variable region (HVR-L2) of SEQ ID NO: 14, a third light chain hypervariable region (HVR-L3) of SEQ ID NO: 15; and/or a first heavy chain hypervariable region (HVR-H1) of SEQ ID NO: 41, a second heavy chain hypervariable region (HVR-H2) of SEQ ID NO: 197, and a third heavy chain hypervariable region (HVR-H3) of SEQ ID NO: 43;

[0060] (g) a first light chain hypervariable region (HVR-L1) of SEQ ID NO: 13, a second light chain hypervariable region (HVR-L2) of SEQ ID NO: 14, a third light chain hypervariable region (HVR-L3) of SEQ ID NO: 15; and/or a first heavy chain hypervariable region (HVR-H1) of SEQ ID NO: 41, a second heavy chain hypervariable region (HVR-H2) of SEQ ID NO: 42, and a third heavy chain hypervariable region (HVR-H3) of SEQ ID NO: 221;

[0061] (h) a first light chain hypervariable region (HVR-L1) of SEQ ID NO: 13, a second light chain hypervariable region (HVR-L2) of SEQ ID NO: 14, a third light chain hypervariable region (HVR-L3) of SEQ ID NO: 15; and/or a first heavy chain hypervariable region (HVR-H1) of SEQ ID NO: 41, a second heavy chain hypervariable region (HVR-H2) of SEQ ID NO: 42, and a third heavy chain hypervariable region (HVR-H3) of SEQ ID NO: 227;

[0062] (i) a first light chain hypervariable region (HVR-L1) of SEQ ID NO: 13, a second light chain hypervariable region (HVR-L2) of SEQ ID NO: 14, a third light chain hypervariable region (HVR-L3) of SEQ ID NO: 15; and/or a first heavy chain hypervariable region (HVR-H1) of SEQ ID NO: 95, a second heavy chain hypervariable region (HVR-H2) of SEQ ID NO: 112, and a third heavy chain hypervariable region (HVR-H3) of SEQ ID NO: 221;

[0063] (j) a first light chain hypervariable region (HVR-L1) of SEQ ID NO: 13, a second light chain hypervariable region (HVR-L2) of SEQ ID NO: 14, a third light chain hypervariable region (HVR-L3) of SEQ ID NO: 15; and/or a first heavy chain hypervariable region (HVR-H1) of SEQ ID NO: 95, a second heavy chain hypervariable region (HVR-H2) of SEQ ID NO: 112, and a third heavy chain hypervariable region (HVR-H3) of SEQ ID NO: 227;

[0064] (k) a first light chain hypervariable region (HVR-L1) of SEQ ID NO: 13, a second light chain hypervariable region (HVR-L2) of SEQ ID NO: 14, a third light chain hypervariable region (HVR-L3) of SEQ ID NO: 15; and/or a first heavy chain hypervariable region (HVR-H1) of SEQ ID NO: 95, a second heavy chain hypervariable region (HVR-H2) of SEQ ID NO: 113, and a third heavy chain hypervariable region (HVR-H3) of SEQ ID NO: 221;

[0065] (l) a first light chain hypervariable region (HVR-L1) of SEQ ID NO: 13, a second light chain hypervariable region (HVR-L2) of SEQ ID NO: 14, a third light chain hypervariable region (HVR-L3) of SEQ ID NO: 15; and/or a first heavy chain hypervariable region (HVR-H1) of SEQ ID NO: 95, a second heavy chain hypervariable region (HVR-H2) of SEQ ID NO: 113, and a third heavy chain hypervariable region (HVR-H3) of SEQ ID NO: 227; or

[0066] (m) a first light chain hypervariable region (HVR-L1) of SEQ ID NO: 37, a second light chain hypervariable region (HVR-L2) of SEQ ID NO: 38, a third light chain hypervariable region (HVR-L3) of SEQ ID NO: 39; and/or a first heavy chain hypervariable region (HVR-H1) of SEQ ID NO: 65, a second heavy chain hypervariable region (HVR-H2) of SEQ ID NO: 66, and a third heavy chain hypervariable region (HVR-H3) of SEQ ID NO: 67.

[0067] In some embodiments, the present disclosure provides an anti-CD96 antibody comprising:

[0068] (a) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 69;

[0069] (b) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 250;

[0070] (c) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 251;

[0071] (d) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 252;

[0072] (e) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 253;

[0073] (f) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 254;

[0074] (g) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 255;

[0075] (h) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 256;

[0076] (i) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 257;

[0077] (j) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 258;

[0078] (k) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 259;

[0079] (l) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 260;

[0080] (m) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 261;

[0081] (n) a light chain variable domain (V_L) amino acid sequence of SEQ ID NO: 68; and/or a heavy chain variable domain (V_H) amino acid sequence of SEQ ID NO: 262;