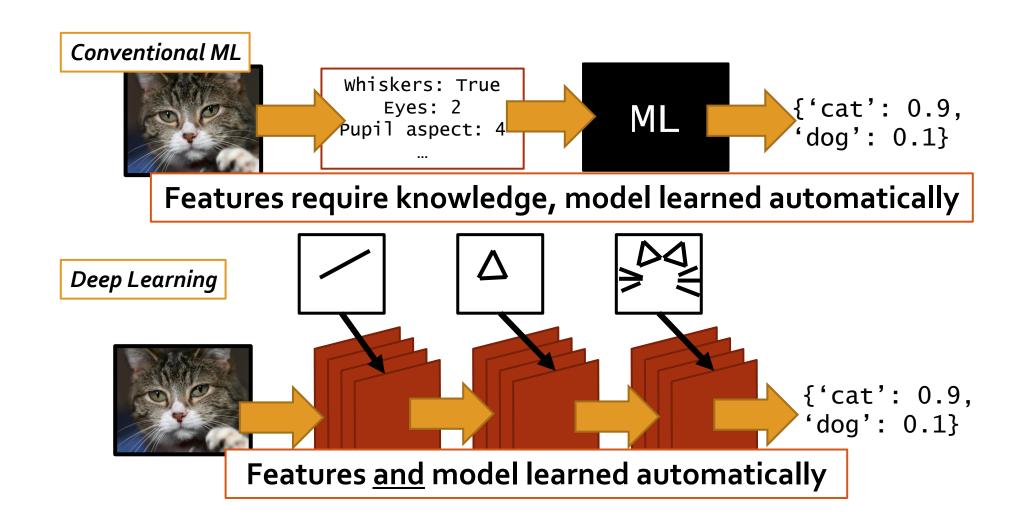
DEEP LEARNING FOR MOLECULAR DATA

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26 January 2021

What is different about deep learning?



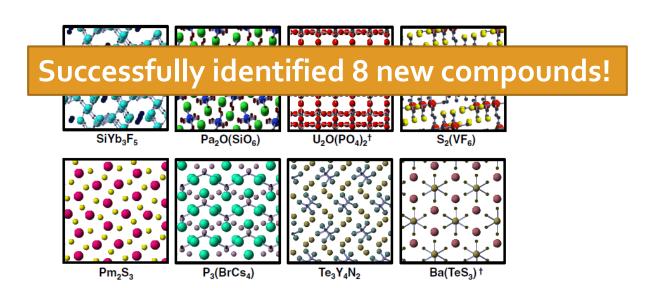
FIRST, WHAT CONVINCED ME DEEP LEARNING WAS WORTHWHILE

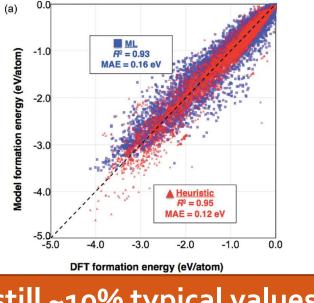
Predicting Crystalline Materials

Goal: Identify new crystalline materials

Model: Given, composition predict ΔH_f

Initial attempt: 2014





Error still ~10% typical values

Can we do even better with deep neural networks?

Ref: Meredig et al. PRB. (2014), 094104. doi: 10.1103/PhysRevB.89.094104

Training Data

Data Source: Open Quantum Materials Database ~470k DFT Calculations

Why OQMD?

- ✓ Open ✓ Large
- Contains unstable materials

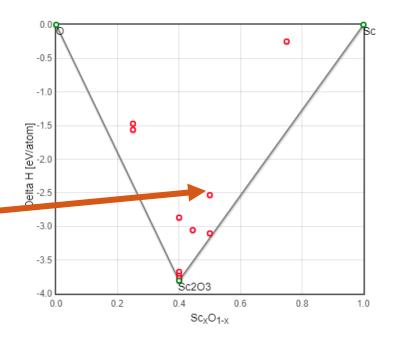
Training Set:

