

## Supplementary Information: Diffusion probabilistic models enhance variational autoencoder for crystal structure generative modeling

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### ENCODERS

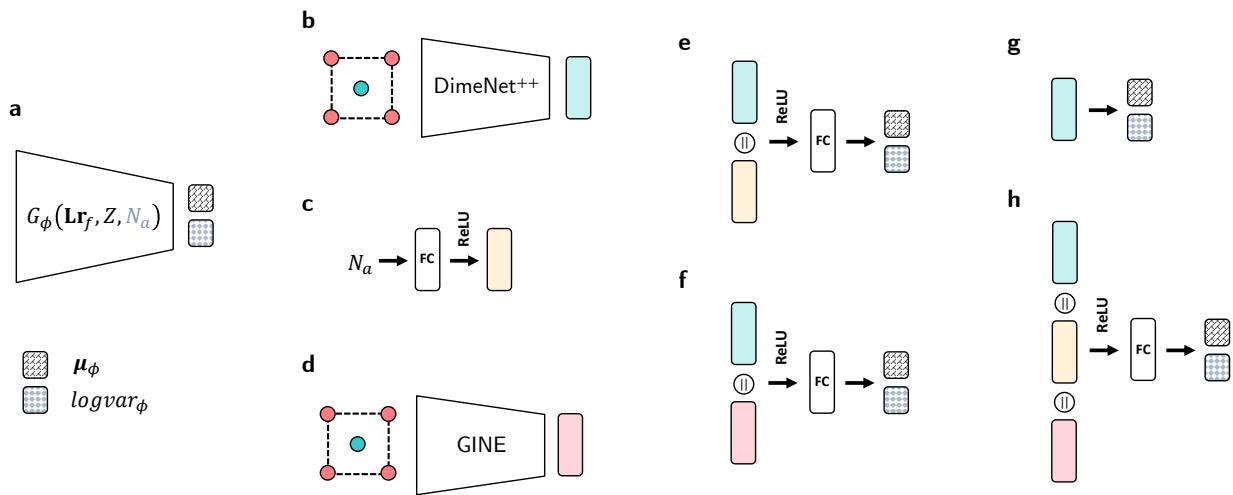


FIG. S1. (a) is the encoder for predicting  $\mu_\phi$  and  $\logvar_\phi$ . (b) and (d) are DimeNet<sup>++</sup> and GINE encoders, respectively, that take pristine crystal structures as inputs. (c) is the  $N_a$  encoder that takes the number of atoms  $N_a$  as an input. (e), (f), (g), and (h) are encoders of DP-CDVAE+ $N_a$ , DP-CDVAE+GINE, DP-CDVAE, and DP-CDVAE+ $N_a$ +GINE models, respectively. Blue, yellow, and pink boxes are the latent features from (b), (c), and (d), respectively, and FC boxes are fully connected layers.

### MODEL EVALUATION

TABLE S1. Reconstruction performance using Eq. 6.

| Models                | Match rate (%) ↑ |           |       | $\langle \delta_{\text{rms}} \rangle \downarrow$ |           |        |
|-----------------------|------------------|-----------|-------|--|-----------|--------|
|                       | Perov-5          | Carbon-24 | MP-20 | Perov-5  | Carbon-24 | MP-20  |
| DP-CDVAE              | 68.72            | 35.86     | 20.40 | 0.0177   | 0.2573    | 0.0715 |
| DP-CDVAE+ $N_a$       | 67.56            | 38.13     | 21.71 | 0.0222   | 0.2719    | 0.0774 |
| DP-CDVAE+GINE         | 49.56            | 35.27     | 19.98 | 0.0822   | 0.2683    | 0.0620 |
| DP-CDVAE+ $N_a$ +GINE | 64.12            | 34.68     | 24.43 | 0.0250   | 0.3212    | 0.0697 |

