Highly accurate protein structure prediction with AlphaFold

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High-level impact

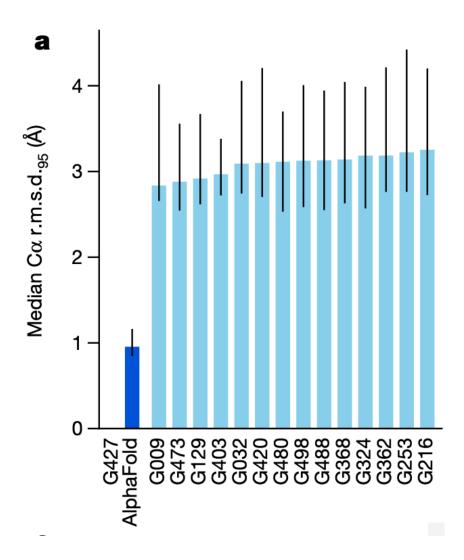
Timeline

- Dec 2018: Alphafold 1 wins CASP
 - CASP: Critical Assessment of protein Structure Prediction
- Nov 2020: Alphafold 2 solves CASP

Impact on drug discovery

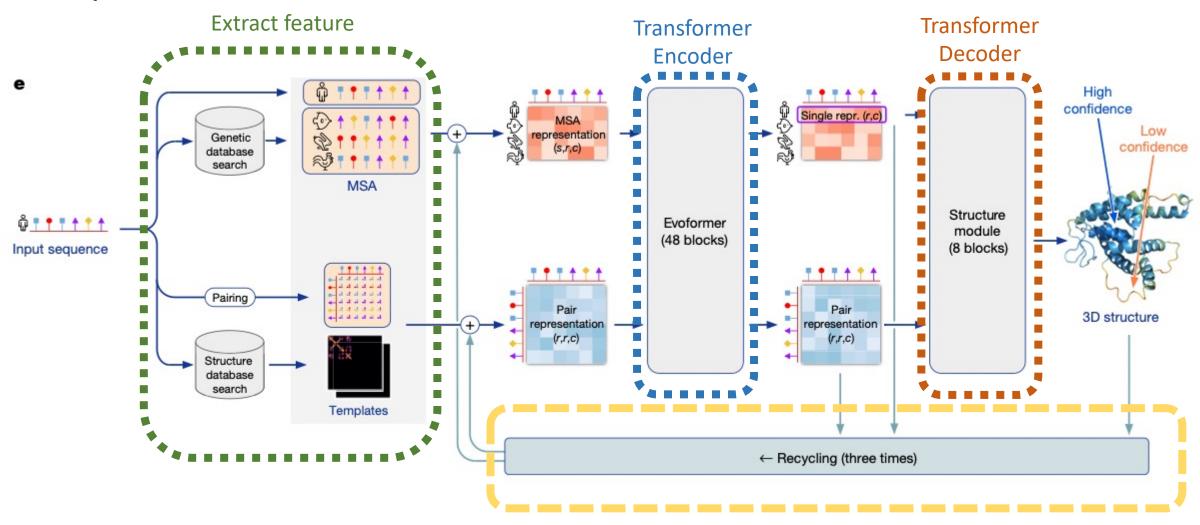
- Universal end-to-end molecular drug discovery pipeline now available

Performance at CASP



First computational method that predict protein structures with atomic accuracy.

Alphafold2



Data Preprocessing

Dataset: 25% PDB dataset + 75% self-distillation Uniclust30 dataset(unlabeled)

MSA: multiple sequence alignments

Template: 3D atom coordinates of homologous structures

MSA Preprocess:

- Filter
- MSA Block Deletion
- MSA Clustering (cluster_profile, extra_msa, mask_msa)
- Residue Cropping(training)

Training & Inference details

MSA resample and ensemble

- MSA block deletion and clustering is stochastic method
- Resampling at training and inference time
- Ensemble at inference time

Recycling

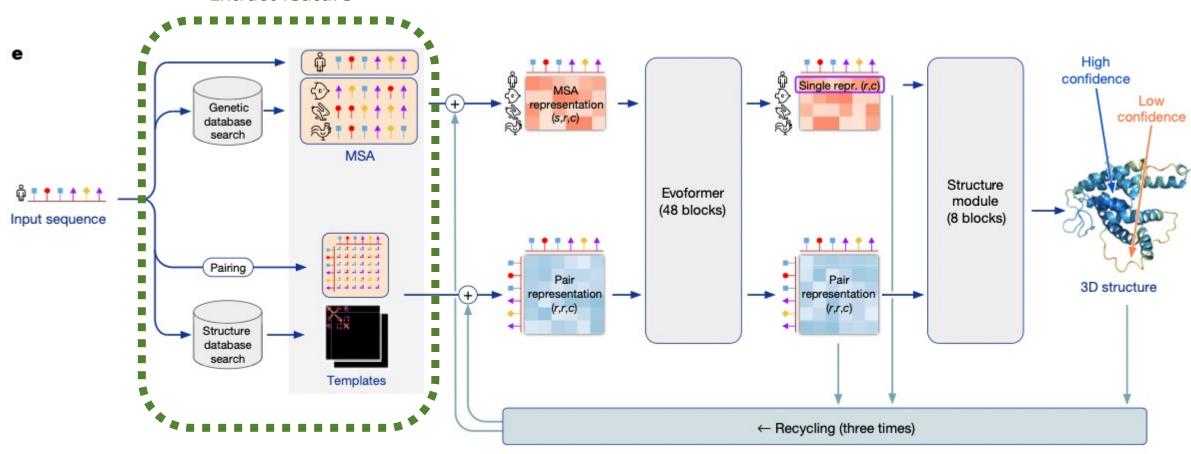
- Random $N \in [0, N_{cycle}]$ at training time
- Fixed N_{cycle} at inference time

