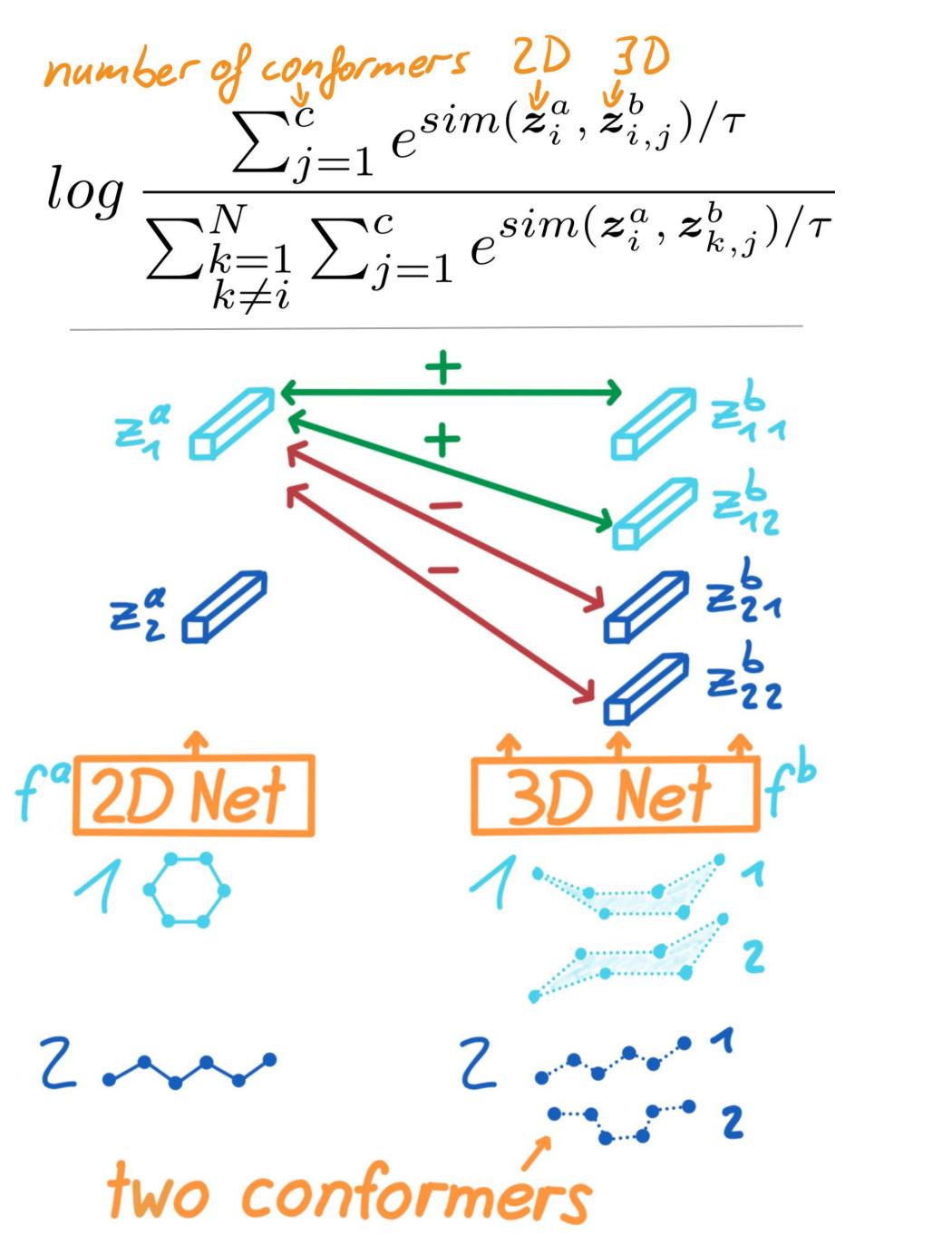
3D Infomax improves GNNs for Molecular Property prediction

Molecular property predictions are better when additionally informed by 3D geometry

However, 3D information is often unavailable. We pre-train a GNN to generate implicit 3D information from 2D inputs which it can use to inform downstream molecular property predictions.

