

# Molecular generative model (1)

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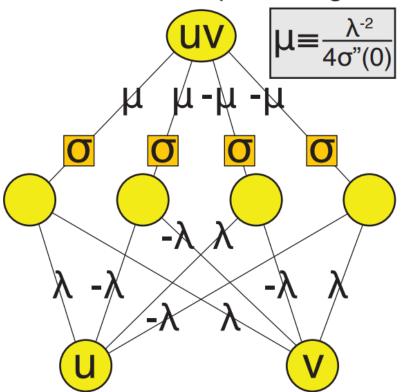
# **Contents**

- Why does deep learning work so well?
- Autoencoder
- Variational autoencoder
- VAE for molecule



MLP can work as a "multiplication gate", with "non-linearity".

Continuous multiplication gate:



Output

$$m(u,v) = \mu \cdot \{ \sigma[\lambda(u+v)] + \sigma[-\lambda(u+v)] - \sigma[\lambda(u-v)] - \sigma[\lambda(-u+v)] \}$$

The nonlinear activation function can be expanded as

$$\sigma(u) \approx \sigma_0 + \sigma_1 \cdot u + \sigma_2 \cdot \frac{u^2}{2}$$

Substituting it into the output function to obtain

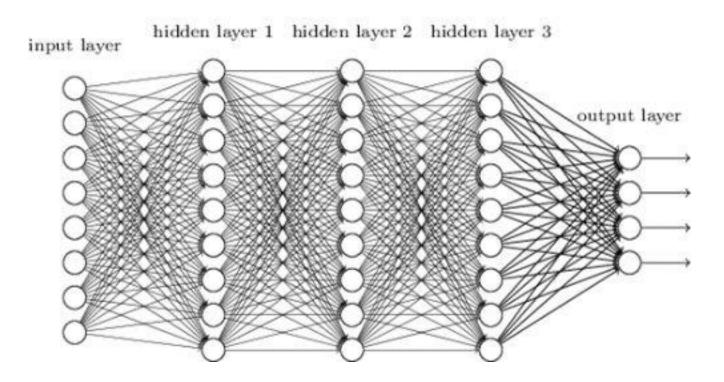
$$m(u,v) = \mu \cdot (\sigma_1 \lambda \cdot [(u+v) + (-u-v) - (u-v) - (-u+v)]$$
  
+  $\sigma_2 \lambda^2 \cdot [(u+v)^2 + (-u-v)^2 - (u-v)^2 - (-u+v)^2)$   
=  $4\mu \cdot \sigma_2 \cdot \lambda^2 \cdot uv$ 

The result is not a linear combination of inputs but multiplication (or nonlinear)!



However, why neural networks need to be deep?

