S+pKa Predictor of Ionization Constants – Recent Progress and Results

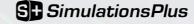
Robert Fraczkiewicz, PhD

Research Fellow



Please note: this presentation, including questions from the audience, is being recorded and may be made available.





A bit of history

- Until 2012 the S+pKa model was exclusively trained on ~11,000 compounds from published literature. This model will be labeled as "v 6.0".
- In 2012 Bayer Pharma AG had shared with us an additional set of ~16,000 compounds with measured pK_a. The resulting "v 7.0" model was trained on combined data and its prediction results were published in 2015.

CHEMICAL INFORMATION

AND MODELING

pubs.acs.org/jcim

Best of Both Worlds: Combining Pharma Data and State of the Art Modeling Technology To Improve in Silico pK_a Prediction

Robert Fraczkiewicz,**,† Mario Lobell,**,‡ Andreas H. Göller,‡ Ursula Krenz,‡ Rolf Schoenneis,‡ Robert D. Clark,† and Alexander Hillisch‡





[†]Simulations Plus, Inc. 42505 10th Street West, Lancaster, California 93534, United States

[‡]Global Drug Discovery, Bayer Pharma AG, Wuppertal, Germany

A bit of history

• S+pKa "v 7.0" has shown dramatic improvements in prediction quality as evaluated in the *Bayer chemical space*. All test sets were external.

Table 1. Performance Statistics of Two Versions of the S+pKa model: One Trained on Public Set Only (marked "v 6.0") and the Other on the Combined Public and Industrial Sets (marked "v 7.0")^a

					M	AE	RM	SE		\mathbb{R}^2
test set	number of compounds	$\begin{array}{c} \text{number of p} K_{\text{a}} \\ \text{values} \end{array}$	average closest Tanimoto similarity to the Industrial Set	fraction of Tanimoto similars (score ≥0.80)	v 6.0	v 7.0	v 6.0	v 7.0	v 6.0	v 7.0
1	4730	5644	0.88	98%	0.82	0.41	1.03	0.58	0.85	0.95
2	8931	9168	0.82	60%	0.79	0.52	1.04	0.71	0.76	0.89
3	12,951	16,404	0.79	45%	0.72	0.50	0.94	0.67	0.87	0.93

[&]quot;External Test Sets 1, 2, and 3 have been described in the Data Sets section. Predictive statistics: MAE = mean absolute error, RMSE = root mean square error, and R^2 = determination coefficient.





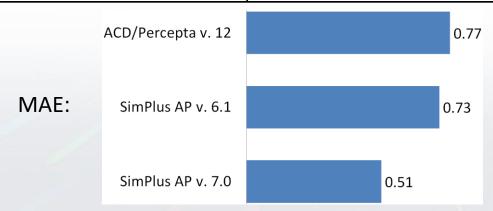
A bit of history

It outperformed competition, too.

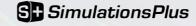
ACD/Percepta v. 12 and ADMET Predictor™ v 6.1 show comparable pK_a prediction accuracy ADMET Predictor™ v 7.0 (after retraining with BTr) shows significantly improved pK_a prediction accuracy

Prediction statistics for 981-compound Bayer test set with 981 exp. pK_a values (subset of newest measurements on 12951 Bayer compounds):

Predicted by	Trained with	MAE	RMSE	R ²
ACD/Percepta v 12	15932 lit pK _a	0.77	1.05	0.84
ADMET Predictor v 6.1	14147 lit pK _a	0.73	0.95	0.86
ADMET Predictor v 7.0	14149 lit pK _a + 19467 Bayer pK _a	0.51	0.67	0.93