Subset: Lowest energy structure

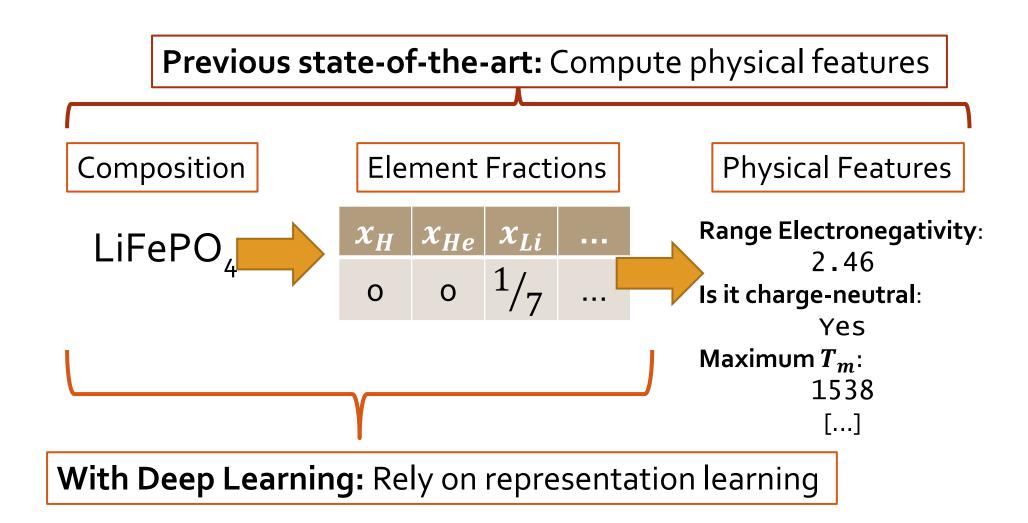
<u>Size:</u> 230336

Input: Composition

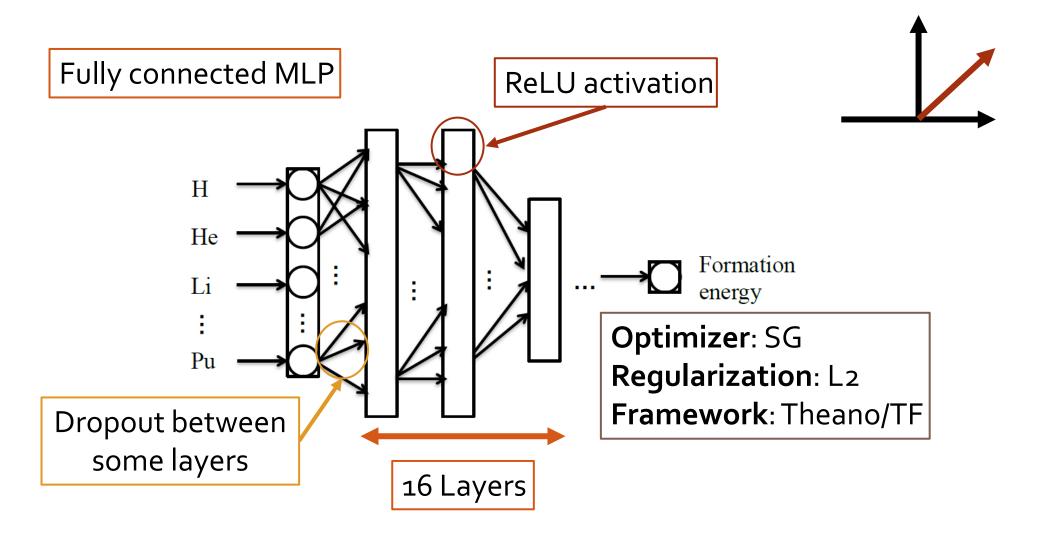
Output: Formation Enthalpy ΔH_f



Input Features

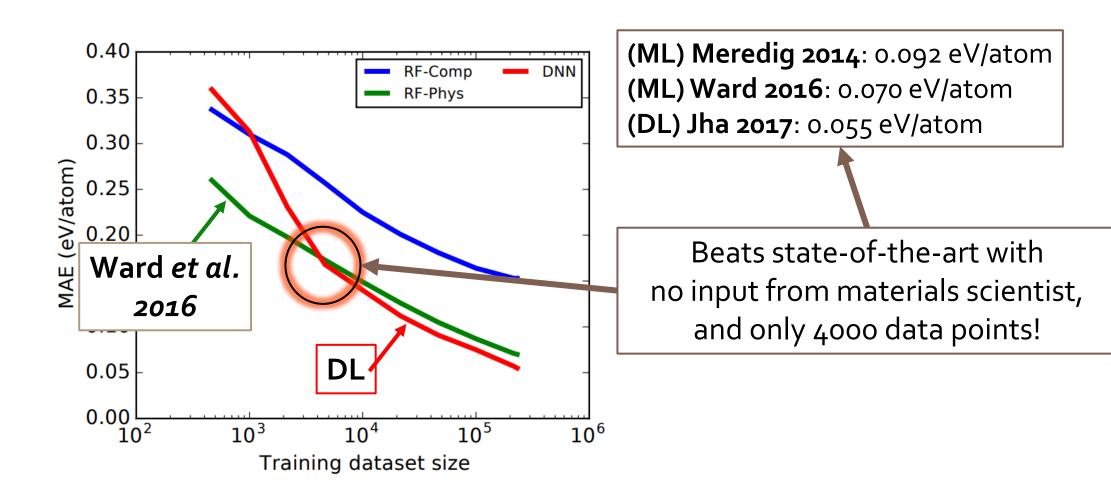


