Text S1. The mathematics formulas for ESM-MSA transformer

1. Masking

For an input multiple sequence alignment (MSA), the masking strategy is performed. Specifically, for each individual sequence in MSA, we randomly sample 15% tokens (amino acids), each of which is changed as a special "masking" token with 80% probability, a randomly-chosen alternate amino acid with 10% probability, and the original input token (i.e., no change) with 10% probability.

2. One-hot encoding

The masked MSA is encoded as three matrices using one-hot encoding from three different views. Specifically, for the *j*-th position of the *i*-th sequence in the masked MSA, we encode it as three one-hot vectors, i.e., x_{ij} , y_{ij} , and z_{ij} , from the views of token type, row position, and column position, respectively.

$$\mathbf{x}_{ij} = (x_{ij1}, x_{ij2}, \dots, x_{ijC_{max}}) \in R^{C_{max}}, x_{ijk} = \begin{cases} 1, \ k = c_{ij} \\ 0, \ k \neq c_{ij} \end{cases}$$
(1)

$$\mathbf{y}_{ij} = (y_{ij1}, y_{ij2}, \dots, y_{ijM_{max}}) \in R^{M_{max}}, y_{ijk} = \begin{cases} 1, \ k = i \\ 0, \ k \neq i \end{cases}$$
(2)

$$\mathbf{z}_{ij} = (z_{ij1}, z_{ij2}, \dots, z_{ijL_{max}}) \in R^{L_{max}}, z_{ijk} = \begin{cases} 1, \ k = j \\ 0, \ k \neq j \end{cases}$$
(3)

where c_{ij} is the index of token type for the *j*-th position of the *i*-th sequence, C_{max} is the number of types of tokens, L_{max} and M_{max} are preset maximum values for sequence length and alignments, respectively. In this work, $C_{max} = 28$ and $L_{max} = M_{max} = 1024$, where 28 types of tokens include 20 common amino acids, 6 non-common amino acids (B, J, O, U, X and Z), 1 gap token, and 1 "masking" token.

According to Eqs. 1-3, the masked MSA can be encoded as three matrices, i.e., X, Y and Z, through one-hot encoding from the view of token type, row position, and column position, respectively, where $X \in R^{M \times L \times C_{max}}$, $Y \in R^{M \times L \times M_{max}}$ and $Z \in R^{M \times L \times L_{max}}$, M is the number of alignments, and L is the length of individual sequence in the masked MSA.

3. Initial embedding

Each one-hot coding matrix is multiplied by a weight matrix to generate the corresponding embedding matrix:

$$H_{token} = XW_{token} = \begin{bmatrix} X[1] \\ X[2] \\ ... \\ X[M] \end{bmatrix} W_{token} = \begin{bmatrix} X[1]W_{token} \\ X[2]W_{token} \\ ... \\ X[M]W_{token} \end{bmatrix} \in R^{M \times L \times D}$$
(4)
$$X[i] \in R^{L \times C_{max}}, W_{token} \in R^{C_{max} \times D}$$
$$H_{row} = XW_{row} = \begin{bmatrix} Y[1] \\ Y[2] \\ ... \\ Y[M] \end{bmatrix} W_{row} = \begin{bmatrix} Y[1]W_{row} \\ Y[2]W_{row} \\ ... \\ Y[M]W_{row} \end{bmatrix} \in R^{M \times L \times D}$$
(5)
$$Y[i] \in R^{L \times M_{max}}, W_{row} \in R^{M_{max} \times D}$$
$$H_{col} = ZW_{col} = \begin{bmatrix} Z[1] \\ Z[2] \\ ... \\ Z[M] \end{bmatrix} W_{col} = \begin{bmatrix} Z[1]W_{col} \\ Z[M]W_{col} \end{bmatrix} \in R^{M \times L \times D}$$
(6)
$$Z[i] \in R^{L \times L_{max}}, W_{col} \in R^{L_{max} \times D}$$

where X[i], Y[i] and Z[i] are the one-hot coding matrices for the *i*-th sequence in the masked MSA from the view of token type, row position, and column position, respectively, H_{token} , H_{row} , and H_{col} are token type-based, row position-based, and column position-based embedding matrices for the masked MSA, respectively, and *D* is the embedding dimension. In this work, D = 768.

