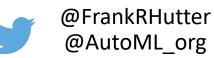


Neural Architecture Search Benchmarks: Past, Present and Future

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Neural architecture search (NAS) is exploding!

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Neural Architecture Search: A Survey

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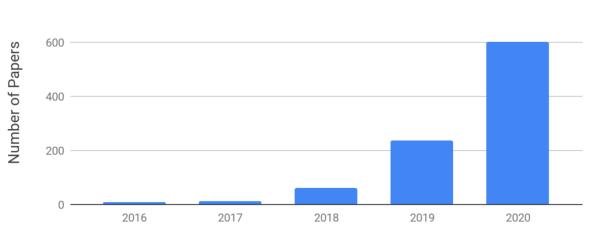
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800

Number of papers on NAS published in conferences, journals and arXiv



Year



The Past

- Scientific best practices in NAS
- First benchmarks
- The Present

• The Future

NAS Had To Overcome Some Childhood Problems 1/2

- Poor performance compared to random search [Li & Talwalkar, 2020, Yu et al, 2020]
- Poor reproducibility [Li & Talwalkar, 2020]
 - Even random seeds are very important
- Training pipeline matters much more than architecture [Yang et al, 2020]
- Poor scientific practices [Lindauer & Hutter, 2020]
 - Inavailability of code

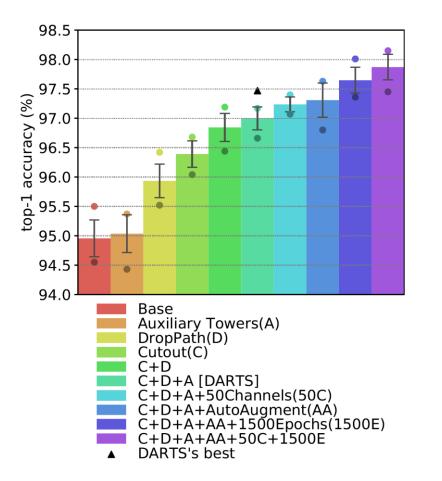
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- Incomparable training code, search spaces, evaluation schemes, etc

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NAS Childhood Problems 2/2

Training pipeline matters much more than architecture



Model	Params	Test error (%)
DenseNet-BC (Huang 1, 2017)	25	3.46
PyramidNet (Han et al., 2007)	JM	3.31
Shake-Shake + c/o (DeVries Saylor, 2017)	26.2M	2.56
PyramidNet + SD (Yamada et a. 918)	26.0M	2.31
ENAS + c/o (Pham et al., 2018)	4.6M	2.89
DARTS + c/o (Liu et al., 2018c)	3.4M	2.83
NASNet-A + c/o (Zoph et al., 20^{10}	27.6M	2.40
PathLevel EAS + c/o (Cai et a^{1} $J18b$)	14.3M	2.30
AmoebaNet-B + c/o (Real, $, 2018$)	9M	2.13
Proxyless-R + c/o (ours)		2.30
Proxyless-G + c/o (out	5.7	2.08

Incomparable

- Different training code (often unavailable)
- Different search spaces
- Different evaluation schemes

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NAS Best Practices Checklist

1. Releasing code

• Not just trained architectures

2. Properly comparing methods

 Proper scientific evaluations, powered by tabular/surrogate benchmarks for statistical significance

3. Reporting important details

• E.g., hyperparameter tuning

Suggestion to reviewers

- Deemphasize final results table on CIFAR-10 (or other datasets),
 - be aware of many confounding factors

[Lindauer and Hutter, JMLR 2020]

The NAS Best Practices Checklist (version 1.0, September 6, 2019) *by Marius Lindauer and Frank Hutter*

Best practices for releasing code

For all experiments you report, check if you released:

- $\hfill\square$ Code for the training pipeline used to evaluate the final architectures
- $\hfill\square$ Code for the search space
- □ The hyperparameters used for the final evaluation pipeline, as well as random seeds
- □ Code for your NAS method
- □ Hyperparameters for your NAS method, as well as random seeds

Note that the easiest way to satisfy the first three of these is to use *existing* NAS benchmarks, rather than changing them or introducing new ones.