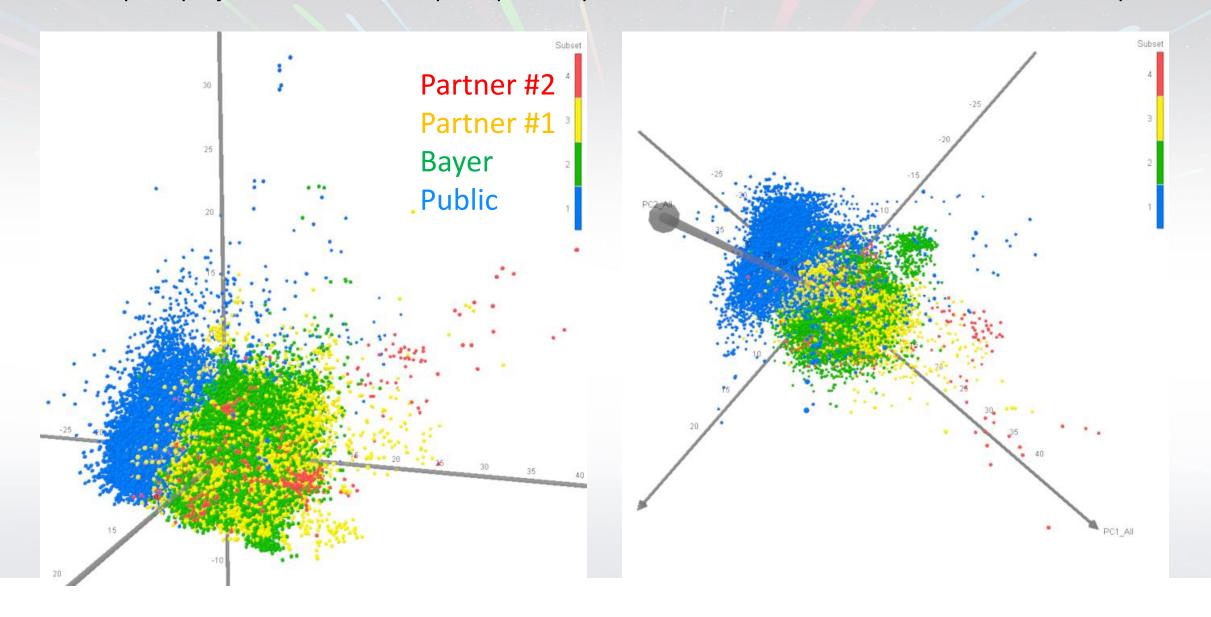
2022

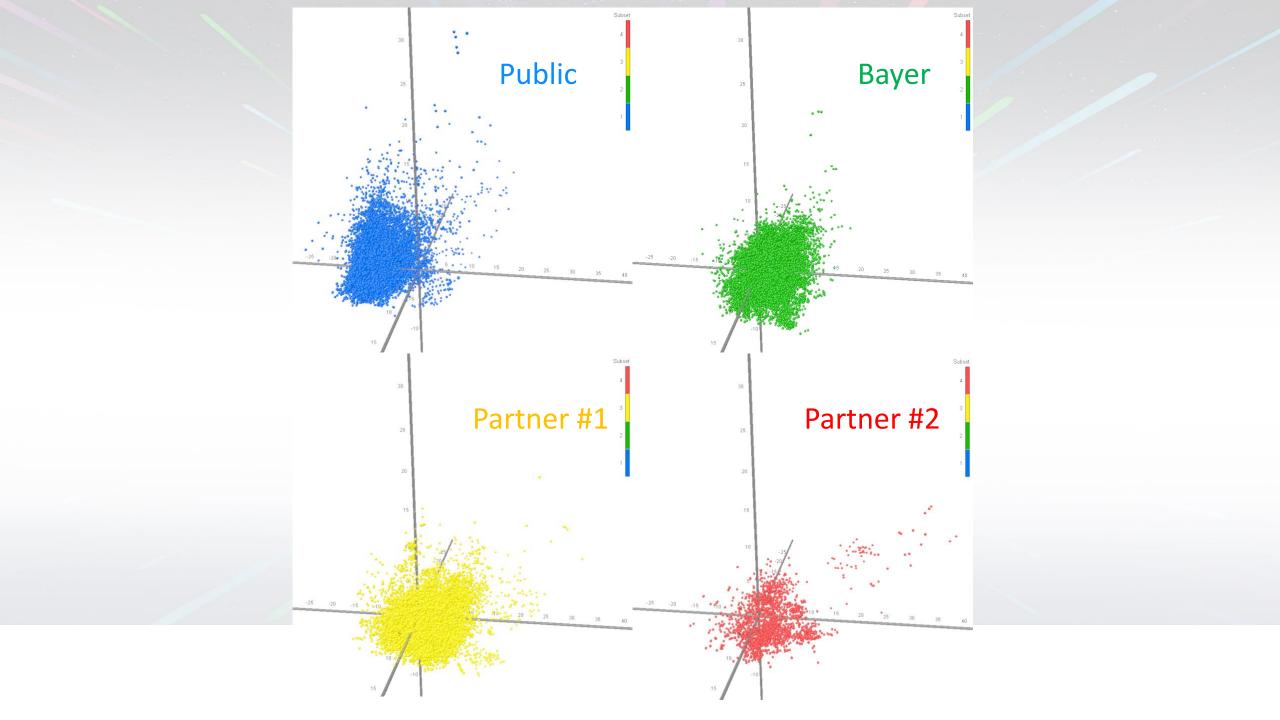
- Two new industrial partners (large pharmaceutical companies; further labeled as "Partner #1" and "Partner #2") have indicated inadequate coverage of their chemical space by the "v 7.0" S+pKa.
- Instead of complaining both partners have shared with us significant amount of experimental pK_a data extracted from their corporate databases.