Three embedding matrices are added as an initial embedding matrix H_{init} :

$$\boldsymbol{H}_{init} = \boldsymbol{H}_{token} + \boldsymbol{H}_{row} + \boldsymbol{H}_{col}, \boldsymbol{H}_{init} \in R^{M \times L \times D}$$
(7)

4. Batch normalization and dropout

The initial embedding matrix H_{init} is fed to the batch normalization layer to generate the corresponding normalized matrix H_1 :

$$\boldsymbol{H}_{1} = BN(\boldsymbol{H}_{init}) = \begin{bmatrix} BN(\boldsymbol{h}_{11}) & \cdots & BN(\boldsymbol{h}_{1L}) \\ \vdots & \ddots & \vdots \\ BN(\boldsymbol{h}_{M1}) & \cdots & BN(\boldsymbol{h}_{ML}) \end{bmatrix}$$
(8)

$$BN(\boldsymbol{h}_{ij}) = \gamma \cdot \frac{\boldsymbol{h}_{ij} - \boldsymbol{u}_{ij}}{\sqrt{\sigma_{ij}^2 + \epsilon}} + \beta, \, \boldsymbol{h}_{ij} \in R^D$$
(9)

where h_{ij} is the initial embedding vector for the *j*-th position of the *i*-th sequence in the masked MSA, u_{ij} and σ_{ij}^2 are mean and variance for h_{ij} , respectively, and γ , β , and ϵ are normalized factors.

The normalized matrix H_1 is fed to dropout layer:

$$\boldsymbol{H}_{1} \leftarrow dropout(\boldsymbol{H}_{1}, r) \tag{10}$$

where r is the rate of neurons which are randomly dropped in each training step, indicating that the corresponding weight vectors will be not optimized.

5. Self-attention

The initial embedding matrix H_1 is fed to the self-attention network with N blocks, each of which consists of three sub-blocks. In this work, N = 12.

The first sub-block consists of a batch normalization layer, a row attention layer, a dropout layer, and a short connection, as follows.

$$\boldsymbol{H}_{k}^{B} = BN(\boldsymbol{H}_{k}) \tag{11}$$

$$\boldsymbol{H}_{k}^{R} = RA(\boldsymbol{H}_{k}^{B}) \tag{12}$$

$$\boldsymbol{H}_{k}^{R} \leftarrow dropout(\boldsymbol{H}_{k}^{R}, r) \tag{13}$$

$$\boldsymbol{F}_{k} = SC(\boldsymbol{H}_{k}, \boldsymbol{H}_{k}^{R}) = \boldsymbol{H}_{k} + \boldsymbol{H}_{k}^{R}$$
(14)

where H_k and F_k are the input and output matrices in the first sub-block of the *k*-th self-attention block, respectively, $BN(\cdot)$ is the batch normalization function (see Eqs. 8-9), $SC(\cdot)$ is the short connection, and $RA(\cdot)$ is the row attention layer (see Eqs. 23-30), H_k , H_k^B , H_k^R , $F_k \in R^{M \times L \times D}$.

The second sub-block consists of a batch normalization layer, a column attention layer, a dropout layer, and a short connection, as follows.

$$\boldsymbol{F}_{k}^{B} = BN(\boldsymbol{F}_{k}) \tag{15}$$

$$\boldsymbol{F}_{k}^{C} = CA(\boldsymbol{F}_{k}^{B}) \tag{16}$$

$$\boldsymbol{F}_{k}^{C} \leftarrow dropout(\boldsymbol{F}_{k}^{C}, r) \tag{17}$$

$$\boldsymbol{U}_{k} = SC(\boldsymbol{F}_{k}, \boldsymbol{F}_{k}^{C}) = \boldsymbol{F}_{k} + \boldsymbol{F}_{k}^{C}$$
(18)

where F_k and U_k are the input and output matrices in the second sub-block of the *k*-th self-attention block, respectively, $CA(\cdot)$ is the column attention layer (see Eqs. 31-39), and F_k^B , F_k^C , $U_k \in \mathbb{R}^{M \times L \times D}$.