Best practices for comparing NAS methods

- □ For all NAS methods you compare, did you use exactly the same NAS benchmark, including the same *dataset* (with the same training-test split), *search space* and *code* for training the architectures and *hyperparameters* for that code?
- □ Did you control for confounding factors (different hardware, versions of DL libraries, different runtimes for the different methods)?
- \Box Did you run ablation studies?
- $\hfill\square$ Did you use the same evaluation protocol for the methods being compared?
- □ Did you compare performance over time?
- □ Did you compare to random search?
- □ Did you perform multiple runs of your experiments and report seeds?
- \Box Did you use tabular or surrogate benchmarks for in-depth evaluations?

Best practices for reporting important details

- □ Did you report how you tuned hyperparameters, and what time and resources this required?
- □ Did you report the time for the entire end-to-end NAS method (rather than, e.g., only for the search phase)?
- □ Did you report all the details of your experimental setup?



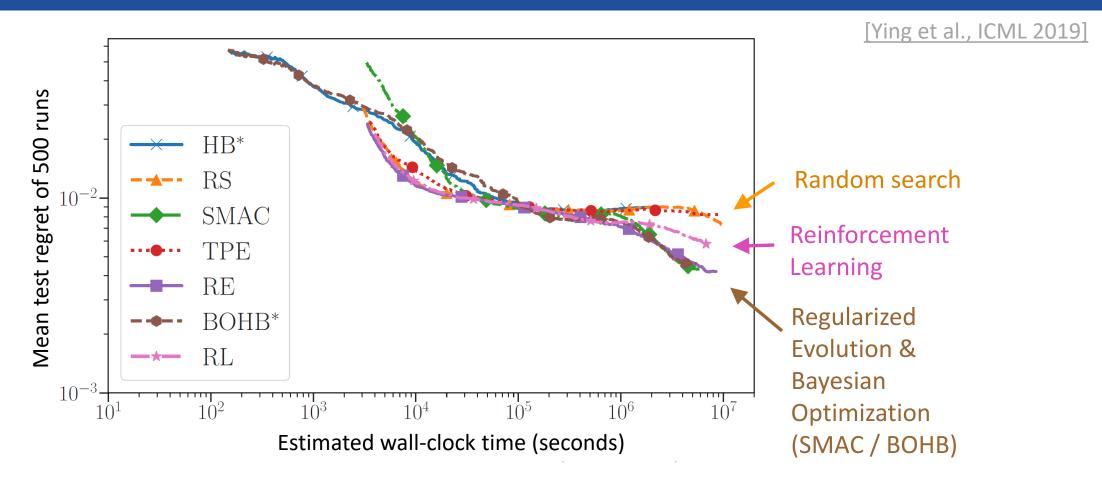
NAS-Bench-101: The First Tabular NAS Benchmark

[Ying et al., ICML 2019]

- Small cell search space that we exhaustively evaluated
 - Enables evaluating a NAS method in minutes on a laptop
 - Enables proper scientific research: multiple runs, robustness studies, etc
 - Fair apples-to-apples evaluations by design (fixed final evaluation pipeline)
 - Of course, source code and scripts are available
- 423k architectures evaluated on CIFAR-10
 - Nobody has to ever run this again
 - Only possible with Google resources (4.000 TPUs for months)
 - One-time cost already far more than amortized

NAS-Bench-101: Comparison of Optimizers

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- Note: SMAC (published 2011) outperforms RL (published 2016)
- Tabular NAS benchmarks finally allow us to do these analyses



- The Past
 - Scientific best practices in NAS
 - First benchmarks
- The Present
 - Surrogate Benchmarks
- The Future

Tabular NAS Benchmarks Really Caught on 🙂

- UNI FREIBURG NAS-Bench-101 [Ying et al, ICML 2019] & NAS-Bench-1Shot1 [Zela et al, ICLR 2020]
 - Up to 423k unique architectures
 - NAS-Bench-201 [Dong & Yang, ICLR 2020]
 - 6466 unique architectures
 - Extension: **NATS-Bench** with 32768 unique architectures
 - NAS-Bench-ASR
 - 8242 unique architectures
 - NAS-Bench-NLP
 - 14322 architectures evaluated
 - But these NAS benchmarks are too small to be realistic 😕
 - E.g., local search is state of the art for such small space, but performs poorly on large ones, such as DARTS [White et al, AutoML 2020]
 - More realistically-sized search spaces
 - E.g., DARTS search space has $\approx 10^{18}$ architectures
 - E.g., FBNet search space is $\approx 10^{21}$ architectures