ElemNet Architecture: Nothing Fancy

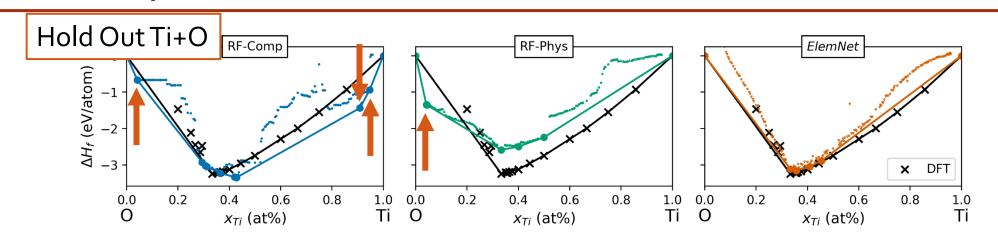


Ref: Jha et al., Sci. Rep. (2020) 17593

Better than conventional learning?

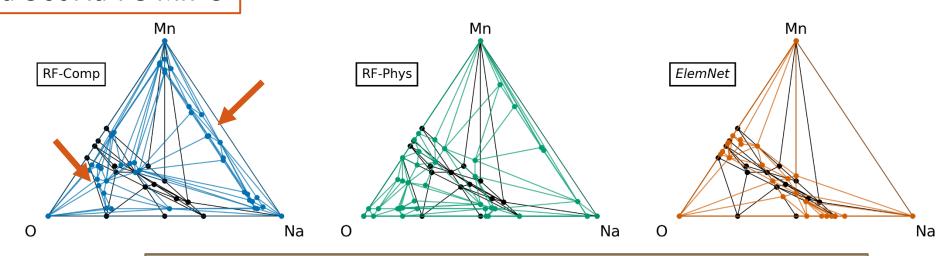


Can DL interpolate between elements?