The last sub-block consists of a batch normalization layer, a feed-forward network, a dropout layer, and a short connection, as follows.

$$\boldsymbol{U}_{k}^{B} = BN(\boldsymbol{U}_{k}) \tag{19}$$

$$\boldsymbol{U}_{k}^{F} = FFN(\boldsymbol{U}_{k}^{B}) \tag{20}$$

$$\boldsymbol{U}_{k}^{F} \leftarrow dropout(\boldsymbol{U}_{k}^{F}, r)$$
(21)

$$\boldsymbol{H}_{k+1} = SC(\boldsymbol{U}_k, \boldsymbol{U}_k^F) = \boldsymbol{U}_k + \boldsymbol{U}_k^F$$
(22)

where U_k and H_{k+1} are the input and output matrices in the third sub-block of the k-th self-attention block, respectively, FFN(.) is the feed-forward network (see Eqs. 40-45), and U_k^B , U_k^F , $H_{k+1} \in \mathbb{R}^{M \times L \times D}$.

(A) Row attention

Each row attention layer consists of m attention heads and a linear unit, where m = 12. In each attention head, the input matrix is multiplied by three weight matrices to generate the corresponding Query, Key, and Value matrices.

$$\boldsymbol{Q}_{kt}^{R} = \boldsymbol{H}_{k}^{B} \boldsymbol{W}_{kt}^{QR} = \begin{bmatrix} \boldsymbol{H}_{k}^{B}[1] \\ \boldsymbol{H}_{k}^{B}[2] \\ \dots \\ \boldsymbol{H}_{k}^{B}[M] \end{bmatrix} \boldsymbol{W}_{kt}^{QR} = \begin{bmatrix} \boldsymbol{H}_{k}^{B}[1] \boldsymbol{W}_{kt}^{QR} \\ \boldsymbol{H}_{k}^{B}[2] \boldsymbol{W}_{kt}^{QR} \\ \dots \\ \boldsymbol{H}_{k}^{B}[M] \boldsymbol{W}_{kt}^{QR} \end{bmatrix} \in R^{M \times L \times (\frac{D}{m})}$$
(23)

$$K_{kt}^{R} = H_{k}^{B} W_{kt}^{KR} = \begin{bmatrix} H_{k}^{B}[1] \\ H_{k}^{B}[2] \\ \dots \\ H_{k}^{B}[M] \end{bmatrix} W_{kt}^{KR} = \begin{bmatrix} H_{k}^{B}[1] W_{kt}^{KR} \\ H_{k}^{B}[2] W_{kt}^{KR} \\ \dots \\ H_{k}^{B}[M] W_{kt}^{KR} \end{bmatrix} \in R^{M \times L \times (\frac{D}{m})}$$
(24)
$$V_{kt}^{R} = H_{k}^{B} W_{kt}^{VR} = \begin{bmatrix} H_{k}^{B}[1] \\ H_{k}^{B}[2] \\ \dots \\ H_{k}^{B}[M] \end{bmatrix} W_{kt}^{VR} = \begin{bmatrix} H_{k}^{B}[1] W_{kt}^{VR} \\ H_{k}^{B}[2] W_{kt}^{VR} \\ \dots \\ H_{k}^{B}[M] W_{kt}^{VR} \end{bmatrix} \in R^{M \times L \times (\frac{D}{m})}$$
(25)
$$H_{k}^{B}[i] \in R^{L \times D}, W_{kt}^{QR}, W_{kt}^{KR}, W_{kt}^{KR} \in R^{D \times (\frac{D}{m})}$$

