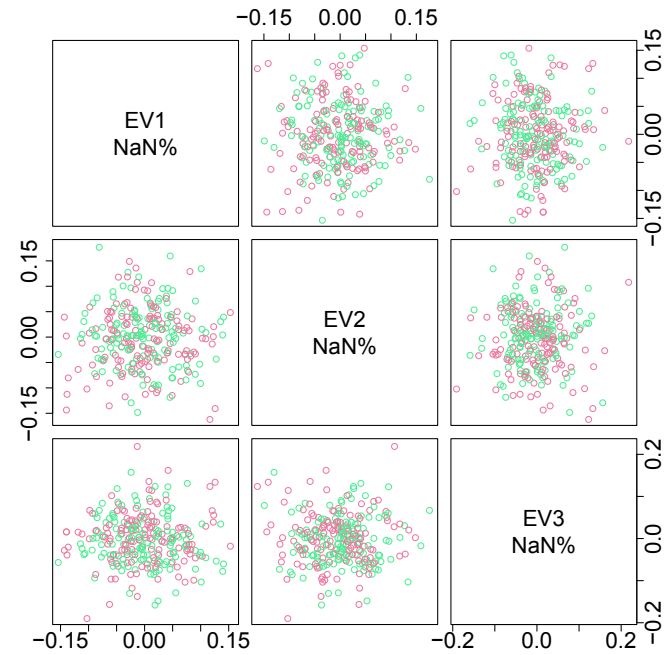
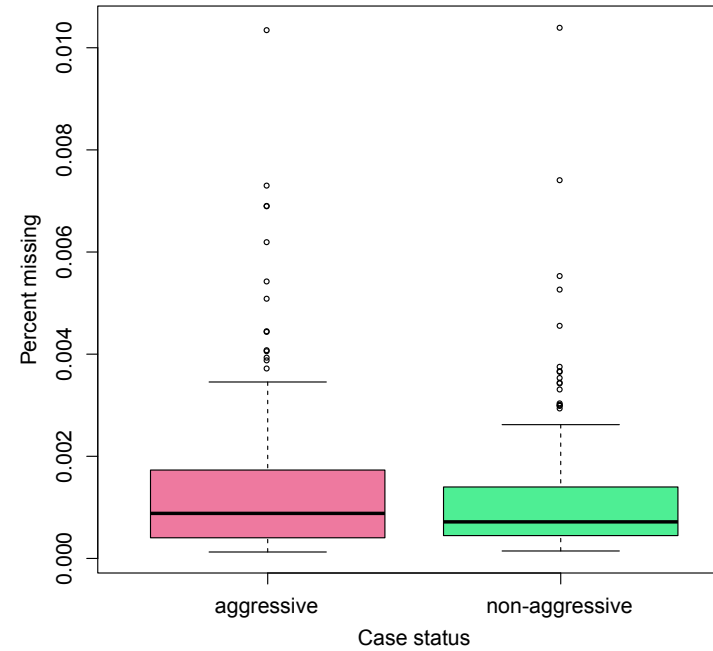
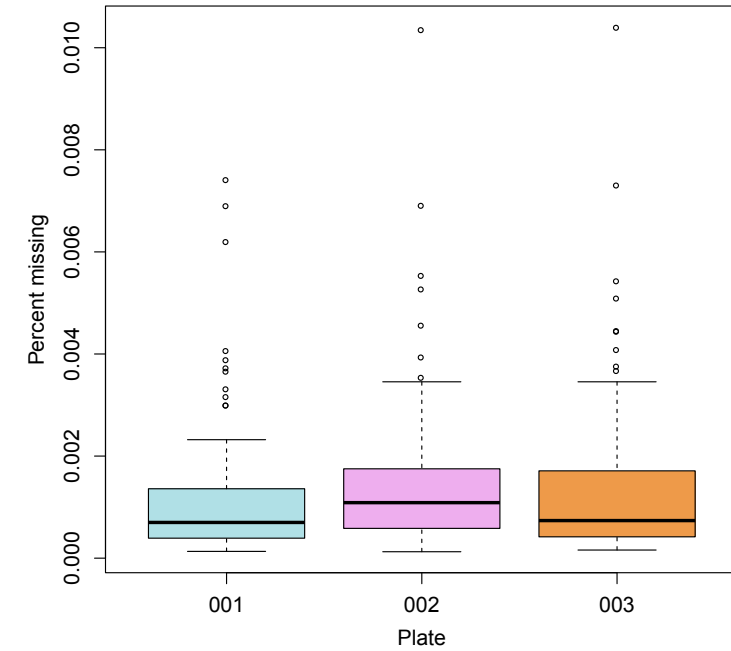


**a)****b)****c)**

**Supplementary Figure 2 – Evaluation of samples for potential batch effect confounders.** **a)** Principal Component Analysis highlighted by case status. Pairs for the first three Principal Components are shown for non-aggressive (green) and aggressive (pink) samples. **b)** Assessment of genotype call rates by case status. **c)** Assessment of genotype call rate by sequencing library preparation workflow plate. No evidence for underlying genetic or artefactual confounding effects was observed between or within phenotype groups.