



# Gilmer et al. (2017)

## Neural Message Passing for Quantum Chemistry

Andreas Holm (HDF957) & Tim  
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UNIVERSITY OF COPENHAGEN

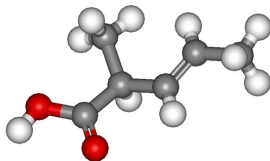


## References (A)

- Main article: *Justin Gilmer, Samuel S. Schoenholz, Patrick F. Riley, Oriol Vinyals, George E. Dahl Neural Message Passing for Quantum Chemistry, International Conference on Machine Learning (2017)*
- Simpler model based on Gilmer: *Bishop and Bishop Deep Learning Foundations and Concepts (2024), chapter 13 Graph Neural Networks*
- QM9 Dataset used by Gilmer: *Raghunathan Ramakrishnan, Pavlo Dral, Matthias Rupp and Anatole Lilienfeld: Quantum chemistry structures and properties of 134 kilo molecules, Scientific Data (2014)*

# The simple model of Bishop (2024) (T)

- Supervised learning, observations are molecules, classified as water soluble/insoluble
- Applying ML to a deterministic rather than stochastic process
- Nodes, attributes  $X$ , edges, adjacency matrix  $A$



From

Chantiers de Science <https://jcmarot.com/2019/02/03/molecules-1-isomeries/>

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$$X = \begin{bmatrix} x_{11}, \dots, x_{1D} \\ \dots \\ x_{N1}, \dots, x_{ND} \end{bmatrix}$$

$$A = \begin{bmatrix} 0, \dots, 1 \\ \dots \\ 1, \dots, 0 \end{bmatrix}$$

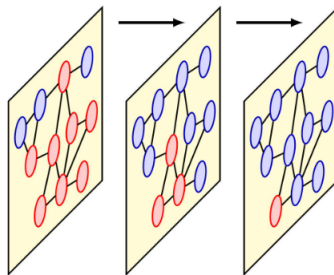
- i.i.d. is out and permutation equivariance is in

## But ... different molecules have different dimensionality? (A)

- Convolution is your friend
- Node (atom)  $x_v$  has a hidden value  $h_v$  that is updated by each layer
- $h_v^{t+1} = \text{Update}(h_v^t, \text{Aggregate}(h_m^t : m \in N(x_v)))$
- The larger the molecule, the deeper the network

13.2. Neural Message-Passing

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Bishop, Figure 13.4

## Gilmer (2017) (T)

- Predict not one but 13 different variables related to molecules
- Two types of known variable values
  - What chemists already know (does not cover all 130.000 molecules)
  - What can be approximated using quantum computing (expensive)
- ML is almost as good as (1) and much less expensive than (2)

No.	Property	Unit	Description
1	tag	—	'gdbn' string to facilitate extraction
2	<i>i</i>	—	Consecutive, 1-based integer identifier
3	<i>A</i>	GHz	Rotational constant
4	<i>B</i>	GHz	Rotational constant
5	<i>C</i>	GHz	Rotational constant
6	$\mu$	D	Dipole moment
7	$\alpha$	$\text{\AA}^3$	Isotropic polarizability
8	$\epsilon_{\text{HOMO}}$	Ha	Energy of HOMO
9	$\epsilon_{\text{LUMO}}$	Ha	Energy of LUMO
10	$\epsilon_{\text{gap}}$	Ha	Gap ( $\epsilon_{\text{LUMO}} - \epsilon_{\text{HOMO}}$ )
11	$\langle R^2 \rangle$	$\text{\AA}^2$	Electronic spatial extent
12	zpve	Ha	Zero point vibrational energy
13	$U_0$	Ha	Internal energy at 0 K
14	$U$	Ha	Internal energy at 298.15 K
15	$H$	Ha	Enthalpy at 298.15 K
16	$G$	Ha	Free energy at 298.15 K
17	$C_v$	$\frac{\text{cal}}{\text{molK}}$	Heat capacity at 298.15 K

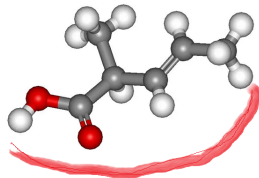
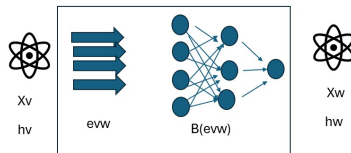
Table 3. Calculated properties. Properties are stored in the order given by the first column.

Ramakrishnan (2014)

## Predict one variable (A)

- Edges become multi-dimensional real-valued vectors
- Message from atom  $x_v$  to  $x_w$ 

$$M(h_v, h_w, e, vw) = B(e_{vw})h_w$$
- Notation suppresses number of layer
- $O(n^2 d^2)$  - but large molecules are rarely cliques
- If the largest molecule in the training data is big, build a deep network and vice versa
- Not robust if test molecules are much larger than those in the training data (no i.i.d.)
- Next: 'attention mechanism' across all atoms



## Gilmer Table 2: Results (T)

	Previous non-graph models					Previous graph models			Gilmer		
	Table 2. Comparison of Previous Approaches (left) with MPNN baselines (middle) and our methods (right)								Graph model	Ensemble	
	Target	BAML	BOB	CM	ECFP4	HDAD	GC	GG-NN	DTNN	enn-s2s	enn-s2s-ens5
Variables	mu	4.34	4.23	4.49	4.82	3.34	0.70	1.22	-	0.30	0.20
	alpha	3.01	2.98	4.33	34.54	1.75	2.27	1.55	-	0.92	0.68
	HOMO	2.20	2.20	3.09	2.89	1.54	1.18	1.17	-	0.99	0.74
	LUMO	2.76	2.74	4.26	3.10	1.96	1.10	1.08	-	0.87	0.65
	gap	3.28	3.41	5.32	3.86	2.49	1.78	1.70	-	1.60	1.23
	R2	3.25	0.80	2.83	90.68	1.35	4.73	3.99	-	0.15	0.14
	ZPVE	3.31	3.40	4.80	241.58	1.91	9.75	2.52	-	1.27	1.10
	U0	1.21	1.43	2.98	85.01	0.58	3.02	0.83	-	0.45	0.33
	U	1.22	1.44	2.99	85.59	0.59	3.16	0.86	-	0.45	0.34
	H	1.22	1.44	2.99	86.21	0.59	3.19	0.81	-	0.39	0.30
	G	1.20	1.42	2.97	78.36	0.59	2.95	0.78	.84 <sup>2</sup>	0.44	0.34
	Cv	1.64	1.83	2.36	30.29	0.88	1.45	1.19	-	0.80	0.62
	Omega	0.27	0.35	1.32	1.47	0.34	0.32	0.53	-	0.19	0.15
	Average	2.17	2.08	3.37	53.97	1.35	2.59	1.36	-	0.68	0.52

- Unit is proportion of the error that chemists make using traditional methods
- ChatGPT was used to mark all errors green if  $< 1$