPCR/amplicon)*1* | 120 | 71.0% | Infection status inferred from species identification |
| *1*Species identification prioritized qPCR melting curves, with amplicon sequencing as backup | | | |