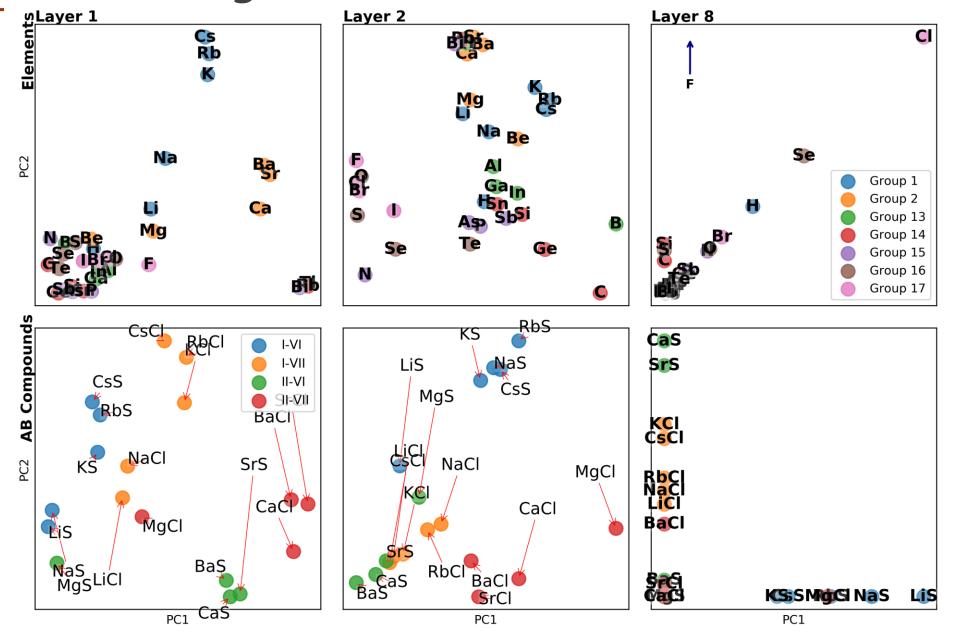
Hold Out Na-Fe-Mn-O

Deep Learning Yields Fewer Spurious Predictions

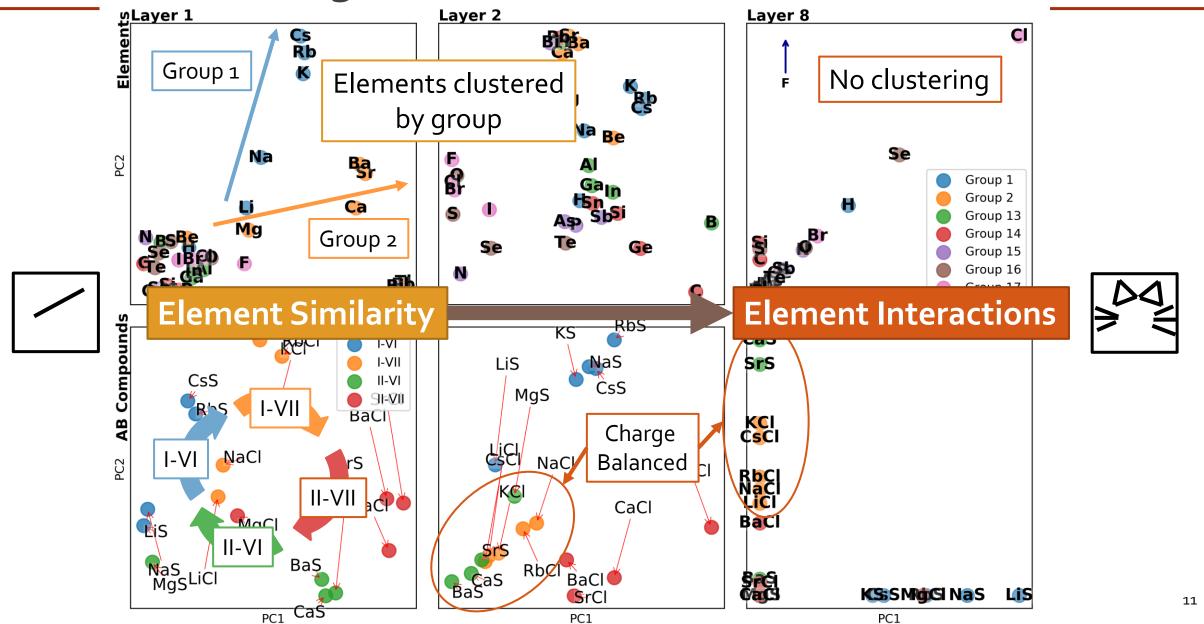


How is this possible? Model hasn't seen Ti-O bonds

How is it working so well?