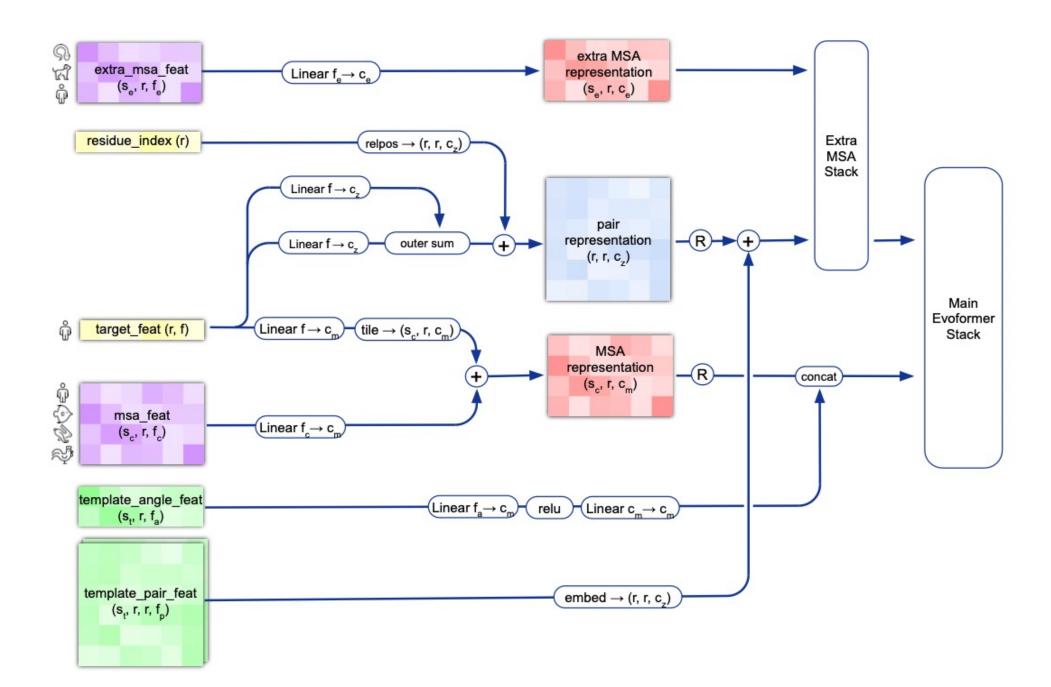
Reducing the memory consumption

- gradient checkpoint at training time
- 'chunk' at inference time

Alphafold2







Algorithm 4 Relative position encoding

$$\begin{aligned} \textbf{def} \ \operatorname{relpos}(\{f_i^{\operatorname{residue_index}}\}, \mathbf{v}_{\operatorname{bins}} &= [-32, -31, \dots, 32]): \\ 1: \ d_{ij} &= f_i^{\operatorname{residue_index}} - f_j^{\operatorname{residue_index}} \\ 2: \ \mathbf{p}_{ij} &= \operatorname{Linear}(\operatorname{one_hot}(d_{ij}, \mathbf{v}_{\operatorname{bins}})) \\ 3: \ \mathbf{return} \ \ \{\mathbf{p}_{ij}\} \end{aligned}$$

Algorithm 17 Template pointwise attention

def TemplatePointwiseAttention($\{\mathbf{t}_{s_t ij}\}, \{\mathbf{z}_{ij}\}, c = 64, N_{\text{head}} = 4$):

1:
$$\mathbf{q}_{ij}^{h} = \text{LinearNoBias}(\mathbf{z}_{ij})$$
 $\mathbf{q}_{ij}^{h} \in \mathbb{R}^{c}, h \in \{1, \dots, N_{\text{head}}\}$
2: $\mathbf{k}_{s_{t}ij}^{h}, \mathbf{v}_{s_{t}ij}^{h} = \text{LinearNoBias}(\mathbf{t}_{s_{t}ij})$ $\mathbf{k}_{s_{t}ij}^{h}, \mathbf{v}_{s_{t}ij}^{h} \in \mathbb{R}^{c}$

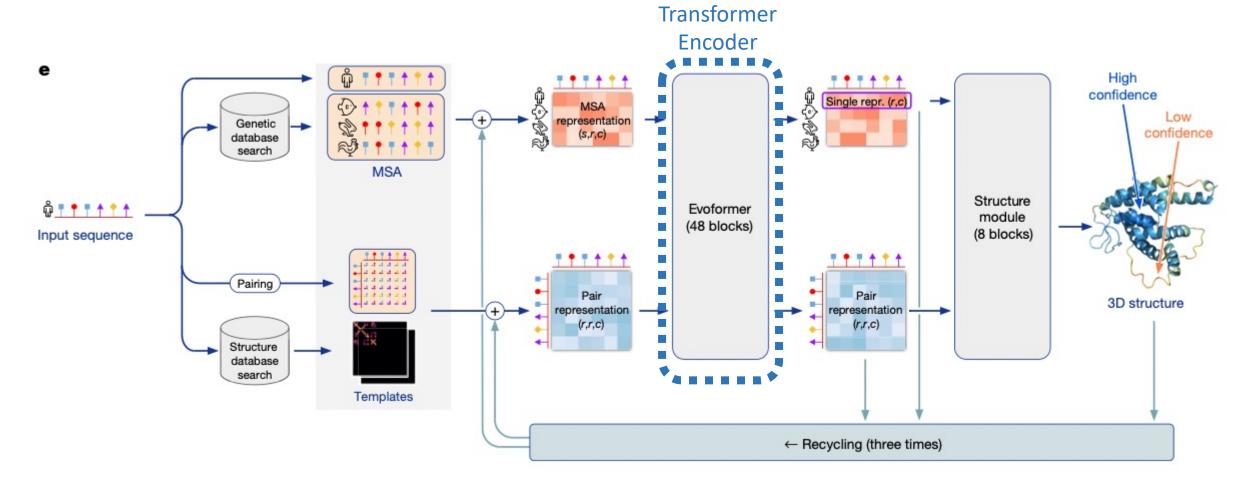
3:
$$a_{s_t i j}^h = \operatorname{softmax}_{s_t} \left(\frac{1}{\sqrt{c}} \ \mathbf{q}_{i j}^{h^{\top}} \mathbf{k}_{s_t i j}^h \right)$$

