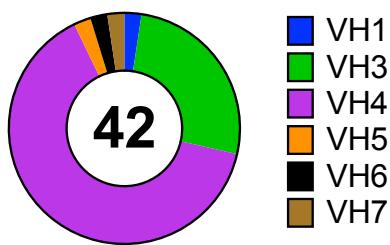
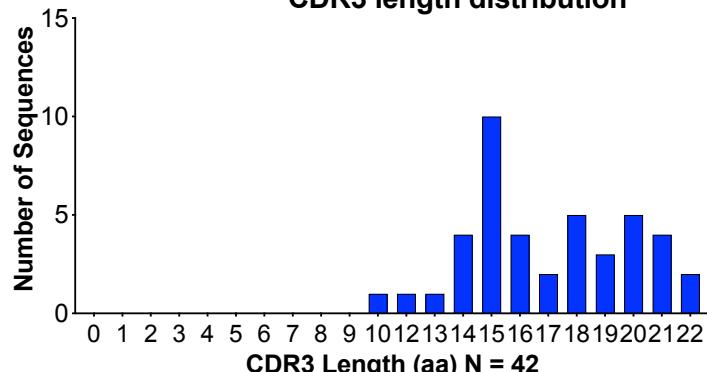


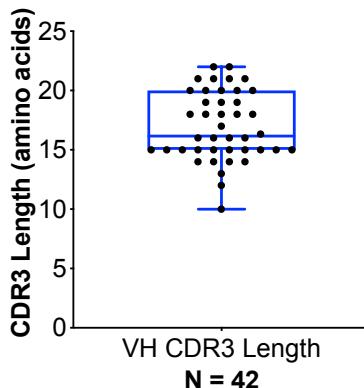
A VH Gene Usage



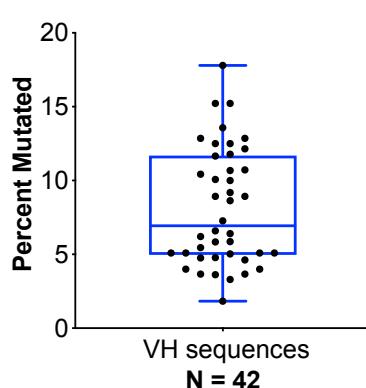
CDR3 length distribution



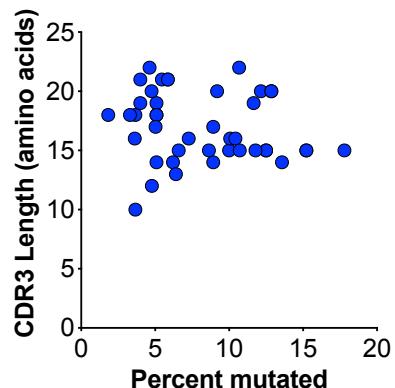
VH CDR3 Length



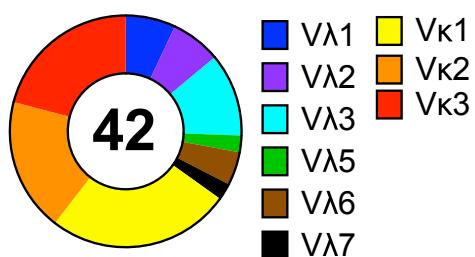
VH Mutation Percentage



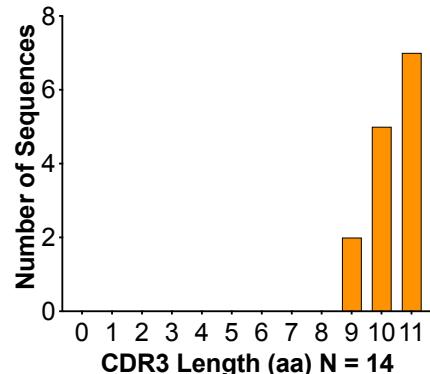
VH Percent Mutated and CDR3 Length



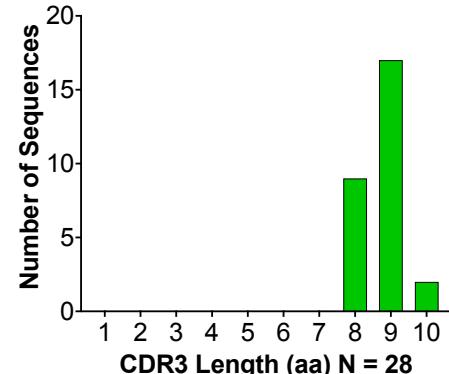
B VL Gene Usage



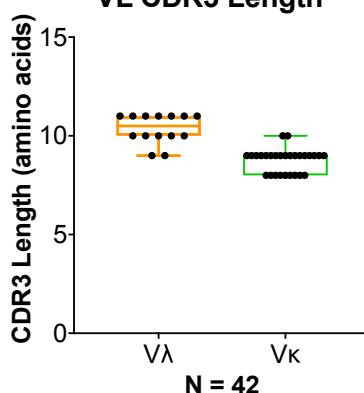
V λ CDR3 Length distribution



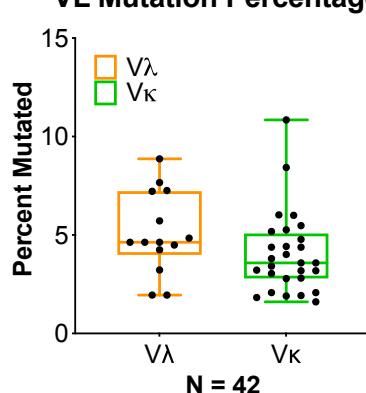
V κ CDR3 Length distribution



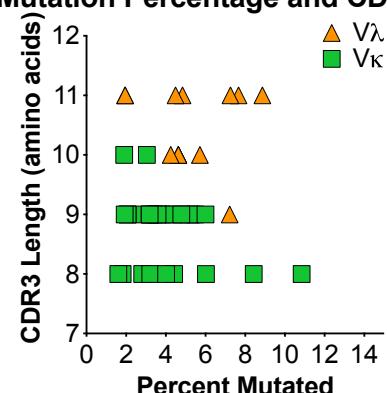
VL CDR3 Length



VL Mutation Percentage



VL Mutation Percentage and CDR3 Length



S6 Fig. Immunogenetic analysis of monoclonal antibody sequences isolated from macaque 80-12 post vaccination. DH845 and DH846 lineage antibodies were among the antibodies isolated from 80-12. Variable gene segment usage, CDR3 lengths, and mutation percentages are shown for the (A) heavy (blue), (B) lambda (orange), and (B) kappa (green) chains.