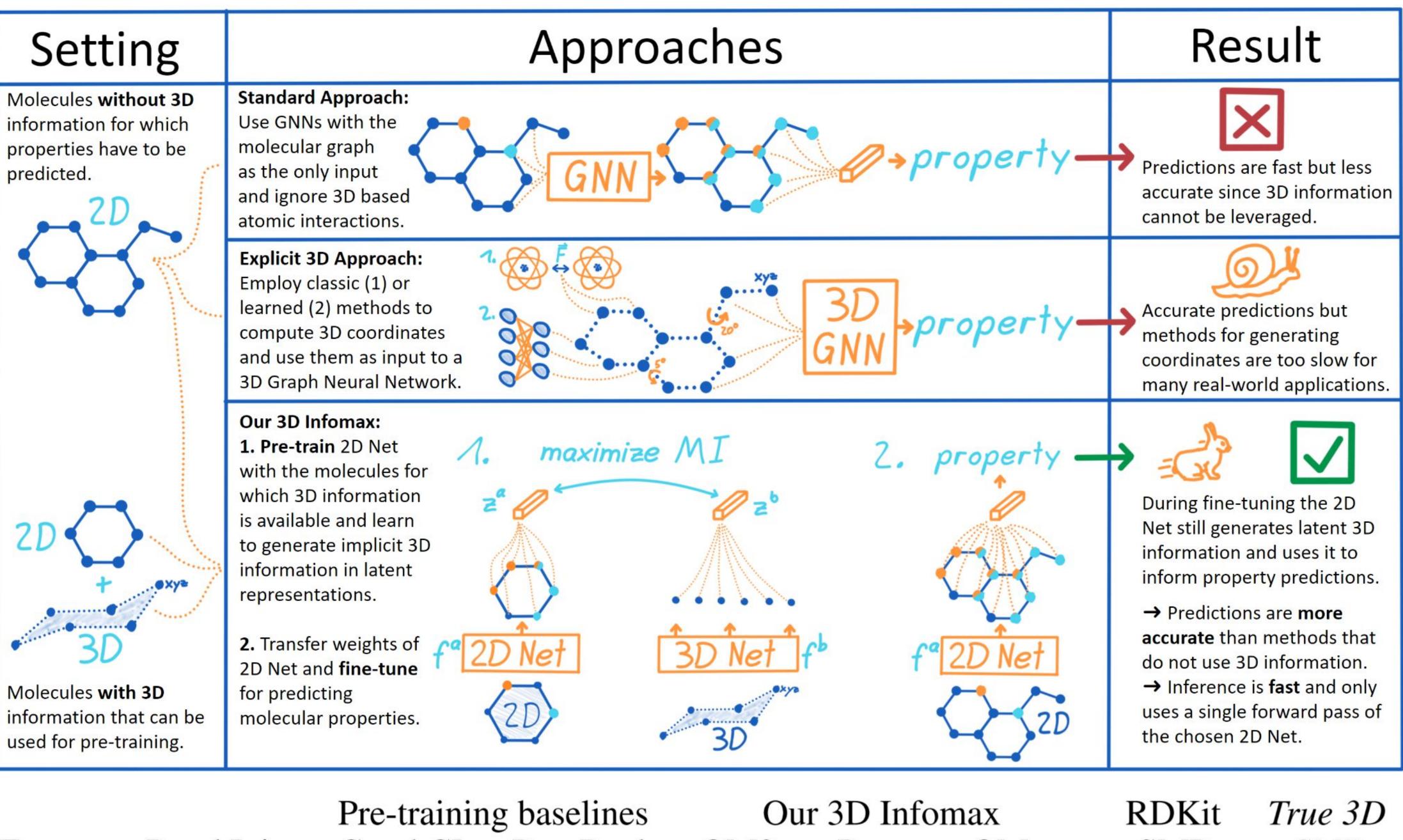
- More accurate than 2D methods but fast
- Highly transferable and robust representations
- Principled SSL: we know that 3D reasoning helps

Multi-conformer contrastive objective:



Hannes Stärk*, Dominique Beaini, Gabriele Corso, Christian Dallago, Prudencio Tossou, Pietro Liò, Stephan Günnemann





		Pre-training baselines		Our 3D Infomax			RDKit	True 3D
Target	Rand Init	GraphCL	PropPred	QM9	Drugs	QMugs	SMP	SMP
μ	$0.4133{\scriptstyle\pm0.003}$	0.3937	0.3975	0.3507	0.3512	0.3668	0.4344	0.0726
lpha	$0.3972{\scriptstyle\pm0.014}$	0.3295	0.3732	0.3268	0.2959	0.2807	0.3020	0.1542
homo	$82.10{\scriptstyle \pm 0.33}$	79.57	93.11	68.96	70.78	70.77	82.51	56.19
lumo	85.72 ± 1.62	80.81	99.84	69.51	71.38	78.10	80.36	43.58
gap	$123.08{\scriptstyle\pm3.98}$	120.08	131.99	101.71	102.59	103.85	114.24	85.10
r2	$22.14{\scriptstyle\pm0.21}$	21.84	29.21	17.39	18.96	18.00	22.63	1.51
ZPVE	$15.08{\scriptstyle\pm2.83}$	12.39	11.17	7.966	9.677	12.06	5.18	2.69
c_v	$0.1670{\scriptstyle \pm 0.004}$	0.1422	0.1795	0.1306	0.1409	0.1208	0.1419	0.0498