where \boldsymbol{H}_{k}^{B} is the input matrix of row attention layer in the *k*-th self-attention block (See Eq. 12), \boldsymbol{Q}_{kt}^{R} , \boldsymbol{K}_{kt}^{R} , and \boldsymbol{V}_{kt}^{R} are Query, Key, and Value matrices in the *t*-th head of the row attention layer in the *k*-th block, respectively, \boldsymbol{W}_{kt}^{QR} , \boldsymbol{W}_{kt}^{KR} , and \boldsymbol{W}_{kt}^{VR} are corresponding weight metrices.

Then, the dot-product between Q_{kt}^R and K_{kt}^R is performed and then normalized by SoftMax function to generate a row attention weight matrix:

$$\boldsymbol{W}_{kt}^{AR} = SoftMax(\frac{\sum_{i=1}^{M} \boldsymbol{Q}_{kt}^{R}[i] \cdot (\boldsymbol{K}_{kt}^{R}[i])^{T}}{\sqrt{MD/m}}) \in R^{L \times L}, \ \boldsymbol{Q}_{kt}^{R}[i], \ \boldsymbol{K}_{kt}^{R}[i] \in R^{L \times (D/m)}$$
(26)
$$\boldsymbol{W}_{kt}^{AR} \leftarrow dropout(\boldsymbol{W}_{kt}^{AR}, r)$$
(27)

where W_{kt}^{AR} is the attention weight matrix in the *t*-th head of the row attention layer in the *k*-th block and measures the correlation for each pair of columns in the masked MSA.

Next, the row attention weight matrix W_{kt}^{AR} is multiplied by Value matrix V_{kt}^{R} to generate the corresponding row attention matrix:

$$\boldsymbol{A}_{kt}^{R} = \boldsymbol{W}_{kt}^{AR} \boldsymbol{V}_{kt}^{R} = \boldsymbol{W}_{kt}^{AR} \begin{bmatrix} \boldsymbol{V}_{kt}^{R}[1] \\ \boldsymbol{V}_{kt}^{R}[2] \\ \dots \\ \boldsymbol{V}_{kt}^{R}[M] \end{bmatrix} = \begin{bmatrix} \boldsymbol{W}_{kt}^{AR} \boldsymbol{V}_{kt}^{R}[1] \\ \boldsymbol{W}_{kt}^{AR} \boldsymbol{V}_{kt}^{R}[2] \\ \dots \\ \boldsymbol{W}_{kt}^{AR} \boldsymbol{V}_{kt}^{R}[M] \end{bmatrix} \in R^{M \times L \times \left(\frac{D}{m}\right)}, \boldsymbol{V}_{kt}^{R}[i] \in R^{L \times \left(\frac{D}{m}\right)}$$
(28)

where A_{kt}^{R} is the attention matrix in the *t*-th head of the row attention layer in the *k*-th block.

Finally, the outputs of all attention heads are concatenated as a new matrix, which is further fed to a linear unit:

$$\boldsymbol{A}_{k}^{R} = \boldsymbol{A}_{k1}^{R} \boldsymbol{A}_{k2}^{R} \dots \boldsymbol{A}_{km}^{R} \in \boldsymbol{R}^{M \times L \times D}$$

$$\boldsymbol{\boldsymbol{A}}_{k}^{R} \boldsymbol{\boldsymbol{A}}_{k1}^{R} \boldsymbol{\boldsymbol{A}}_{k2}^{R} \dots \boldsymbol{\boldsymbol{A}}_{km}^{R} \boldsymbol{\boldsymbol{A}$$