 - Partner #1 has provided ~19,000 compounds
 - Partner #2 has provided ~2,400 compounds
- From August 2022 until November 2022 we were busy rebuilding the S+pKa model with the new data appended to public+Bayer set. The resulting newest version carries the "v 10.5" label.





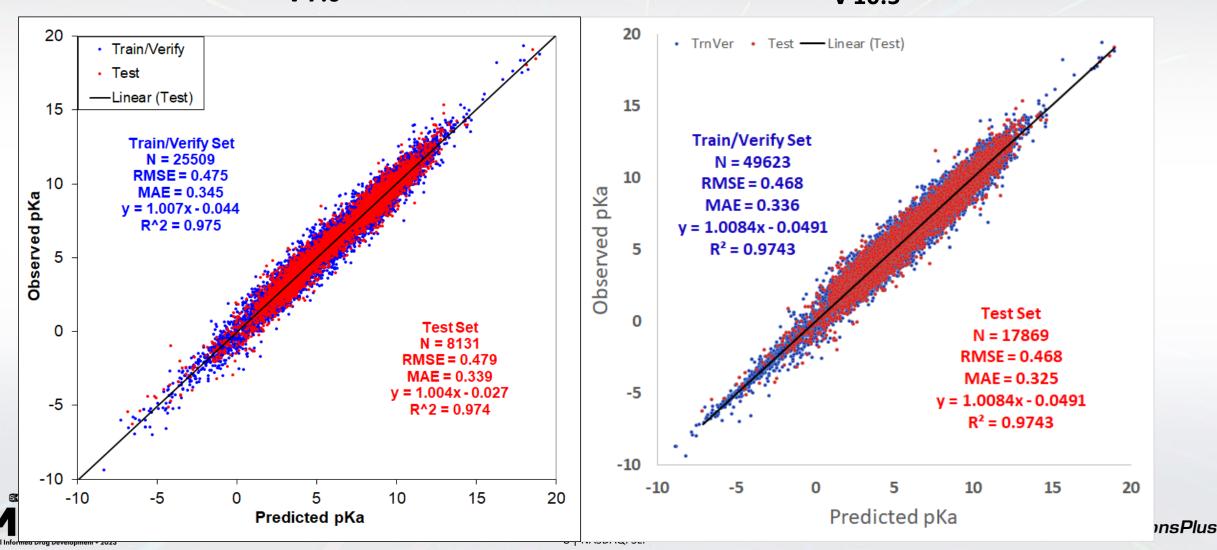
Chemical space projected on the first 3 principal components of the ADMET Predictor molecular descriptors matrix