#### Deep neural network





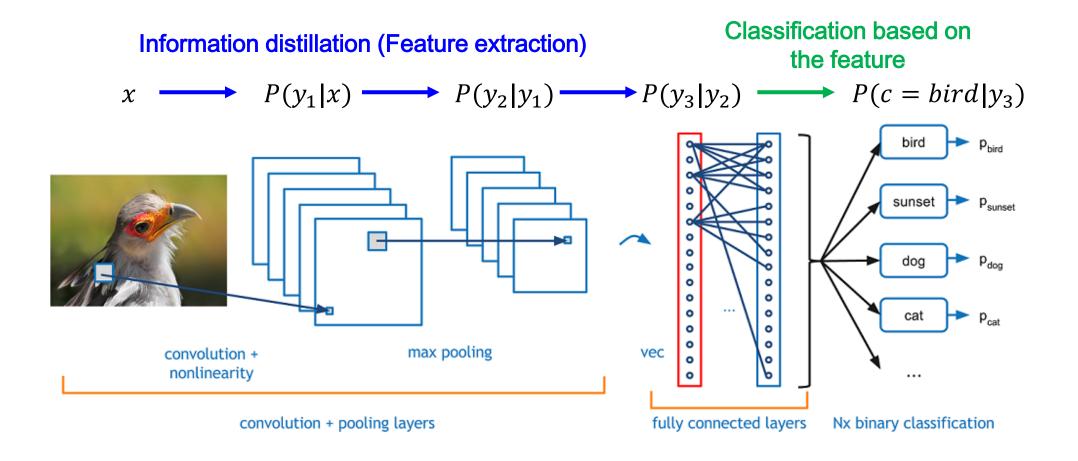
#### Sufficient statistics

Given P(y|x), a sufficient statistic T(x) is defined by the equation P(y|x) = P(y|T(x)) and has played an important role in statistics for almost a century. All the information about y contained in x is contained in the sufficient statistics.

A *minimal sufficient statistic* is some sufficient statistic  $T_*$  which is a sufficient statistic for all other sufficient statistics. This means that if T(y) is sufficient, then there exists some function f such that  $T_*(y) = f(T(y))$ .  $T_*$  can be thought of as an information distiller, optimally compressing the data so as to retain all information relevant to determining y and discarding all irrelevant information.



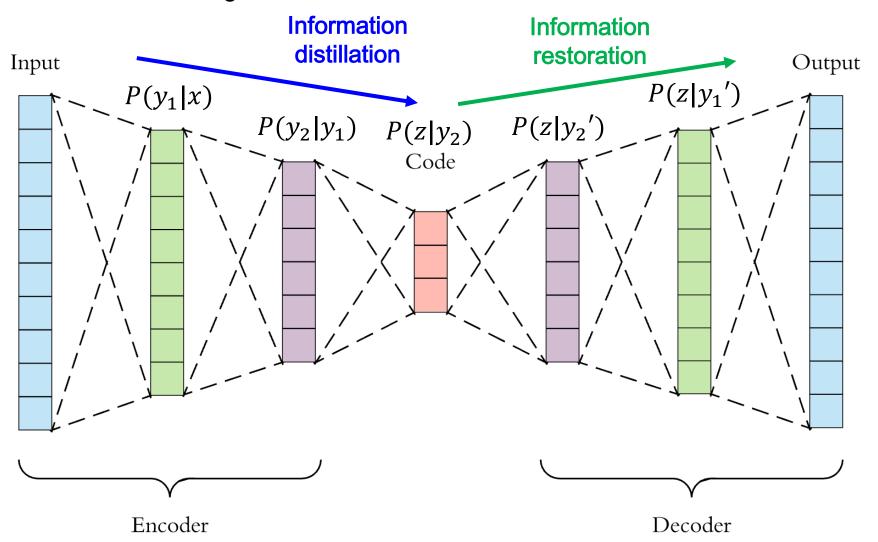
#### Sufficient statistics





## Autoencoder

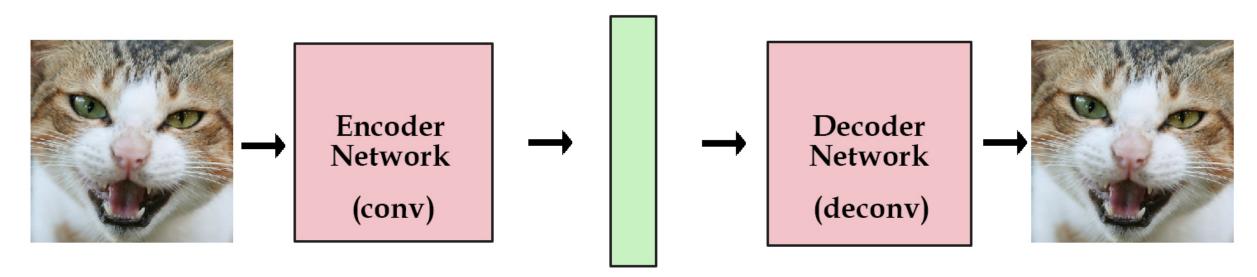
Can we extract the feature without given labels?