$$\boldsymbol{H}_{k}^{R} = \boldsymbol{A}_{k}^{R}\boldsymbol{W}_{k}^{R} + \boldsymbol{b}_{k}^{R} = \begin{bmatrix} \boldsymbol{A}_{k}^{[1]} \\ \boldsymbol{A}_{k}^{R}[2] \\ \dots \\ \boldsymbol{A}_{k}^{R}[M] \end{bmatrix} \boldsymbol{W}_{k}^{R} + \boldsymbol{b}_{k}^{R} = \begin{bmatrix} \boldsymbol{A}_{k}^{[1]} \boldsymbol{W}_{k} \\ \boldsymbol{A}_{k}^{R}[2] \boldsymbol{W}_{k}^{R} \\ \dots \\ \boldsymbol{A}_{k}^{R}[M] \boldsymbol{W}_{k}^{R} \end{bmatrix} + \boldsymbol{b}_{k}^{R} \in R^{M \times L \times D} \quad (30)$$
$$\boldsymbol{W}_{k}^{R} \in R^{D \times D}, \boldsymbol{A}_{k}^{R}[i] \in R^{L \times D}$$

where \boldsymbol{H}_{k}^{R} in the output matrix of row attention layer in the *k*-th attention block (See Eq. 12), and \boldsymbol{W}_{k}^{R} and \boldsymbol{b}_{k}^{R} are weight matrix and bias in the linear unit, respectively.

(B) Column attention

Each column attention layer consists of m attention heads and a linear unit. In each attention head, the input matrix is multiplied by three weight matrices to generate the corresponding Query, Key, and Value matrices.

$$\mathbf{Q}_{kt}^{C} = \mathbf{F}_{k}^{B} \mathbf{W}_{kt}^{QC} = \begin{bmatrix} \mathbf{F}_{k}^{B} [1] \\ \mathbf{F}_{k}^{B} [2] \\ \dots \\ \mathbf{F}_{k}^{B} [M] \end{bmatrix} \mathbf{W}_{kt}^{QC} = \begin{bmatrix} \mathbf{F}_{k}^{B} [1] \mathbf{W}_{kt}^{QC} \\ \mathbf{F}_{k}^{B} [2] \mathbf{W}_{kt}^{QC} \\ \dots \\ \mathbf{F}_{k}^{B} [M] \mathbf{W}_{kt}^{QC} \end{bmatrix} \in R^{M \times L \times (\frac{D}{m})}$$

$$\mathbf{K}_{kt}^{C} = \mathbf{F}_{k}^{B} \mathbf{W}_{kt}^{KC} = \begin{bmatrix} \mathbf{F}_{k}^{B} [1] \\ \mathbf{F}_{k}^{B} [2] \\ \dots \\ \mathbf{F}_{k}^{B} [M] \end{bmatrix} \mathbf{W}_{kt}^{KC} = \begin{bmatrix} \mathbf{F}_{k}^{B} [1] \mathbf{W}_{kt}^{KC} \\ \mathbf{F}_{k}^{B} [2] \mathbf{W}_{kt}^{KC} \\ \dots \\ \mathbf{F}_{k}^{B} [M] \mathbf{W}_{kt}^{KC} \end{bmatrix} \in R^{M \times L \times (\frac{D}{m})}$$

$$\mathbf{V}_{kt}^{C} = \mathbf{F}_{k}^{B} \mathbf{W}_{kt}^{VC} = \begin{bmatrix} \mathbf{F}_{k}^{B} [1] \\ \mathbf{F}_{k}^{B} [2] \\ \dots \\ \mathbf{F}_{k}^{B} [M] \end{bmatrix} \mathbf{W}_{kt}^{VC} = \begin{bmatrix} \mathbf{F}_{k}^{B} [1] \mathbf{W}_{kt}^{VC} \\ \mathbf{F}_{k}^{B} [2] \mathbf{W}_{kt}^{VC} \\ \mathbf{F}_{k}^{B} [2] \mathbf{W}_{kt}^{VC} \\ \mathbf{F}_{k}^{B} [2] \mathbf{W}_{kt}^{VC} \end{bmatrix} \in R^{M \times L \times (\frac{D}{m})}$$