4:
$$\mathbf{o}_{ij}^h = \sum_{s_t} a_{s_t ij}^h \mathbf{v}_{s_t ij}^h$$

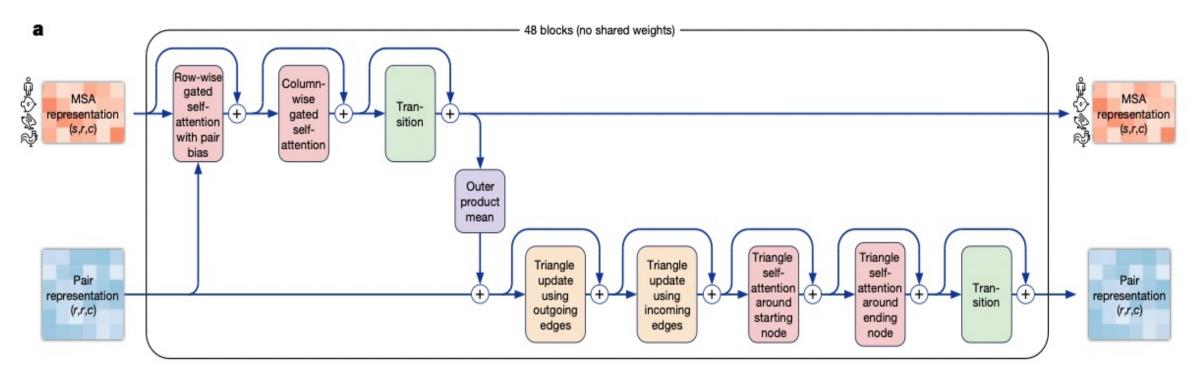
5:
$$\{\tilde{\mathbf{z}}_{ij}\} = \operatorname{Linear}\left(\operatorname{concat}_h(\{\mathbf{o}_{ij}^h\})\right)$$

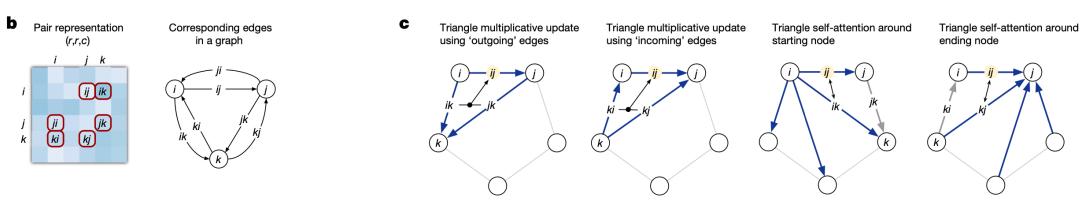
6: **return**
$$\{\tilde{\mathbf{z}}_{ij}\}$$

Alphafold2

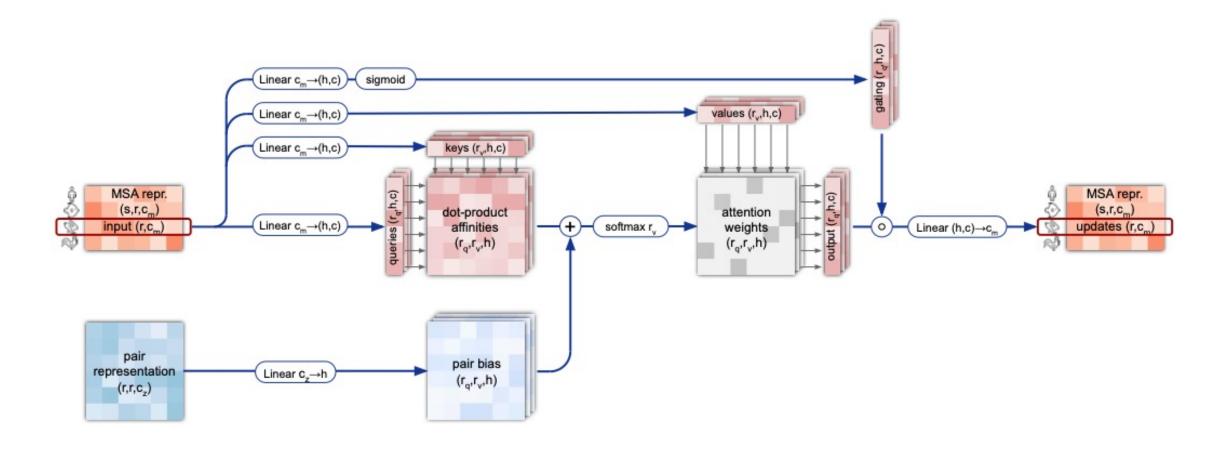


Evoformer





Row-wise gated self-attention



Supplementary Figure 2 MSA row-wise gated self-attention with pair bias. Dimensions: s: sequences, r: residues, c: channels, h: heads.