#### Autoencoder

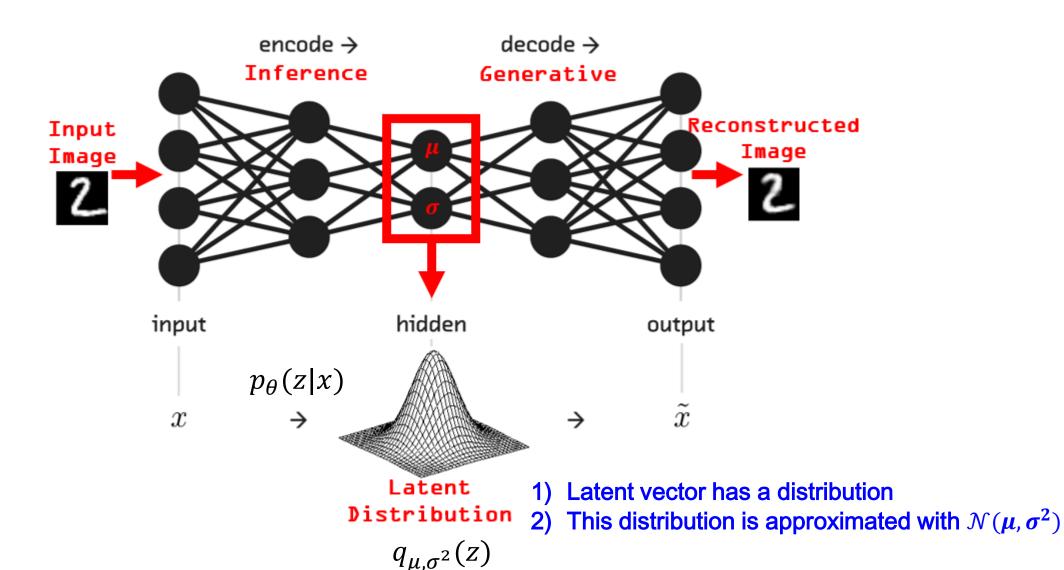
Can we extract the feature without given labels?



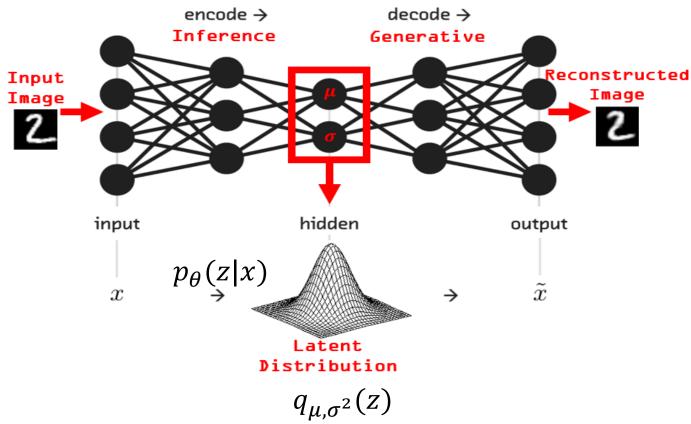
latent vector / variables

: contains the essential information of given input









Our minimization objective

1. Reconstruction of the image

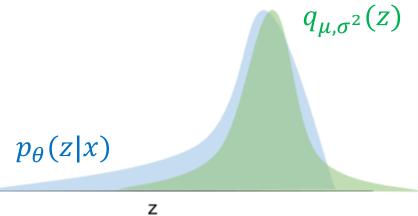
$$\mathcal{L}_{reconst} = \|x - \tilde{x}\|^2$$