Surrogates: Going Beyond the Limits of Tabular NAS Benchmarks

Problem

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- For any realistically-sized search space, there is no hope to evaluate it exhaustively to compute a table

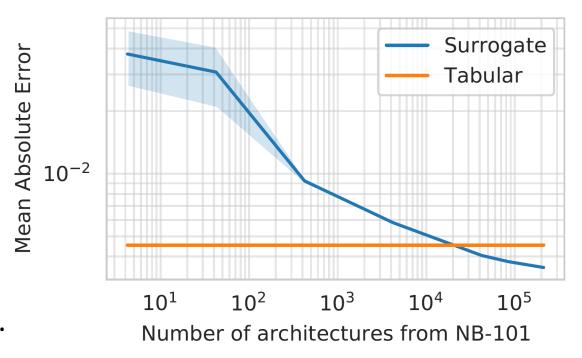
• NAS Surrogate Benchmark Methodology:

- Evaluate a subset of architectures
- Fit a model to those; use model predictions in lieu of the real/tabular benchmark
- Many previous works already used a surrogate model to predict the performance of untested architectures
 - All works on Bayesian optimization (SMAC, BOHB, NAS-BOWL, BANANAS, ...)
 - All works on "predictor-based NAS" (NPE-NAS, BRP-NAS, etc)
- The difference is in how we use the model:
 - not to speed up search, but to define a benchmark
 - Search algorithms only have a blackbox interface to the surrogate benchmark, just like for a tabular benchmark
 - Any improvements in surrogate modelling will improve surrogate NAS benchmarks



- The evaluations in a table come with a certain error due to the noise of SGD
 - Many NAS benchmarks quantify this error with 3-5 repetitions for (some subset of) the architectures
- From a machine learning perspective
 - A tabular NAS benchmark predicts a noisy function f(A) by evaluating at
 A a few times and returning the mean
 - This makes an independence assumption, not using data for similar architectures
 - A good model should do better than that ...
 - And indeed, it does \bigcirc

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Surr-NAS-Bench-DARTS (SNB-DARTS, aka NAS-Bench-301)

[Siems et al, arXiv 2021]

- Evaluated 50.000 architectures in the DARTS search space using different optimizers
- Evaluated broad range of regression models to fit this data
- Best regression models
 - Gradient boosting (XGB/LGB)
 - Graph convolutional networks
- Estimation errors lower than error due to noise in a single run of SGD

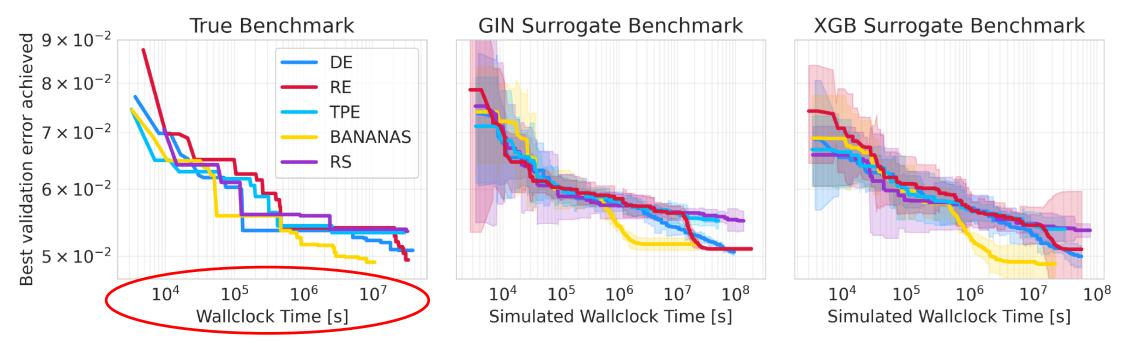
Optimizer			# Evaluations
Discrete		RS	24047
	Evolution	DE	7275
	Evolution	RE	4639
		TPE	6741
	BO	BANANAS	2243
		COMBO	745
One-Shot		DARTS	2053
		GDAS	234
		RANDOM-WS	198
		PC-DARTS	149