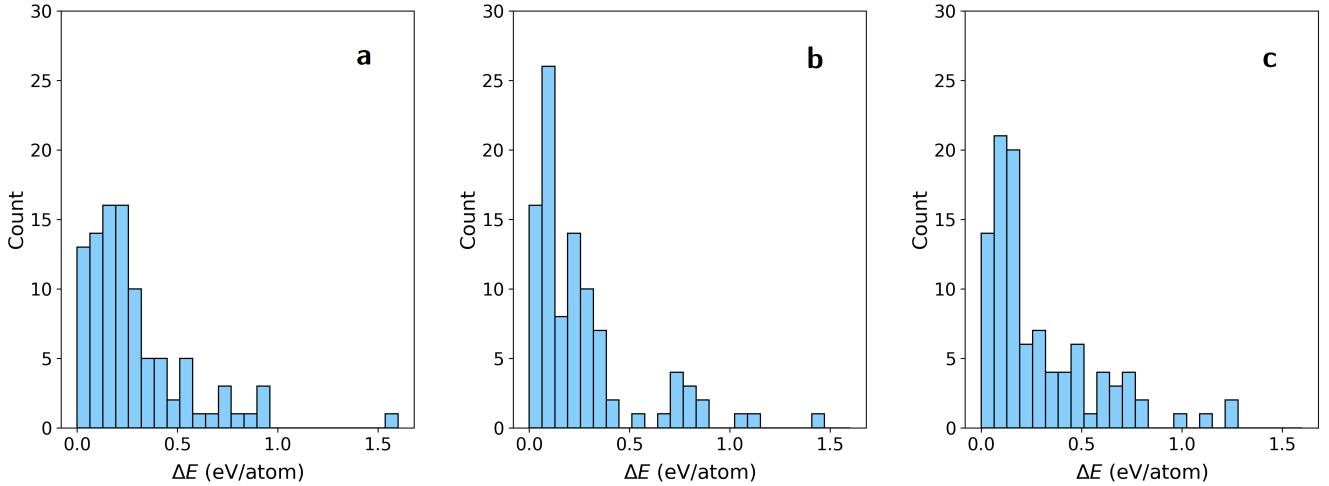


FIG. S2. Histograms of energy difference between generated and relaxed structures of (a) CDVAE, (b) CDVAE+Fourier, and (c) DP-CDVAE models. The modes (highest counting number) of CDVAE, CDVAE+Fourier, and DP-CDVAE models are 19.2 – 32.0, 68.0 – 128, and 64.0 – 128 eV/atom, respectively, where the bin size is 64.0 meV/atom.

## FOURIER FEATURES

We appended the Fourier features to the node attributes for the input of the diffusion network. The Fourier features are the concatenation of  $\sin(2^n\pi \mathbf{r}_t)$  and  $\cos(2^n\pi \mathbf{r}_t)$  where  $n \in \{n_{\min}, \dots, n_{\max}\}$ . For every DP-CDVAE model in the main text,  $n_{\min} = 3$  and  $n_{\max} = 8$ .

## LOSS FUNCTIONS

Similar to the original CDVAE, the total loss function to train the model consists of 5 sub-loss functions: the Kullback–Leibler divergence loss ( $\mathcal{L}_{KLD}$ ) for training  $\mu_\phi$  and  $\logvar_\phi$ , the lattice loss ( $\mathcal{L}_{latt}$ ) for training lattice parameters, the composition loss ( $\mathcal{L}_{comp}$ ) for training  $\mathbf{A}_z$ , the loss for training the number of atoms ( $\mathcal{L}_{N_a}$ ), and the loss from the diffusion network ( $\mathcal{L}_{diff}$ ) for training  $\epsilon_\theta$  and  $\mathbf{A}_\theta$ . In particular, the total loss is

$$\mathcal{L} = \mathcal{L}_{diff} + \lambda_1 \mathcal{L}_{KLD} + \lambda_2 \mathcal{L}_{latt} + \lambda_3 \mathcal{L}_{comp} + \lambda_4 \mathcal{L}_{N_a}, \quad (\text{S1})$$

where  $\lambda_1$ ,  $\lambda_2$ ,  $\lambda_3$ , and  $\lambda_4$  are tunable loss scaling factors,  $\mathcal{L}_{latt}$  is computed from the mean square error of lattice parameters, and  $\mathcal{L}_{comp}$  and  $\mathcal{L}_{N_a}$  are cross-entropy losses.

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