2. Variational inference of the latent distribution

$$\mathcal{L}_{VI} = \mathrm{KL}(q_{\mu,\sigma^2}(z)||p_{\theta}(z|x))$$

3. Total loss

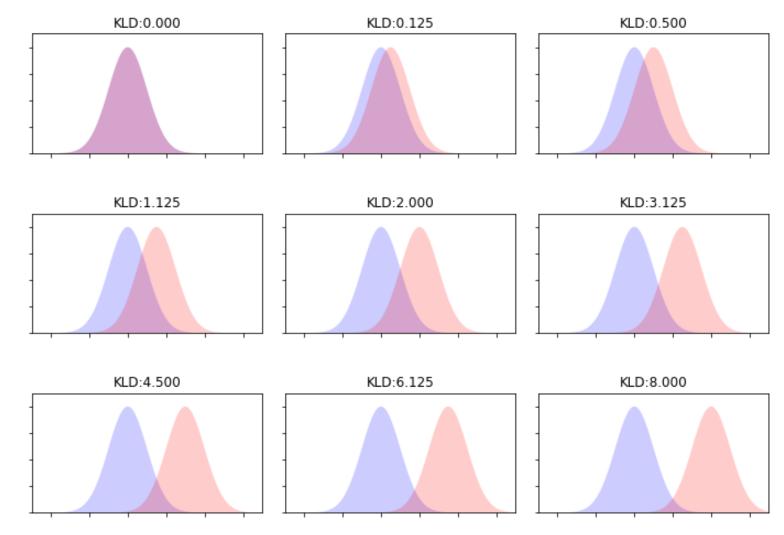
$$\mathcal{L}_{total} = \mathcal{L}_{reconstr} + \mathcal{L}_{VI}$$





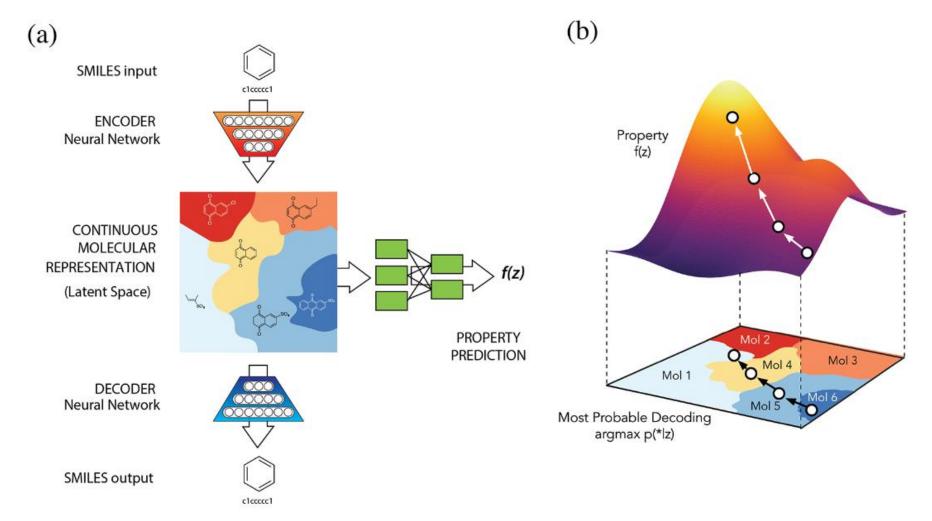
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 $\mathrm{KL}(q_{\mu,\sigma^2}(z)||p_{\theta}(z|x))$ 