$$(32)$$

$$\boldsymbol{F}_{k}^{B}[i] \in R^{L \times D}, \boldsymbol{W}_{kt}^{QC}, \boldsymbol{W}_{kt}^{KC}, \boldsymbol{W}_{kt}^{VC} \in R^{D \times (\frac{D}{m})}$$

where F_k^B is the input matrix of column attention layer in the *k*-th self-attention block (see Eq. 16), Q_{kt}^C , K_{kt}^C , and V_{kt}^C are Query, Key, and Value matrices in the *t*-th head of column attention layer in the *k*-th block, respectively, W_{kt}^{QC} , W_{kt}^{KC} , and W_{kt}^{VC} are corresponding weight metrices.

th block.

Then, the dot-product between Q_{kt}^{C} and K_{kt}^{C} is performed and then normalized by SoftMax function to generate an attention weight matrix:

$$\boldsymbol{W}_{kt}^{AC} = SoftMax\left(\frac{\boldsymbol{Q}_{kt}^{C}(\boldsymbol{K}_{kt}^{C})^{T}}{\sqrt{D/m}}\right) \in R^{M \times L \times M}$$
(34)

$$\boldsymbol{W}_{kt}^{AC} \leftarrow dropout(\boldsymbol{W}_{kt}^{AC}, r)$$
(35)

$$\boldsymbol{Q}_{kt}^{C}(\boldsymbol{K}_{kt}^{C})^{T} = \left[\boldsymbol{Q}_{kt}^{C}[:,1,:] \; \boldsymbol{Q}_{kt}^{C}[:,2,:] \; \dots \; \boldsymbol{Q}_{kt}^{C}[:,L,:]\right] \cdot \left[\boldsymbol{K}_{kt}^{C}[:,1,:] \; \boldsymbol{K}_{kt}^{C}[:,2,:] \; \dots \; \boldsymbol{K}_{kt}^{C}[:,L,:]\right]^{T} =$$

 $\left[\boldsymbol{Q}_{kt}^{C}[:,1,:] \cdot \boldsymbol{K}_{kt}^{C}[:,1,:]^{T} \, \boldsymbol{Q}_{kt}^{C}[:,2,:] \cdot \boldsymbol{K}_{kt}^{C}[:,2,:]^{T} \dots \boldsymbol{Q}_{kt}^{C}[:,L,:] \cdot \boldsymbol{K}_{kt}^{C}[:,L,:]^{T}\right] \in \boldsymbol{R}^{M \times L \times M}$ (36)

$$\boldsymbol{Q}_{kt}^{C}[:,j,:], \boldsymbol{K}_{kt}^{C}[:,j,:] \in R^{M \times \left(\frac{D}{m}\right)}, \boldsymbol{Q}_{kt}^{C}[:,j,:] \cdot \boldsymbol{K}_{kt}^{C}[:,j,:]^{T} \in R^{M \times M}$$

where W_{kt}^{AC} is the attention weight matrix in the *t*-th head of column attention layer in the *k*-th block, and $W_{kt}^{AC}[:, j, :]$ measures the correlation for each pair of alignments at the *j*-th position.

Next, the column attention weight matrix W_{kt}^{AC} is multiplied by Value matrix V_{kt}^{C} to generate the corresponding column attention matrix:

$$A_{kt}^{C} = W_{kt}^{AC} V_{kt}^{C} = \left[W_{kt}^{AC} [:, 1, :] W_{kt}^{AC} [:, 2, :] \dots W_{kt}^{AC} [:, L, :] \right] \cdot \left[V_{kt}^{C} [:, 1, :] V_{kt}^{C} [:, 2, :] \dots V_{kt}^{C} [:, L, :] \right] = \left[W_{kt}^{AC} [:, 1, :] V_{kt}^{C} [:, 2, :] \dots V_{kt}^{C} [:, 2, :] \dots V_{kt}^{C} [:, 2, :] \right]$$