Model	Te	Test		
1110401	R^2	sKT		
LGBoost	0.892	0.816		
XGBoost	0.832	0.817		
GIN	0.832	0.778		
NGBoost	0.810	0.759		
μ -SVR	0.709	0.677		
MLP (Path enc.)	0.704	0.697		
RF	0.679	0.683		
ϵ -SVR	0.675	0.660		

Benchmarking NAS Methods on SNB-DARTS

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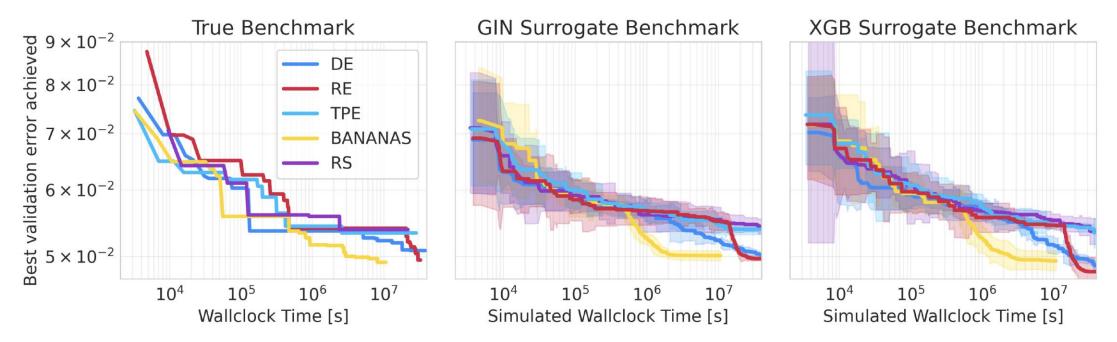
[Siems et al, arXiv 2021]



- Actual wallclock time required when run sequentially: > 1 GPU year, per run
- Surrogate benchmark: many orders of magnitude faster
- Note: performance is smoother on the surrogates, since we could only afford 1 run on the true benchmark so far

Ablation: Only Using Random Samples to Generate SNB-DARTS

[Siems et al, arXiv 2021]



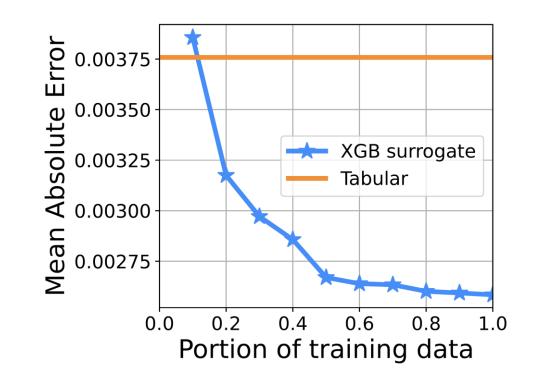
- Randomly-gathered training data suffices
 - At least to obtain truthful performance trajectories
 - Predictive performance for top-performing architectures a bit weaker
- Advantage:

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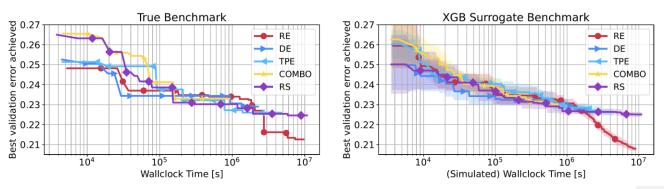
No bias possible towards the optimizers used to generate the training data



- Evaluated 25.000 random architectures in the FBNet search space
- Surrogate model: XGBoost
- Again, estimation errors lower than error due to noise in a single run of SGD



• Again, truthful trajectory plots



[Siems et al, arXiv 2021]



- The Past
 - Scientific best practices in NAS
 - First benchmarks
- The Present
 - Surrogate Benchmarks
 - Many new benchmarks
- 中 The Future