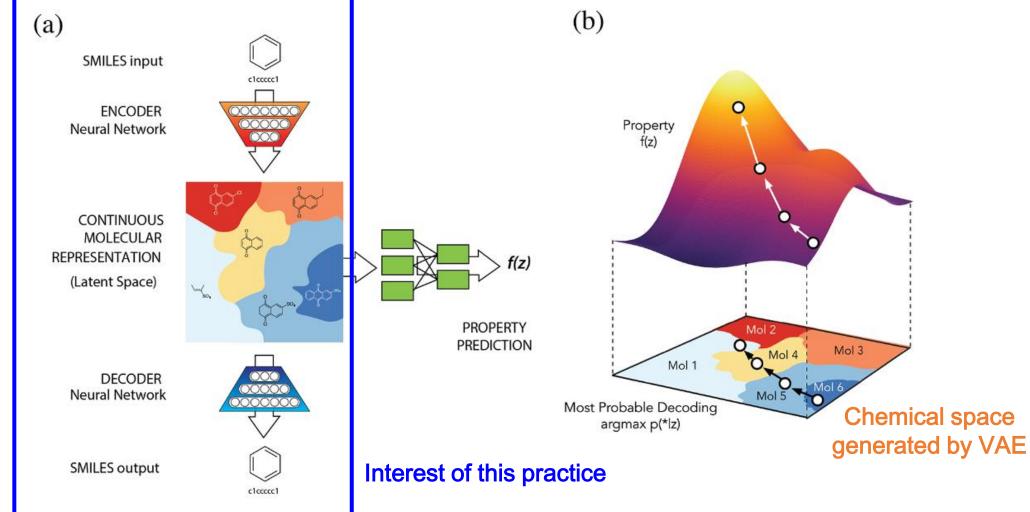


#### ChemicalVAE





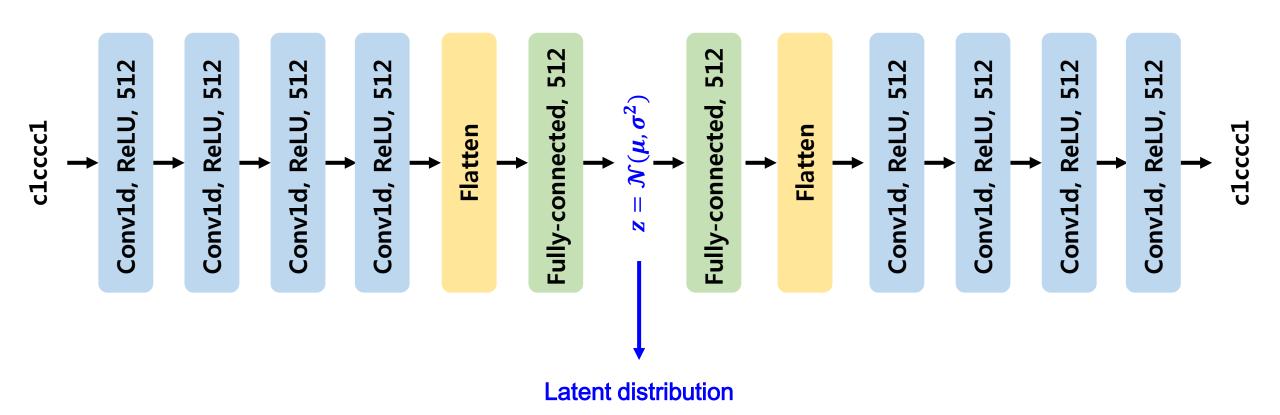
#### ChemicalVAE





Gómez-Bombarelli, Rafael, et al. "Automatic chemical design using a data-driven continuous representation of molecules." ACS central science 4.2 (2018): 268-276. 13