Algorithm 7 MSA row-wise gated self-attention with pair bias

def MSARowAttentionWithPairBias($\{\mathbf{m}_{si}\}, \{\mathbf{z}_{ij}\}, c = 32, N_{\text{head}} = 8$):

Input projections

1:
$$\mathbf{m}_{si} \leftarrow \text{LayerNorm}(\mathbf{m}_{si})$$

2:
$$\mathbf{q}_{si}^h, \mathbf{k}_{si}^h, \mathbf{v}_{si}^h = \text{LinearNoBias}(\mathbf{m}_{si})$$

3:
$$b_{ij}^h = \text{LinearNoBias}(\text{LayerNorm}(\mathbf{z}_{ij}))$$

4:
$$\mathbf{g}_{si}^h = \operatorname{sigmoid} \left(\operatorname{Linear}(\mathbf{m}_{si}) \right)$$

Attention

5:
$$a_{sij}^h = \operatorname{softmax}_j \left(\frac{1}{\sqrt{c}} \mathbf{q}_{si}^h^\top \mathbf{k}_{sj}^h + b_{ij}^h \right)$$

6:
$$\mathbf{o}_{si}^h = \mathbf{g}_{si}^h \odot \sum_j a_{sij}^h \mathbf{v}_{sj}^h$$

7:
$$\tilde{\mathbf{m}}_{si} = \operatorname{Linear}\left(\operatorname{concat}_h(\mathbf{o}_{si}^h)\right)$$

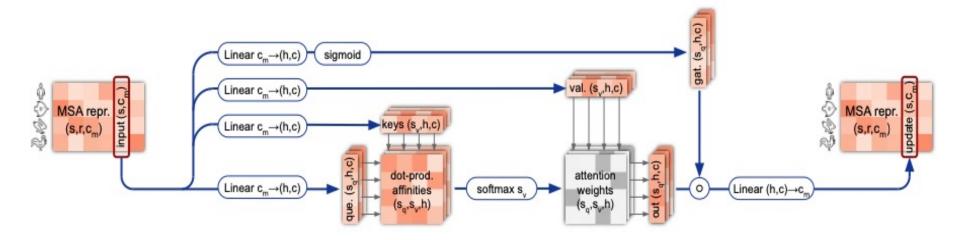
8: **return**
$$\{\tilde{\mathbf{m}}_{si}\}$$

$$\mathbf{q}_{si}^h, \mathbf{k}_{si}^h, \mathbf{v}_{si}^h \in \mathbb{R}^c, \ h \in \{1, \dots, N_{\text{head}}\}$$

$$\mathbf{g}_{si}^h \in \mathbb{R}^c$$

$$ilde{\mathbf{m}}_{si} \in \mathbb{R}^{c_m}$$

Column-wise gated self-attention



Supplementary Figure 3 | MSA column-wise gated self-attention. Dimensions: s: sequences, r: residues, c: channels, h: heads.

Algorithm 8 MSA column-wise gated self-attention

def MSAColumnAttention($\{\mathbf{m}_{si}\}, c = 32, N_{\text{head}} = 8$):

Input projections

1:
$$\mathbf{m}_{si} \leftarrow \text{LayerNorm}(\mathbf{m}_{si})$$

2:
$$\mathbf{q}_{si}^h, \mathbf{k}_{si}^h, \mathbf{v}_{si}^h = \text{LinearNoBias}(\mathbf{m}_{si})$$

3:
$$\mathbf{g}_{si}^h = \operatorname{sigmoid} \left(\operatorname{Linear}(\mathbf{m}_{si}) \right)$$

$\mathbf{q}_{si}^h, \mathbf{k}_{si}^h, \mathbf{v}_{si}^h \in \mathbb{R}^c, \ h \in \{1, \dots, N_{\text{head}}\}$ $\mathbf{g}_{si}^h \in \mathbb{R}^c$

Attention

4:
$$a_{sti}^h = \operatorname{softmax}_t \left(\frac{1}{\sqrt{c}} \mathbf{q}_{si}^{h^{\top}} \mathbf{k}_{ti}^h \right)$$

5:
$$\mathbf{o}_{si}^h = \mathbf{g}_{si}^h \odot \sum_t a_{sti}^h \mathbf{v}_{st}^h$$

6:
$$\tilde{\mathbf{m}}_{si} = \operatorname{Linear}\left(\operatorname{concat}_h(\mathbf{o}_{si}^h)\right)$$

7: **return**
$$\{\tilde{\mathbf{m}}_{si}\}$$

$$ilde{\mathbf{m}}_{si} \in \mathbb{R}^{c_m}$$

Algorithm 13 Triangular gated self-attention around starting node

def TriangleAttentionStartingNode($\{\mathbf{z}_{ij}\}, c = 32, N_{\text{head}} = 4$):

Input projections

1:
$$\mathbf{z}_{ij} \leftarrow \text{LayerNorm}(\mathbf{z}_{ij})$$

2:
$$\mathbf{q}_{ij}^h, \mathbf{k}_{ij}^h, \mathbf{v}_{ij}^h = \text{LinearNoBias}(\mathbf{z}_{ij})$$

3:
$$b_{ij}^h = \text{LinearNoBias}(\mathbf{z}_{ij})$$

4:
$$\mathbf{g}_{ij}^h = \operatorname{sigmoid} \left(\operatorname{Linear}(\mathbf{z}_{ij}) \right)$$

Attention

5:
$$a_{ijk}^h = \operatorname{softmax}_k \left(\frac{1}{\sqrt{c}} \mathbf{q}_{ij}^{h^{\top}} \mathbf{k}_{ik}^h + \mathbf{b}_{jk}^h \right)$$