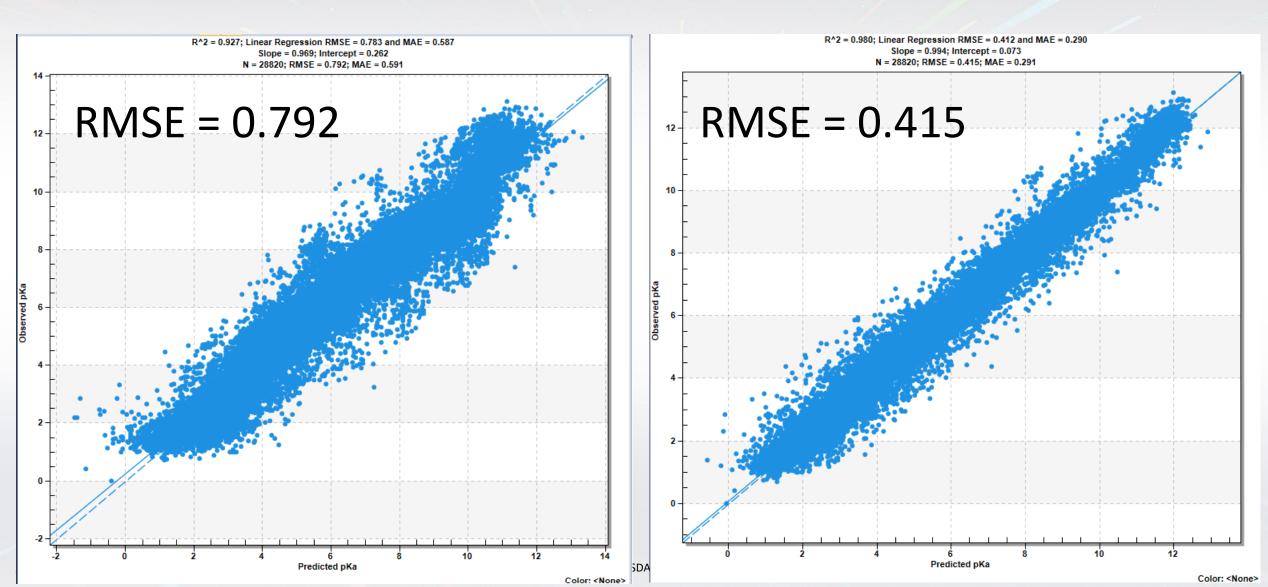
"v 7.0" vs. "v 10.5" performance

It's apples vs. oranges, but the relative improvement is welcome
 v 7.0



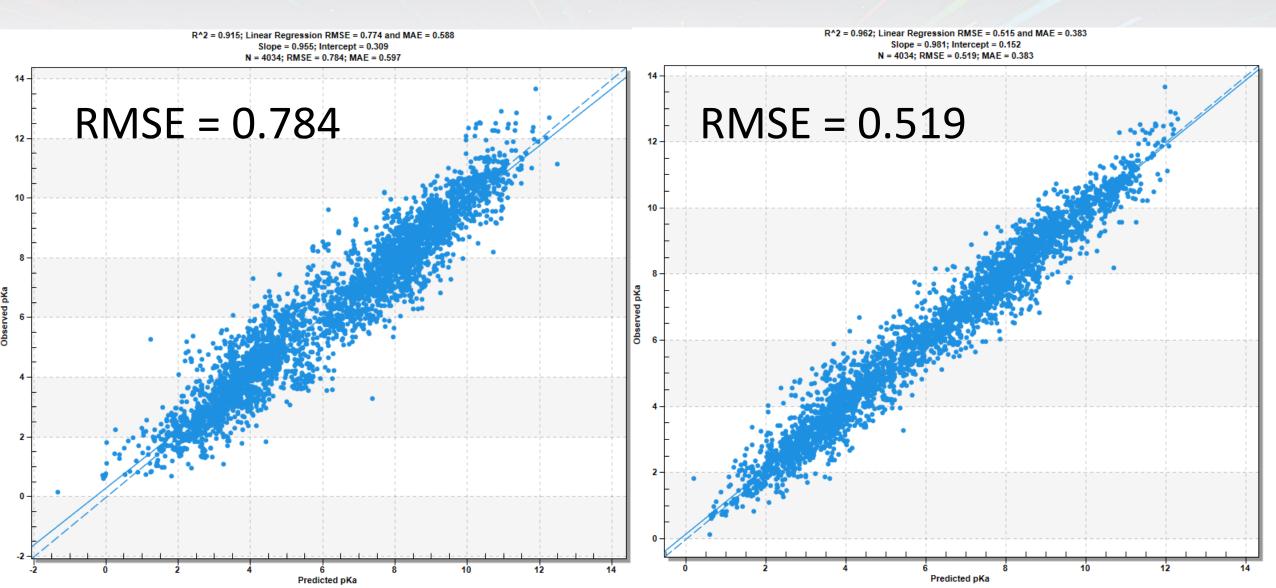
"v 7.0" and "v 10.5" models vs. received data

