

Figure S2

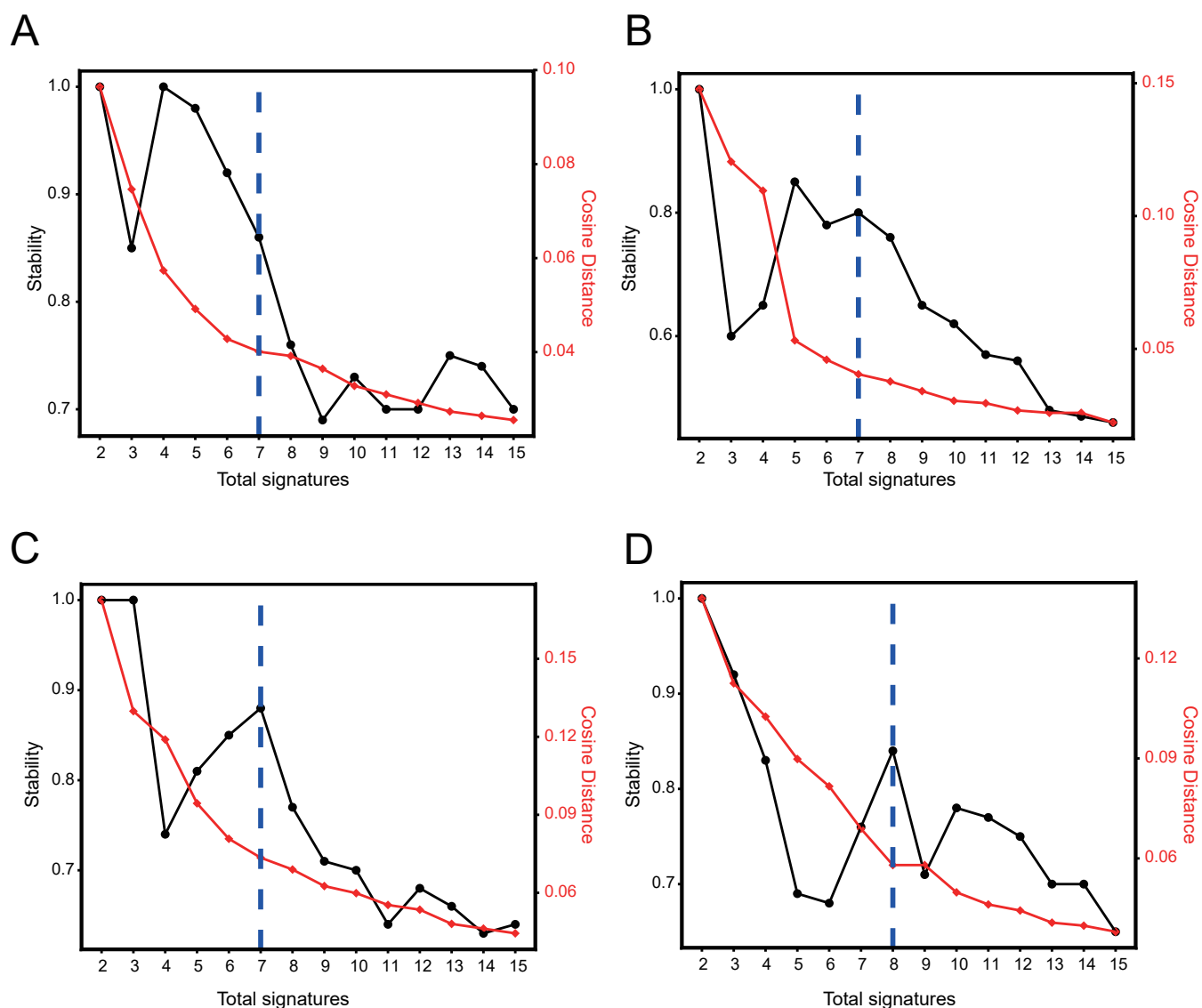


Figure S2. Determining the number of CNA signatures.

The abscissa is the number of signatures, the y-axis on the left is the stability of each signature, and the right is the similarity between signatures. Signatures have stability and low similarity. A-D are single-cell sWGS data (A), bulk sWGS (B), bulk WGS (C), and bulk SNP array (D) data.