6:
$$\mathbf{o}_{ij}^h = \mathbf{g}_{ij}^h \odot \sum_k a_{ijk}^h \mathbf{v}_{ik}^h$$

7:
$$\tilde{\mathbf{z}}_{ij} = \operatorname{Linear}\left(\operatorname{concat}_h(\mathbf{o}_{ij}^h)\right)$$

8: **return**
$$\{\tilde{\mathbf{z}}_{ij}\}$$

$$\mathbf{q}_{ij}^h, \mathbf{k}_{ij}^h, \mathbf{v}_{ij}^h \in \mathbb{R}^c, \ h \in \{1, \dots, N_{\text{head}}\}$$

$$\mathbf{g}_{ij}^h \in \mathbb{R}^c$$

$$\tilde{\mathbf{z}}_{ij} \in \mathbb{R}^{c_z}$$

Algorithm 14 Triangular gated self-attention around ending node

def TriangleAttentionEndingNode($\{\mathbf{z}_{ij}\}, c = 32, N_{\text{head}} = 4$):

Input projections

1:
$$\mathbf{z}_{ij} \leftarrow \text{LayerNorm}(\mathbf{z}_{ij})$$

2:
$$\mathbf{q}_{ij}^h, \mathbf{k}_{ij}^h, \mathbf{v}_{ij}^h = \text{LinearNoBias}(\mathbf{z}_{ij})$$

3:
$$b_{ij}^h = \text{LinearNoBias}(\mathbf{z}_{ij})$$

4:
$$\mathbf{g}_{ij}^h = \operatorname{sigmoid} \left(\operatorname{Linear}(\mathbf{z}_{ij}) \right)$$

Attention

5:
$$a_{ijk}^h = \operatorname{softmax}_k \left(\frac{1}{\sqrt{c}} \ \mathbf{q}_{ij}^{h^\top} \ \mathbf{k}_{kj}^h + b_{ki}^h \right)$$

6:
$$\mathbf{o}_{ij}^h = \mathbf{g}_{ij}^h \odot \sum_k a_{ijk}^h \mathbf{v}_{kj}^h$$

7:
$$\tilde{\mathbf{z}}_{ij} = \operatorname{Linear}\left(\operatorname{concat}_{h}\left(\mathbf{o}_{ij}^{h}\right)\right)$$

8: **return**
$$\{\tilde{\mathbf{z}}_{ij}\}$$

$$\mathbf{q}_{ij}^h, \mathbf{k}_{ij}^h, \mathbf{v}_{ij}^h \in \mathbb{R}^c, \ h \in \{1, \dots, N_{\text{head}}\}$$

$$\mathbf{g}_{ij}^h \in \mathbb{R}^c$$

$$\tilde{\mathbf{z}}_{ij} \in \mathbb{R}^{c_z}$$

Algorithm 11 Triangular multiplicative update using "outgoing" edges

def TriangleMultiplicationOutgoing($\{\mathbf{z}_{ij}\}, c = 128$):

1: $\mathbf{z}_{ij} \leftarrow \text{LayerNorm}(\mathbf{z}_{ij})$

2:
$$\mathbf{a}_{ij}, \mathbf{b}_{ij} = \operatorname{sigmoid} \left(\operatorname{Linear}(\mathbf{z}_{ij}) \right) \odot \operatorname{Linear}(\mathbf{z}_{ij})$$

3:
$$\mathbf{g}_{ij} = \text{sigmoid} \left(\text{Linear}(\mathbf{z}_{ij}) \right)$$

4:
$$\tilde{\mathbf{z}}_{ij} = \mathbf{g}_{ij} \odot \operatorname{Linear}(\operatorname{LayerNorm}(\sum_{k} \mathbf{a}_{ik} \odot \mathbf{b}_{jk}))$$

5: **return**
$$\{\tilde{\mathbf{z}}_{ij}\}$$

 $\mathbf{a}_{ij}, \mathbf{b}_{ij} \in \mathbb{R}^c$

$$\mathbf{g}_{ij} \in \mathbb{R}^{c_z}$$

$$ilde{\mathbf{z}}_{ij} \in \mathbb{R}^{c_z}$$

Algorithm 12 Triangular multiplicative update using "incoming" edges

def TriangleMultiplicationIncoming($\{\mathbf{z}_{ij}\}, c = 128$):

1:
$$\mathbf{z}_{ij} \leftarrow \text{LayerNorm}(\mathbf{z}_{ij})$$

2:
$$\mathbf{a}_{ij}, \mathbf{b}_{ij} = \operatorname{sigmoid} \left(\operatorname{Linear}(\mathbf{z}_{ij}) \right) \odot \operatorname{Linear}(\mathbf{z}_{ij})$$

3:
$$\mathbf{g}_{ij} = \operatorname{sigmoid} \left(\operatorname{Linear}(\mathbf{z}_{ij}) \right)$$