$$V_{kt}^{C} [:, 1, :] W_{kt}^{AC} [:, 2, :] \cdot V_{kt}^{C} [:, 2, :] \dots W_{kt}^{AC} [:, L, :] \cdot V_{kt}^{C} [:, L, :] \right] \in \mathbb{R}^{M \times L \times (\frac{D}{m})}$$
(37)

$$W_{kt}^{AC}[:,j,:] \in \mathbb{R}^{M \times M}, V_{kt}^{C}[:,j,:] \in \mathbb{R}^{M \times (\frac{D}{m})}, W_{kt}^{AC}[:,j,:] \cdot V_{kt}^{C}[:,j,:] \in \mathbb{R}^{M \times (\frac{D}{m})}$$

where A_{kt}^{C} is the attention matrix in the *t*-th head of column attention layer in the *k*-

Finally, the outputs of all attention heads are concatenated as a new matrix, which is further fed to a linear unit:

$$\boldsymbol{A}_{k}^{C} = \boldsymbol{A}_{k1}^{C} \boldsymbol{A}_{k2}^{C} \dots \boldsymbol{A}_{km}^{C} \in \boldsymbol{R}^{M \times L \times D}$$

$$\boldsymbol{\boldsymbol{\Gamma}} \boldsymbol{A}^{C} \boldsymbol{\boldsymbol{\Gamma}} \boldsymbol{\boldsymbol{1}} \boldsymbol{\boldsymbol{1}} \boldsymbol{\boldsymbol{\Gamma}} \qquad \boldsymbol{\boldsymbol{\Gamma}} \boldsymbol{A}^{C} \boldsymbol{\boldsymbol{\Gamma}} \boldsymbol{\boldsymbol{1}} \boldsymbol{\boldsymbol{1}} \boldsymbol{\boldsymbol{W}}^{C} \boldsymbol{\boldsymbol{1}} \boldsymbol{\boldsymbol{1}}$$

$$(38)$$

$$F_{k}^{C} = A_{k}^{C} W_{k}^{C} + b_{k}^{C} = \begin{bmatrix} A_{k}^{[1]} \\ A_{k}^{C}[2] \\ ... \\ A_{k}^{C}[M] \end{bmatrix} W_{k}^{C} = \begin{bmatrix} A_{1}^{[1]} W_{k} \\ A_{2}^{C}[2] W_{k}^{C} \\ ... \\ A_{k}^{C}[M] W_{k}^{C} \end{bmatrix} + b_{k}^{C} \in \mathbb{R}^{M \times L \times D}$$
(39)
$$W_{k}^{C} \in \mathbb{R}^{D \times D}, A_{k}^{C}[i] \in \mathbb{R}^{L \times D}$$

where F_k^c in the output matrix of column attention layer in the k-th attention block, (See Eq. 16), and W_k^c and b_k^c are weight matrix and bias in the linear unit, respectively.

(C) Feed-forward network

$$\boldsymbol{T}_{k}^{F} = gelu(\boldsymbol{U}_{k}^{B}\boldsymbol{W}_{k}^{1} + \boldsymbol{b}_{k}^{1}) \in R^{M \times L \times D_{1}}$$

$$\tag{40}$$

$$\boldsymbol{T}_{k}^{F} \leftarrow dropout(\boldsymbol{T}_{k}^{F}, r) \tag{41}$$

$$\boldsymbol{U}_{k}^{F} = \boldsymbol{T}_{k}^{F} \boldsymbol{W}_{k}^{2} + \boldsymbol{b}_{k}^{2} \in R^{M \times L \times D}$$