**ChemicalVAE** 





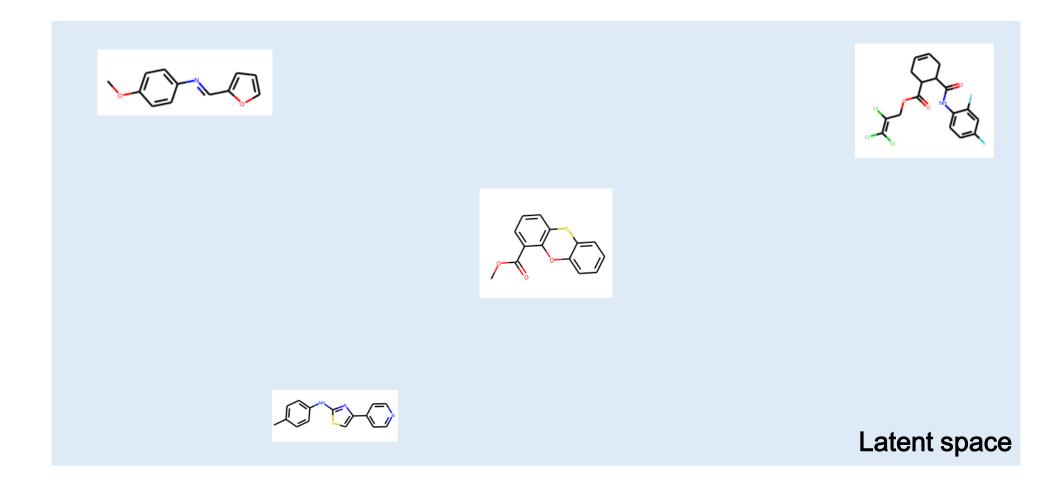
#### ChemicalVAE

Input SMILES	Reconstructed SMILES
CC(C)(C)c1ccccc1OCC(=O)Nc1cccc(C(=O)[O-])c1	CC(C)(C)c1ccccc1OCC(=O)Nc1cccc(C(=O)[O-])c1
COc1ccc(-c2nnc(SCC(=O)OC3CCCCC3)n2C)cc1	COc1ccc(-c2nnc(SCC(=O)OC3CCCCC3)n2C)cc1
C=C(C)C(=O)OCC <b>S</b> C	C=C(C)C(=O)OCC <b>C</b> C
CNc1oc(-c2ccc(Cl)cc2)nc1S(=O)(=O)c1ccc(C)cc1	COcloc(-c2ccc(Cl)cc2)nc1S(=O)(=O)c1ccc(C)cc1
CC(=O)N(c1ccccc1C)S(=O)(=O)c1ccc(CI)cc1	CC(=O)N(c1cccc1C)S(=O)(=O)c1ccc(CI)cc1
O = C(COC(=O)c1cccc(S(=O)(=O)N2CCCCC2)c1)Nc1ccc(F)cc1F	O = C(COC(=O)c1cccc(S(=O)(=O)N2CCCCC2)c1)Nc1ccc(F)cc1F
CCOc1ccc(N2C3=[N+](CCCCC3)CC2(O)c2ccc([N+](=O)[O-])cc2)cc1	CCOc1ccc(N2C3=[N+](CCCCC3)CC2(O)c2ccc([N+](=O)[O-])cc2)cc1
CC(C)(C)c1csc(NC(=O)COc2ccc(F)cc2)n1	CC(C)(C)c1csc(NC(=O)COc2ccc(F)cc2)n1
O=C1CN(C(=O)C=Cc2ccc3c(c2)OCO3)c2ccccc2 <b>N</b> 1	O=C1CN(C(=O)C=Cc2ccc3c(c2)OCO3)c2ccccc21
COC(=O)c1ccccc1NC(=O)CSc1ccc(C)cc1C	COC(=O)c1ccccc1NC(=O)CSc1ccc(C)cc1C