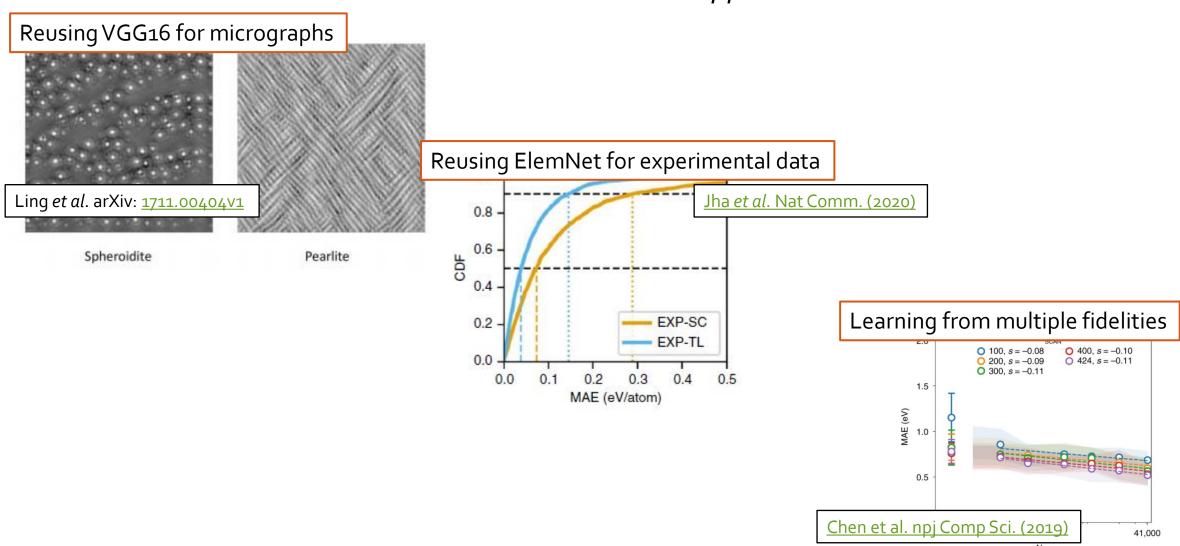


How is it working so well?



Big Opportunity: Transfer Learning

Features learned on one dataset applicable to others



Deep Learning is not Cure-All

Table 3: Summary of performances(test subset): conventional methods versus graph-based methods. Graph-based models outperform conventional methods on 11/17 datasets.

Rost performances -

Rost performances

	Category	Dataset	\mathbf{Metric}	Best performances -	Best performances -
	Category		Metric	conventional methods	graph-based methods
		QM7	MAE	KRR(CM): 10.22	DTNN: 8.75
	Quantum Mechanics	QM7b	MAE	KRR(CM): 1.05	DTNN: 1.77*
_ , ,		<i>C</i> 1 .		Multitask: 0.0150	MPNN: 0.0143
For benchmark problems of learning				Multitask(CM): 4.35	DTNN: 2.35
				XGBoost: 0.99	MPNN: 0.58
from molecular data,				XGBoost: 1.74	MPNN: 1.15
CON				XGBoost: 0.799	GC: 0.655
conventional ML better for 6/17 cases ${}^{\!$				Logreg: 0.129	GC: 0.136
		MUV	AUC-PRC	Multitask: 0.184	Weave: 0.109
	Biophysics	HIV	AUC-ROC	KernelSVM: 0.792	GC: 0.763
		BACE	AUC-ROC	RF: 0.867	Weave: 0.806
		PDBbind(full)	RMSE	RF(grid): 1.25	GC: 1.44
	BBBP	AUC-ROC	KernelSVM: 0.729	GC: 0.690	
Physiology		Tox21	AUC-ROC	KernelSVM: 0.822	GC: 0.829
		ToxCast	AUC-ROC	Multitask: 0.702	Weave: 0.742
		SIDER	AUC-ROC	RF: 0.684	GC: 0.638
		ClinTox	AUC-ROC	Bypass: 0.827	Weave: 0.832

^{*} As discussed in section 4.4, DTNN outperforms KRR(CM) on 14/16 tasks in QM7b while the mean-MAE is skewed due to different magnitudes of labels.