$$\tag{42}$$

$$gelu(x) = x \emptyset(x) \tag{43}$$

$$\boldsymbol{U}_{k}^{B}\boldsymbol{W}_{k}^{1} = \begin{bmatrix} \boldsymbol{U}_{k}^{B}[1] \\ \boldsymbol{U}_{k}^{B}[2] \\ \dots \\ \boldsymbol{U}_{k}^{B}[M] \end{bmatrix} \boldsymbol{W}_{k}^{1} = \begin{bmatrix} \boldsymbol{U}_{k}^{B}[1]\boldsymbol{W}_{k}^{1} \\ \boldsymbol{U}_{k}^{B}[2]\boldsymbol{W}_{k}^{1} \\ \dots \\ \boldsymbol{U}_{k}^{B}[M]\boldsymbol{W}_{k}^{1} \end{bmatrix} \in R^{M \times L \times D_{1}}$$
(44)

$$\boldsymbol{T}_{k}^{F}\boldsymbol{W}_{k}^{2} = \begin{bmatrix} \boldsymbol{T}_{k}^{F}[1] \\ \boldsymbol{T}_{k}^{F}[2] \\ \dots \\ \boldsymbol{T}_{k}^{F}[M] \end{bmatrix} \boldsymbol{W}_{k}^{2} = \begin{bmatrix} \boldsymbol{T}_{k}^{F}[1]\boldsymbol{W}_{k}^{2} \\ \boldsymbol{T}_{k}^{F}[2]\boldsymbol{W}_{k}^{2} \\ \dots \\ \boldsymbol{T}_{k}^{F}[M]\boldsymbol{W}_{k}^{2} \end{bmatrix} \in R^{M \times L \times D}$$
(45)

$$\boldsymbol{U}_{k}^{B}[i] \in R^{L \times D}, \boldsymbol{W}_{k}^{1} \in R^{D \times D_{1}}, \boldsymbol{T}_{k}^{F}[i] \in R^{L \times D_{1}}, \boldsymbol{W}_{k}^{2} \in R^{D_{1} \times D}, D_{1} = 3072$$

where U_k^B and U_k^F are the input and output matrices of feed-forward network in the k-th self-attention block, respectively, (see Eq. 20), W_k^1 and W_k^2 are weight matrices, b_k^1 and b_k^2 are bias, and $\phi(x)$ is the integral of Gaussian Distribution for x.

6. Output layer

The output of the last attention layer is fed to a fully connected layer with SoftMax function to generate a probability matrix:

$$\boldsymbol{P} = SoftMax(\boldsymbol{H}_{N}\boldsymbol{W}_{N} + \boldsymbol{b}_{N}) \in R^{M \times L \times C_{max}}$$
(46)
$$\boldsymbol{\Gamma} \boldsymbol{H}_{n}[1]\boldsymbol{W}_{n}]$$

$$\boldsymbol{H}_{N}\boldsymbol{W}_{N} = \begin{bmatrix} \boldsymbol{H}_{N}[1] \boldsymbol{W}_{N} \\ \boldsymbol{H}_{N}[2] \boldsymbol{W}_{N} \\ \vdots \\ \boldsymbol{H}_{N}[M] \boldsymbol{W}_{N} \end{bmatrix}, \boldsymbol{H}_{N}[i] \in R^{L \times D}, \boldsymbol{W}_{N} \in R^{D \times C_{max}}$$
(47)

where H_N is the outputted embedding matrix in the *N*-th attention block, W_N and b_N are weight matrix and bias, respectively, and the P(i, j, c) indicates the probability that the *j*-th position of the *i*-th sequence in the masked MSA is predicted as the *c*-th type of amino acid.

7. Loss function

For an individual MSA, the loss function is designed as:

$$Loss_{msa} = \frac{1}{M} \cdot \sum_{i=1}^{M} \{ \frac{1}{|mask(i)|} \cdot \sum_{j \in mask(i)} -logP_{i,j,c(i,j)} \}$$
(48)

where *M* is the number of alignments, mask(i) is a set of masking position in the *i*-th sequence, |mask(i)| is the number of elements in mask(i), c(i, j) is the type index of amino acid for the *j*-th position in the *i*-th sequence before masking, and - $logP_{i,j,c(i,j)}$ is negative log likelihood of the true amino acid at the *j*-th position in the *i*-th sequence under condition of masking.