Recunstruction accuracy (mol) = 70%, accuracy(character) > 99%

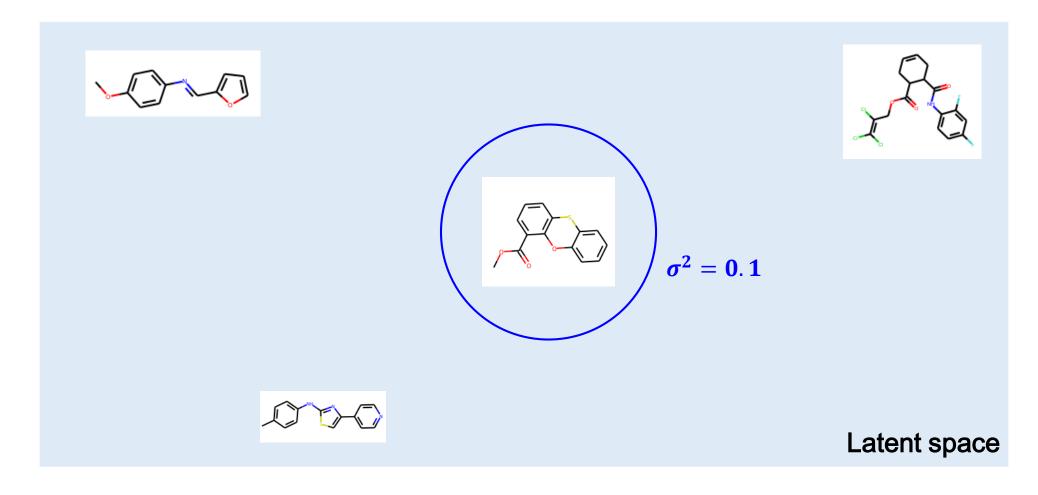


#### ChemicalVAE

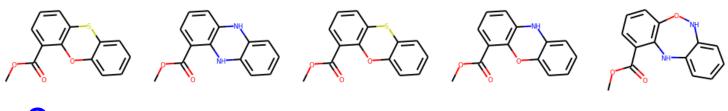




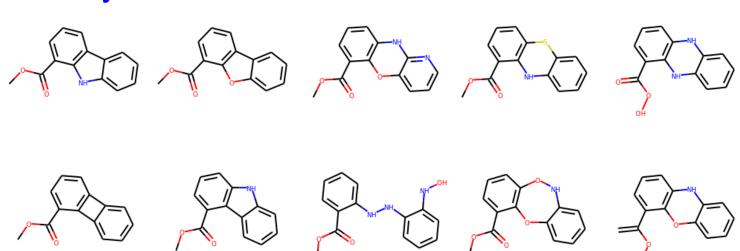
#### Searching latent space

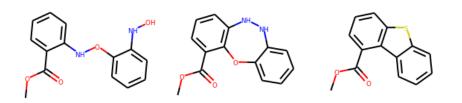






#### Query

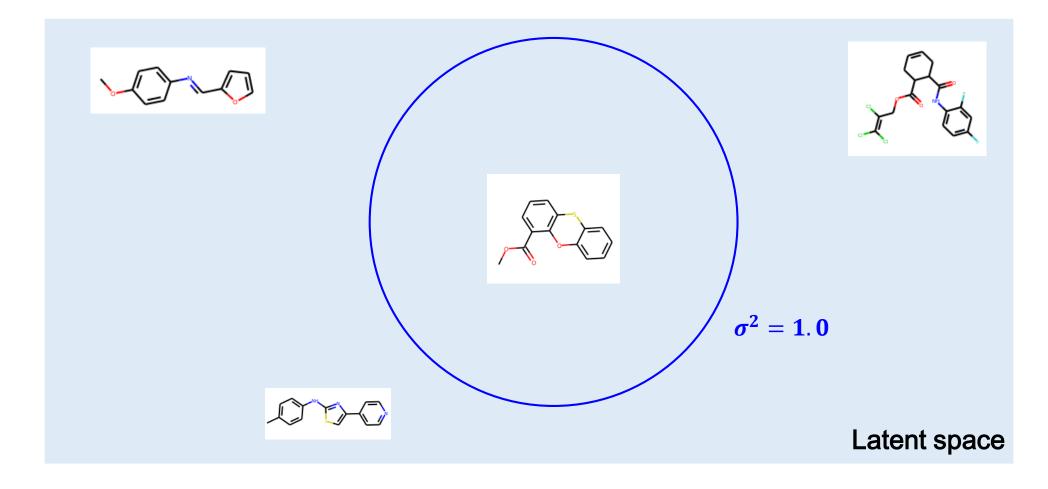




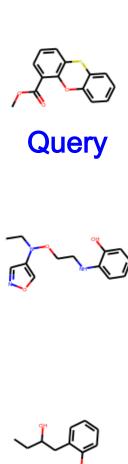
 $\sigma^2 = 0.1$ 

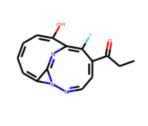
$$\sigma^2 = 0.1$$

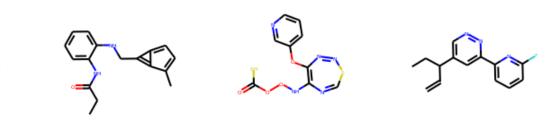
#### Searching latent space

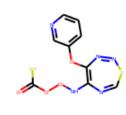


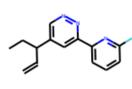


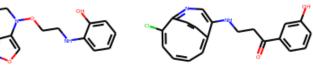


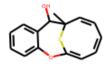


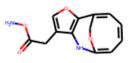








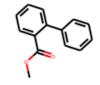


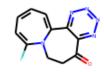








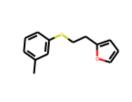


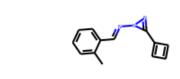


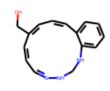




$$\sigma^2=1.0$$





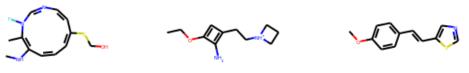


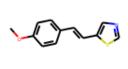


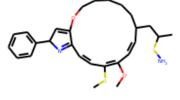
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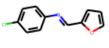


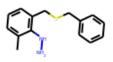


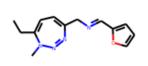


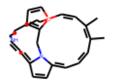


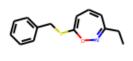


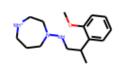


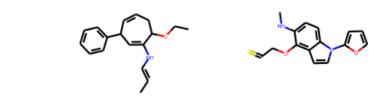