Ref: Wu et al. Chem Sci. (2017) 10.1039/C7SC02664A

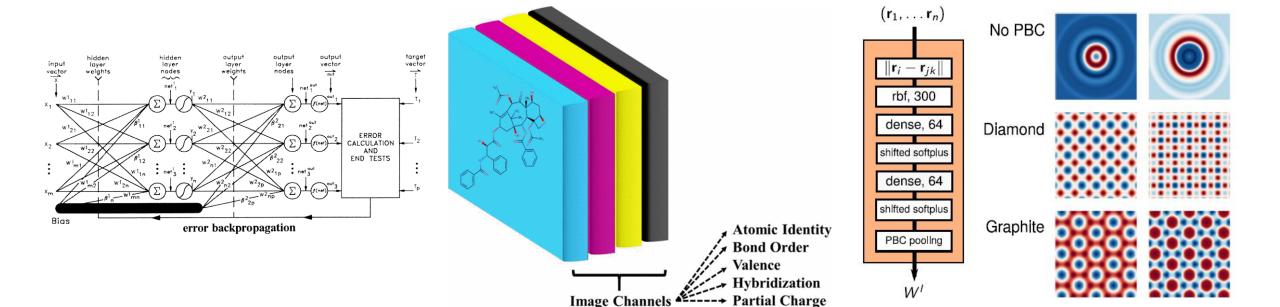
MANY APPROACHES FOR DEEP LEARNING WITH MOLECULAR DATA

How do we do deep learning with molecules?

As a normal supervised learner

Repurposing Image Models

Message-Passing Networks

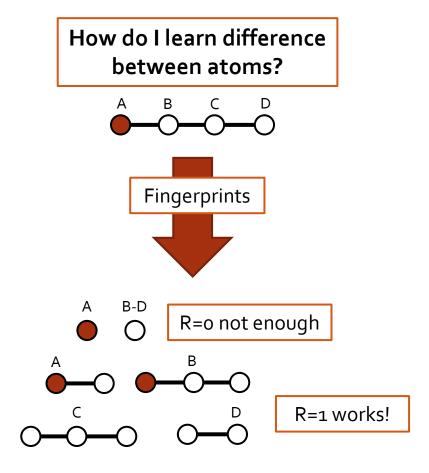


Ex: <u>Ulmer et al. CTPS. (1998), 311.</u>

Ex: Goh et al. IEEE WACV. (2018)

Ex: Schütt et al. JCP, (2018)

Our focus: Message-Passing Networks

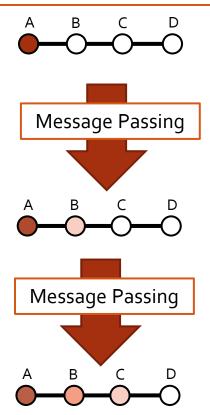


Problems:

- 1. Combinatorics: Many features!
- 2. No similarity between groups (is A:B > B:C)

Our focus: Message-Passing Networks

How do I learn difference between atoms?



Atomic fingerprints are learned by "convolution"

Message Passing is Two Steps

Step 1: Message generation

$$m(B) = f_M(\overset{\mathsf{B}}{\bigcirc}, \overset{\mathsf{A}}{\bullet}, -) + f_M(\overset{\mathsf{B}}{\bigcirc}, \overset{\mathsf{C}}{\bigcirc}, -) = \bigcirc$$

Step 2: Update

$$B^{\{n+1\}} = f_{II}(O, O) = O$$

Solution: Message passing gives continuous features
1. Combinatorics: Many features! Fixed feature count
2. No similarity between groups (is A:B > B:C) Distances

New Challenge: Learning update and message functions

Formalizing Message Passing Neural Networks

A generalized form of neural networks for graph data, introduced by Gilmer et al. (Google)

Existing strategies mostly variants of

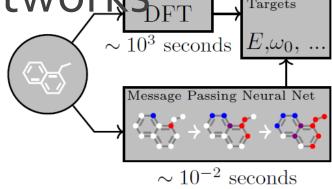


Figure 1. A Message Passing Neural Network predicts quantum properties of an organic molecule by modeling a computationally expensive DFT calculation.

•
$$m_v^{t+1} = \sum_{w \in N(v)} \mathbf{M_t}(h_v^t, h_w^t, e_{vw})$$

• $m_v^{t+1} = \sum_{w \in N(v)} M_t(h_v^t, h_w^t, e_{vw})$ 1. Gather messages from neighboring nodes