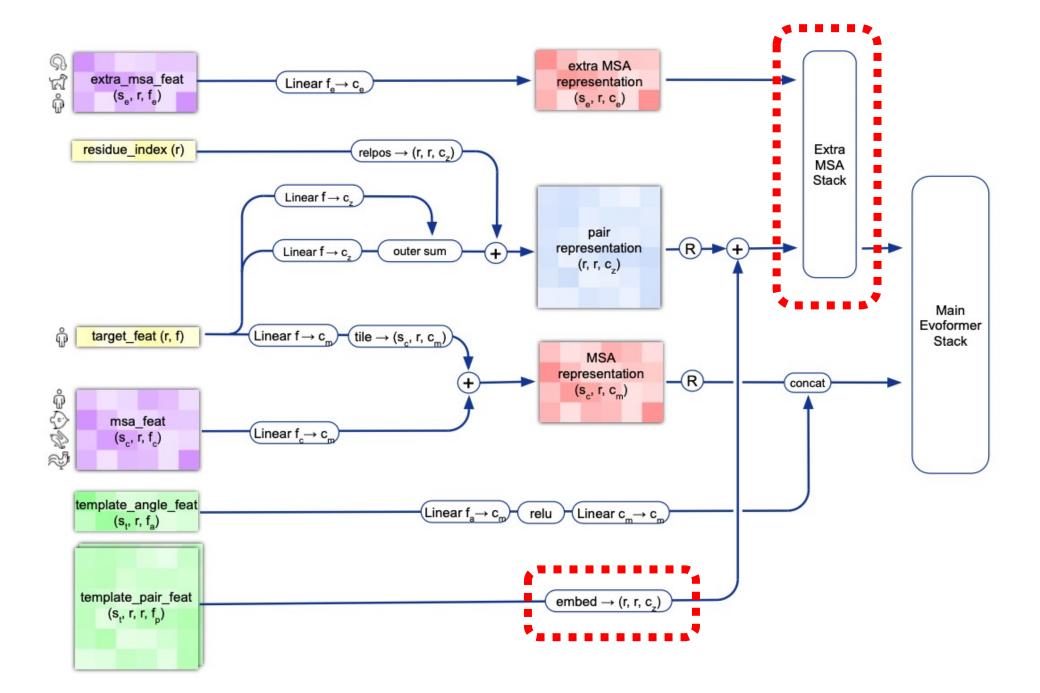
4:
$$\tilde{\mathbf{z}}_{ij} = \mathbf{g}_{ij} \odot \operatorname{Linear}(\operatorname{LayerNorm}(\sum_{k} \mathbf{a}_{ki} \odot \mathbf{b}_{kj}))$$

5: **return**
$$\{\tilde{\mathbf{z}}_{ij}\}$$

 $\mathbf{a}_{ij}, \mathbf{b}_{ij} \in \mathbb{R}^c$

$$\mathbf{g}_{ij} \in \mathbb{R}^{c_z}$$

$$ilde{\mathbf{z}}_{ij} \in \mathbb{R}^{c_z}$$



Algorithm 16 Template pair stack

```
def TemplatePairStack(\{\mathbf{t}_{ij}\}, N_{\text{block}} = 2):

1: for all l \in [1, ..., N_{\text{block}}] do

2: \{\mathbf{t}_{ij}\} += DropoutRowwise<sub>0.25</sub>(TriangleAttentionStartingNode(\{\mathbf{t}_{ij}\}, c = 64, N_{\text{head}} = 4))

3: \{\mathbf{t}_{ij}\} += DropoutColumnwise<sub>0.25</sub>(TriangleAttentionEndingNode(\{\mathbf{t}_{ij}\}, c = 64, N_{\text{head}} = 4))

4: \{\mathbf{t}_{ij}\} += DropoutRowwise<sub>0.25</sub>(TriangleMultiplicationOutgoing(\{\mathbf{t}_{ij}\}, c = 64))

5: \{\mathbf{t}_{ij}\} += DropoutRowwise<sub>0.25</sub>(TriangleMultiplicationIncoming(\{\mathbf{t}_{ij}\}, c = 64))

6: \{\mathbf{t}_{ij}\} += PairTransition(\{\mathbf{t}_{ij}\}, n = 2)

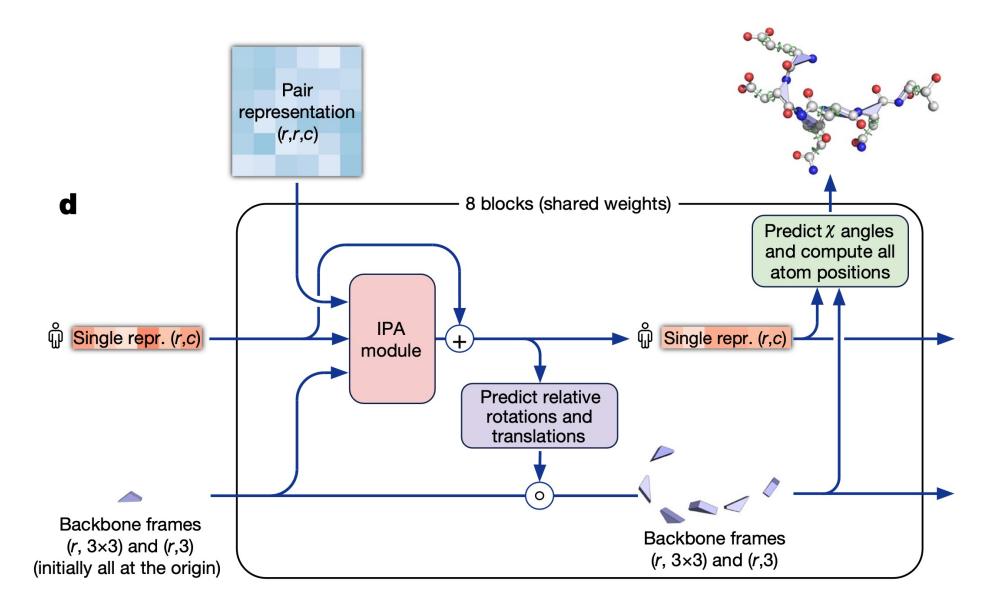
7: end for

8: return LayerNorm(\{\mathbf{t}_{ij}\})
```