- Goal: Discover Entirely New Architectures with NAS
 - All the important architectures in deep learning were found manually
 - I hope that this will change over the next years
- We need:
 - Reliable & Efficient NAS Methods
 - Robust zero-cost proxies
 - Robust one-shot models
 - Efficient blackbox methods
 - Powerful search spaces
 - E.g., hierarchical spaces
 - NAS methods that are compatible with arbitrary search spaces



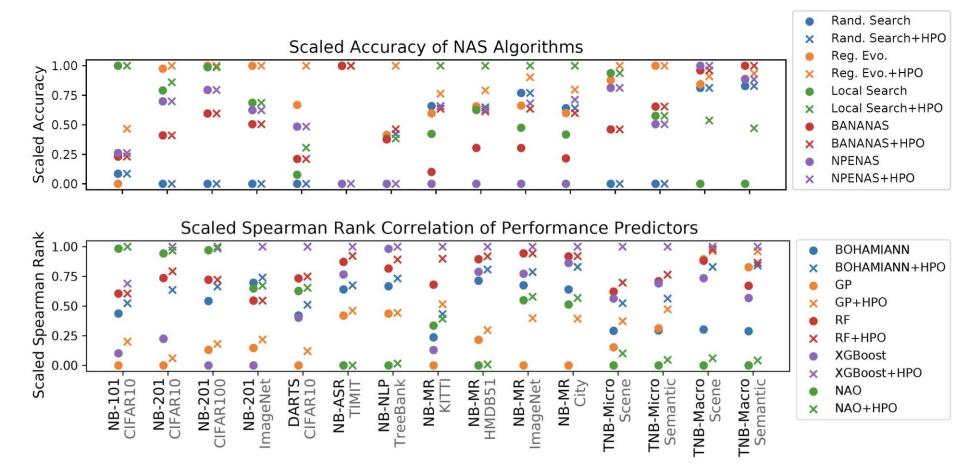
• Problem: Existing NAS Benchmarks are very different

- E.g., NAS-Bench-101 has the operations in the nodes while NAS-Bench-201 has them in the edges
- As a result, NAS Algorithms often hardcoded the search space in their code
- NASlib helps to unify the interface to different NAS benchmarks
- This allows access to 25 (!) different NAS benchmarks
 - For the cost of a single implementation

	Size	Queryable						
Benchmark		Tab.	Surr.	LCs	Macro	Туре	#Tasks	NAS-Bench-Suite
NAS-Bench-101	423k	1				Image class.	1	✓
NAS-Bench-201	6k	1		1		Image class.	3	✓
NAS-Bench-NLP	10^{53}			1		NLP	1	✓
NAS-Bench-1Shot1	364k	1				Image class.	1	✓
NAS-Bench-301	10^{18}		1			Image class.	1	✓
NAS-Bench-ASR	8k	1			1	ASR	1	✓
NAS-Bench-MR	10^{23}		1		1	Var. CV	4	✓
TransNAS-Bench	7k	1		1	1	Var. CV	14	✓
NAS-Bench-111	423k		1	1		Image class.	1	✓
NAS-Bench-311	10^{18}		1	1		Image class.	1	✓
NAS-Bench-NLP11	10^{53}		1	1		NLP	1	✓

Do we really need so many different NAS benchmarks?

- Do weSadly, yes
 - Conclusions on "just" NAS-Bench-101 & 3 NAS-Bench-201 datasets can be misleading



 Tuning NAS algorithm hyperparameters on one benchmark can lead to poor performance on others



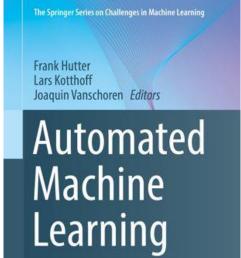
- There are tons of tabular NAS benchmarks by now
 - These enable scientific evaluations with minimal compute (i.e., carbon emissions)
- Surrogates are the path to realistic search spaces
 - They can even model performance more truthfully than tabular benchmarks
- NAS-Bench-Suite has 25 queryable NAS benchmarks
 - Available through a unified interface in NASLib (https://github.com/automl/NASLib)

• More information: <u>http://automl.org</u>

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Book on AutoML: <u>http://automl.org/book</u>





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I'm looking for additional great postdocs!