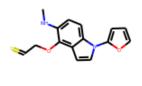


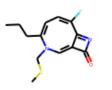












$$\sigma^2 = 1.0$$

# Questions

- 1. What is the meaning of latent vectors?
  - → It may contains the "structural information", whose original representation is SMILES.



# **Questions**

- 1. What is the meaning of latent vectors?
  - → It may contains the "structural information", whose original representation is SMILES.
- 2. Is using SMILES relevant?

C1(C2=NC(NC3=CC=CC=C3)=NC=C2)=CN=CC=C1

$$CC(C=CC=C1)=C1NC2=NC=CC(C3=CN=CC=C3)=N2$$

**Gleevec (Imatinib)** 

HN N

CC(C=CC(NC(C1=CC=C(CN2CCN(C)CC2)C=C1)=O)

KAIST =C3)=C3NC4=NC=CC(C5=CN=CC=C5)=N4

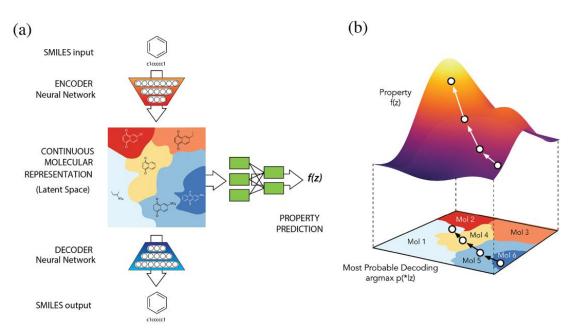
## Questions

- 1. What is the meaning of latent vectors?
  - → It may contains the "structural information", whose original representation is SMILES.
- 2. Is using SMILES relevant?
  - → Graph generative model might be better
- 3. How can we generate the molecules with desired properties?
  - → Conditional VAE, training VAE jointly with a property controler

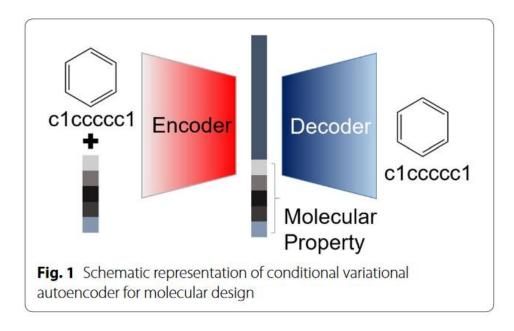


## **Next time**

#### Conditional VAE, training VAE jointly with a property controler



Gómez-Bombarelli, Rafael, et al. "Automatic chemical design using a data-driven continuous representation of molecules." *ACS central science* 4.2 (2018): 268-276.



Lim, Jaechang, et al. "Molecular generative model based on conditional variational autoencoder for de novo molecular design." *arXiv preprint arXiv:1806.05805* (2018).