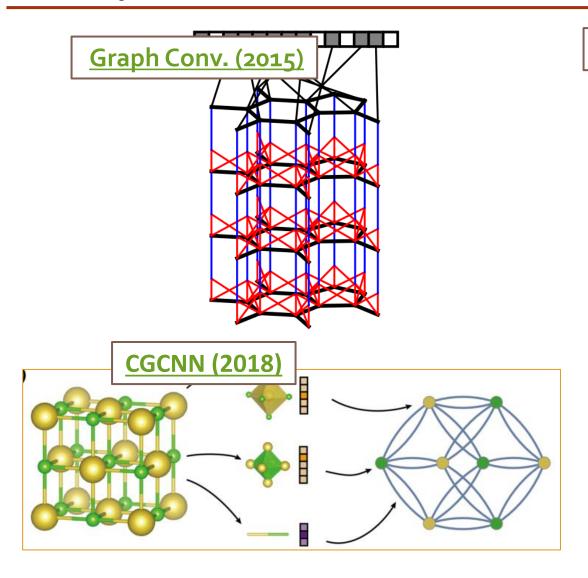
•
$$h_v^{t+1} = U_t(h_v^t, m_v^{t+1})$$

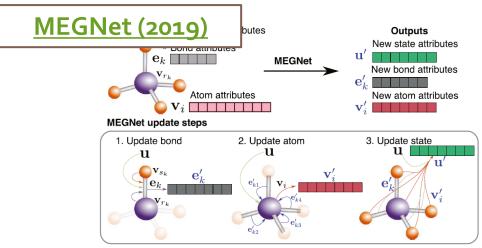
2. **Update** node state given messages

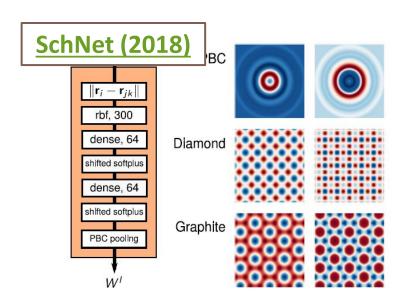
•
$$\hat{y} = \mathbf{R}(\{h_v^T | v \in G\})$$

3. **Readout** graph properties given node states

Many variations of MPNNs

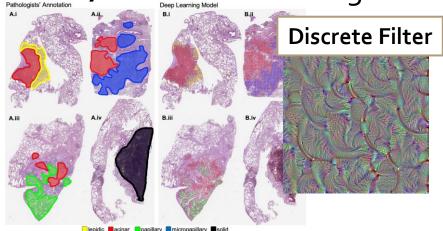






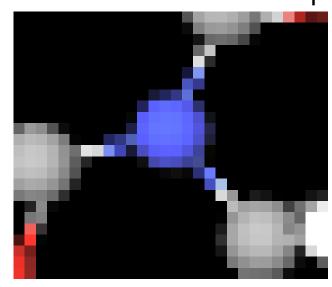
SchNet: Continuous Convolutions

Opportunity: Convolutions are great!

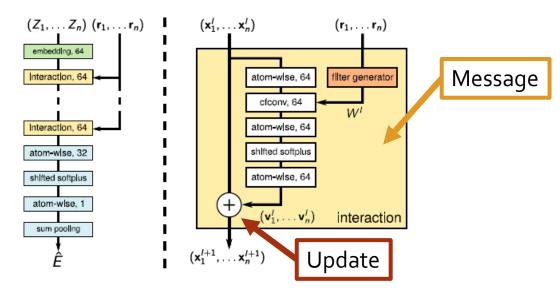


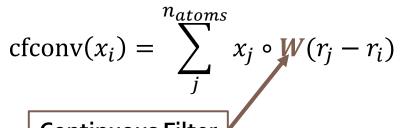
Wei et al. Sci Rep. (2019), 3358

Problem: Atoms are not pixel



Key Innovation: "Continuous Convolutions"







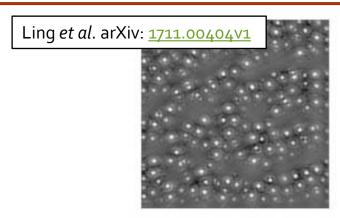


Ref: Schütt et al. JCP, (2018)

Take-home Points

Deep Learning has many advantages...

- Learning from sparse features
- High model complexity -> High accuracy
- Transfer learning



Spheroidite

Major method for molecules: Message Passing Networks

- Automatically learning fingerprints
- Combination of "message," "update" and "readout"
- Many variations in the literature

Practical Exercises: Learn how to implement these in TensorFlow