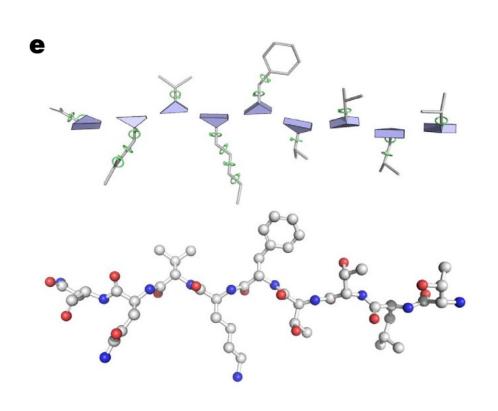
Algorithm 18 Extra MSA stack

```
def ExtraMsaStack(\{\mathbf{e}_{s_ei}\}, \{\mathbf{z}_{ij}\}, N_{block} = 4):
 1: for all l \in [1, \dots, N_{\mathrm{block}}] do
         MSA stack
         \{\mathbf{e}_{s_ei}\} += DropoutRowwise<sub>0.15</sub>(MSARowAttentionWithPairBias(\{\mathbf{e}_{s_ei}\}, \{\mathbf{z}_{ij}\}, \frac{c=8}{}))
         \{\mathbf{e}_{s_e i}\} += MSAColumn Global Attention(\{\mathbf{e}_{s_e i}\}
         \{\mathbf{e}_{s_e i}\} += MSATransition(\{\mathbf{e}_{s_e i}\})
         Communication
  #
         \{\mathbf{z}_{ij}\} += OuterProductMean(\{\mathbf{e}_{s_ei}\})
         Pair stack
         \{\mathbf{z}_{ij}\} += DropoutRowwise<sub>0.25</sub>(TriangleMultiplicationOutgoing(\{\mathbf{z}_{ij}\}))
 6:
         \{\mathbf{z}_{ij}\} += DropoutRowwise<sub>0.25</sub>(TriangleMultiplicationIncoming(\{\mathbf{z}_{ij}\}))
         \{\mathbf{z}_{ij}\} += DropoutRowwise<sub>0.25</sub>(TriangleAttentionStartingNode(\{\mathbf{z}_{ij}\}))
 8:
         \{\mathbf{z}_{ij}\} += DropoutColumnwise<sub>0.25</sub>(TriangleAttentionEndingNode(\{\mathbf{z}_{ij}\}))
         \{\mathbf{z}_{ij}\} += PairTransition(\{\mathbf{z}_{ij}\})
11: end for
12: return \{\mathbf{z}_{ij}\}
```

Structure Module



Frame and torsion angle



- Backbone: $\{T_i = (R_i, t_i)\}$ map from local frame to global frame

- Side chain: the torsion angles are the only degrees of freedom, while all bond angles and bond lengths are fully rigid.

3D Equivariance

12: **return** $\{\tilde{\mathbf{s}}_i\}$

Algorithm 22 Invariant point attention (IPA)

 $\mathbf{def} \ \operatorname{InvariantPointAttention}(\{\mathbf{s}_i\}, \{\mathbf{z}_{ij}\}, \{T_i\}, N_{\mathsf{head}} = 12, c = 16, N_{\mathsf{query points}} = 4, N_{\mathsf{point values}} = 8):$ 1: $\mathbf{q}_i^h, \mathbf{k}_i^h, \mathbf{v}_i^h = \text{LinearNoBias}(\mathbf{s}_i)$ $\mathbf{q}_i^h, \mathbf{k}_i^h, \mathbf{v}_i^h \in \mathbb{R}^c, h \in \{1, \dots, N_{\text{head}}\}$ 2: $\vec{\mathbf{q}}_i^{hp}, \vec{\mathbf{k}}_i^{hp} = \text{LinearNoBias}(\mathbf{s}_i)$ $\vec{\mathbf{q}}_i^{hp}, \vec{\mathbf{k}}_i^{hp}, \in \mathbb{R}^3, \ p \in \{1, \dots, N_{\text{query points}}\}, \ \text{units: nanometres}$ $\vec{\mathbf{v}}_i^{hp} \in \mathbb{R}^3, \ p \in \{1, \dots, N_{\text{point values}}\}, \text{ units: nanometres}$ 3: $\vec{\mathbf{v}}_i^{hp} = \text{LinearNoBias}(\mathbf{s}_i)$ 4: $b_{ij}^h = \text{LinearNoBias}(\mathbf{z}_{ij})$ 5: $w_C = \sqrt{\frac{2}{9N_{\text{query points}}}}$, 6: $w_L = \sqrt{\frac{1}{3}}$ 7: $a_{ij}^h = \operatorname{softmax}_j \left(w_L \left(\frac{1}{\sqrt{c}} \mathbf{q}_i^{h^{\top}} \mathbf{k}_j^h + b_{ij}^h - \frac{\gamma^h w_C}{2} \sum_p \left\| \mathbf{T}_i \circ \vec{\mathbf{q}}_i^{hp} - \mathbf{T}_j \circ \vec{\mathbf{k}}_j^{hp} \right\|^2 \right) \right)$ 8: $\tilde{\mathbf{o}}_i^h = \sum_j a_{ij}^h \mathbf{z}_{ij}$ 9: $\mathbf{o}_i^h = \sum_i a_{ij}^h \mathbf{v}_i^h$ 10: $\vec{\mathbf{o}}_i^{hp} = T_i^{-1} \circ \sum_j a_{ij}^h \left(T_j \circ \vec{\mathbf{v}}_j^{hp} \right)$ 11: $\tilde{\mathbf{s}}_i = \operatorname{Linear}\left(\operatorname{concat}_{h,p}(\tilde{\mathbf{o}}_i^h, \mathbf{o}_i^h, \tilde{\mathbf{o}}_i^{hp}, \left\|\vec{\mathbf{o}}_i^{hp}\right\|)\right)$

Algorithm 23 Backbone update

def BackboneUpdate(\mathbf{s}_i):

1:
$$b_i, c_i, d_i, \vec{\mathbf{t}}_i = \text{Linear}(\mathbf{s}_i)$$

 $b_i, c_i, d_i \in \mathbb{R}, \ \vec{\mathbf{t}}_i \in \mathbb{R}^3$

Convert (non-unit) quaternion to rotation matrix.

