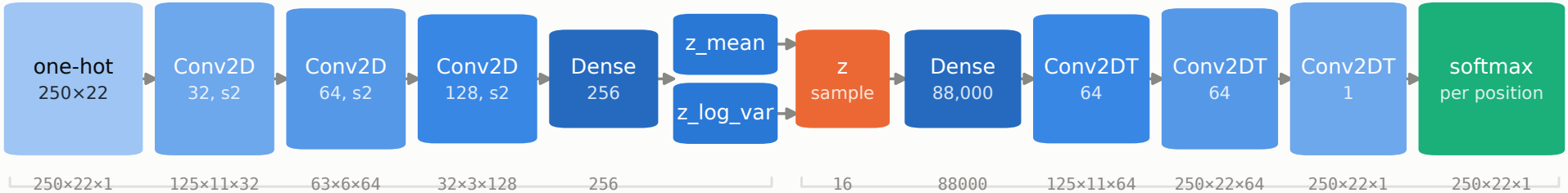
Convolutional VAE over one-hot antibody sequences

latent dimension 16 · 250 aa window · 22-letter alphabet



Strides of 2 halve each spatial axis; the decoder mirrors the encoder and ends on a softmax across the alphabet axis, so every position is a distribution over 22 residues.