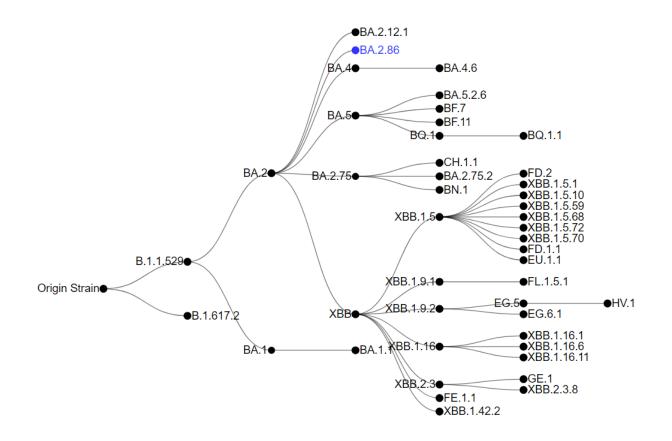
SUPPLEMENTARY FIGURE 2. SARS-CoV-2 Omicron variant lineage relationship diagram\* illustrating relative relationship of BA.2.86<sup>+</sup> with other SARS-CoV-2 Omicron variants<sup>§</sup>



\* Lines in this diagram do not reflect genomic distance but rather the relationships between lineages. BA.2.86 has been added among current circulating lineages with prevalence at or above 1% since the emergence of Omicron.

<sup>+</sup> BA.2.86 contains its sublineages including BA.2.86.1, BA.2.86.2, BA.2.86.3, JN.1, JN.2, JN.3, and JQ.1.

<sup>§</sup> BA.2.86 has not yet been included on COVID Data Tracker because it is not above 1% of currently circulating lineages. Current (September 21, 2023) circulating lineages above 1%: EG.5, FL.1.5.1, XBB.1.16, XBB.1.16.6, HV.1, XBB.2.3, XBB.1.16.1, XBB.1.5.70, XBB.1.16.11, XBB, XBB.1.5, XBB.1.9.1, GE.1, EG.6.1, and XBB.1.5.72.