2:
$$(a_i, b_i, c_i, d_i) \leftarrow (1, b_i, c_i, d_i) / \sqrt{1 + b_i^2 + c_i^2 + d_i^2}$$

3:
$$R_i = \begin{pmatrix} a_i^2 + b_i^2 - c_i^2 - d_i^2 & 2b_ic_i - 2a_id_i & 2b_id_i + 2a_ic_i \\ 2b_ic_i + 2a_id_i & a_i^2 - b_i^2 + c_i^2 - d_i^2 & 2c_id_i - 2a_ib_i \\ 2b_id_i - 2a_ic_i & 2c_id_i + 2a_ib_i & a_i^2 - b_i^2 - c_i^2 + d_i^2 \end{pmatrix}$$

4:
$$T_i = (R_i, \vec{\mathbf{t}}_i)$$

5: return
$$T_i$$

R: orthogonal matrix, norm=1

Algorithm 24 Compute all atom coordinates

def computeAllAtomCoordinates $(T_i, \vec{\alpha}_i^f, F_i^{\text{aatype}})$:

1:
$$\hat{\vec{\alpha}}_i^f = \vec{\alpha}_i^f / \|\vec{\alpha}_i^f\|$$

2:
$$(\vec{\omega}_i, \vec{\phi}_i, \vec{\psi}_i, \vec{\chi}_{1_i}, \vec{\chi}_{2_i}, \vec{\chi}_{3_i}, \vec{\chi}_{4_i}) = \hat{\vec{\alpha}}_i^f$$

Make extra backbone frames.

3:
$$r_i = F_i^{\text{aatype}}$$

4:
$$T_{i1} = T_i \circ T_{r_i,(\omega \to bb)}^{\text{lit}} \circ \text{makeRotX}(\vec{\omega}_i)$$

5:
$$T_{i2} = T_i \circ T^{\text{lit}}_{r_i,(\phi \to \text{bb})} \circ \text{makeRotX}(\vec{\phi_i})$$

6:
$$T_{i3} = T_i \circ T^{\text{lit}}_{r_i,(\psi \to \text{bb})} \circ \text{makeRotX}(\vec{\psi_i})$$

Make side chain frames (chain them up along the side chain).

7:
$$T_{i4} = T_i \circ T^{\text{lit}}_{r_i,(\chi_1 \to \text{bb})} \circ \text{makeRotX}(\vec{\chi}_{1_i})$$

8:
$$T_{i5} = T_{i4} \circ T_{r_i,(\chi_2 \to \chi_1)}^{\text{lit}} \circ \text{makeRotX}(\vec{\chi}_{2_i})$$

9:
$$T_{i6} = T_{i5} \circ T_{r_i,(\chi_3 \to \chi_2)}^{\text{lit}} \circ \text{makeRotX}(\vec{\chi}_{3_i})$$

10:
$$T_{i7} = T_{i6} \circ T^{\text{lit}}_{r_i,(\chi_4 \to \chi_3)} \circ \text{makeRotX}(\vec{\chi}_{4_i})$$

Map atom literature positions to the global frame.

11:
$$\vec{\mathbf{x}}_i^a = \operatorname{concat}_{f,a'} \left(\{ T_i^f \circ \vec{\mathbf{x}}_{r_i,f,a'}^{\operatorname{lit}} \} \right)$$

12: **return**
$$T_i^f, \vec{\mathbf{x}}_i^a$$

Loss

- FAPE: scores a set of predicted atom coordinates under a set of predicted local frames against the corresponding ground truth atom coordinates and ground truth local frames
- Auxiliary loss: FAPE + torsion Angle loss
- pLDDT(confidence loss): the per-residue IDDT-Cα scores.
- TM score: assessing global structure of predicted protein

Recycle

Algorithm 32 Embedding of Evoformer and Structure module outputs for recycling

def RecyclingEmbedder($\{\mathbf{m}_{1i}\}, \{\mathbf{z}_{ij}\}, \{\vec{\mathbf{x}}_i^{\mathbb{C}^{\beta}}\}$):

Embed pair distances of backbone atoms:

1:
$$d_{ij} = \left\| ec{\mathbf{x}}_i^{\mathbf{C}^eta} - ec{\mathbf{x}}_j^{\mathbf{C}^eta}
ight\|$$

 C^{α} used for glycin

2:
$$\mathbf{d}_{ij} = \text{Linear}(\text{one_hot}(d_{ij}, \mathbf{v}_{\text{bins}} = [3\% \text{ Å}, 5\% \text{ Å}, \dots, 21\% \text{ Å}]))$$

 $\mathbf{d}_{ij} \in \mathbb{R}^{c_z}$

Embed output Evoformer representations:

3:
$$\tilde{\mathbf{z}}_{ij} = \mathbf{d}_{ij} + \text{LayerNorm}(\mathbf{z}_{ij})$$

4:
$$\tilde{\mathbf{m}}_{1i} = \text{LayerNorm}(\mathbf{m}_{1i})$$

5: **return**
$$\{\tilde{\mathbf{m}}_{1i}\}, \{\tilde{\mathbf{z}}_{ij}\}$$

Inference time

```
-V100 GPU
```

```
-seq_len= 256 4.8min
```

$$-seq_len = 2500 18h$$

Ablation Study