• Data from Partner #1. "v 7.0" RMSE = 0.792, "v 10.5" RMSE = 0.415



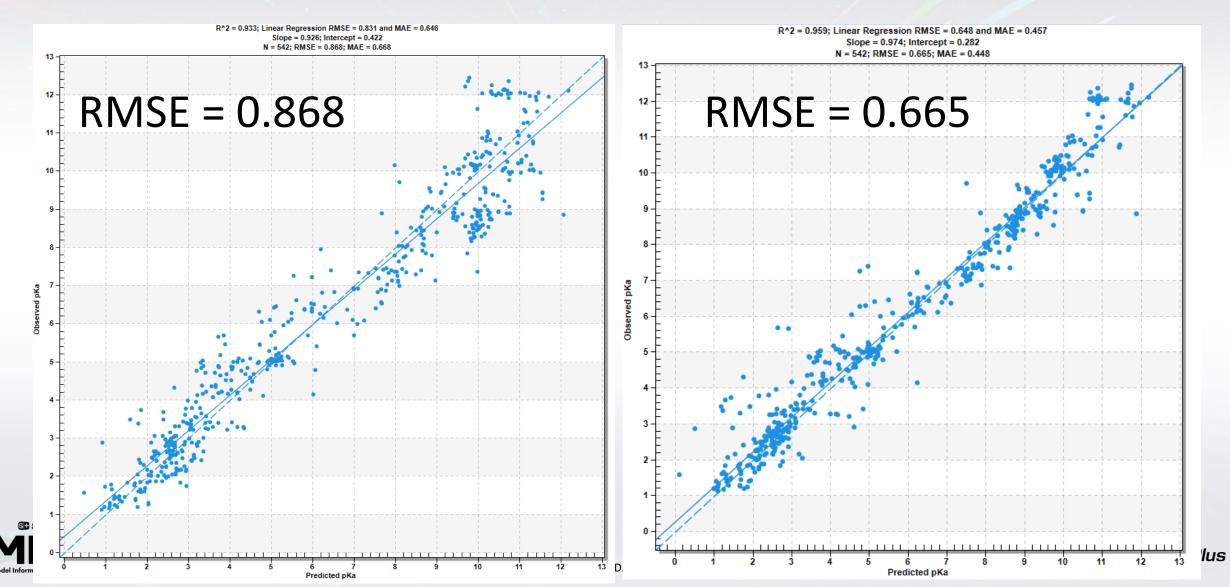
"v 7.0" and "v 10.5" models vs. received data

Data from Partner #2. "v 7.0" RMSE = 0.784, "v 10.5" RMSE = 0.519



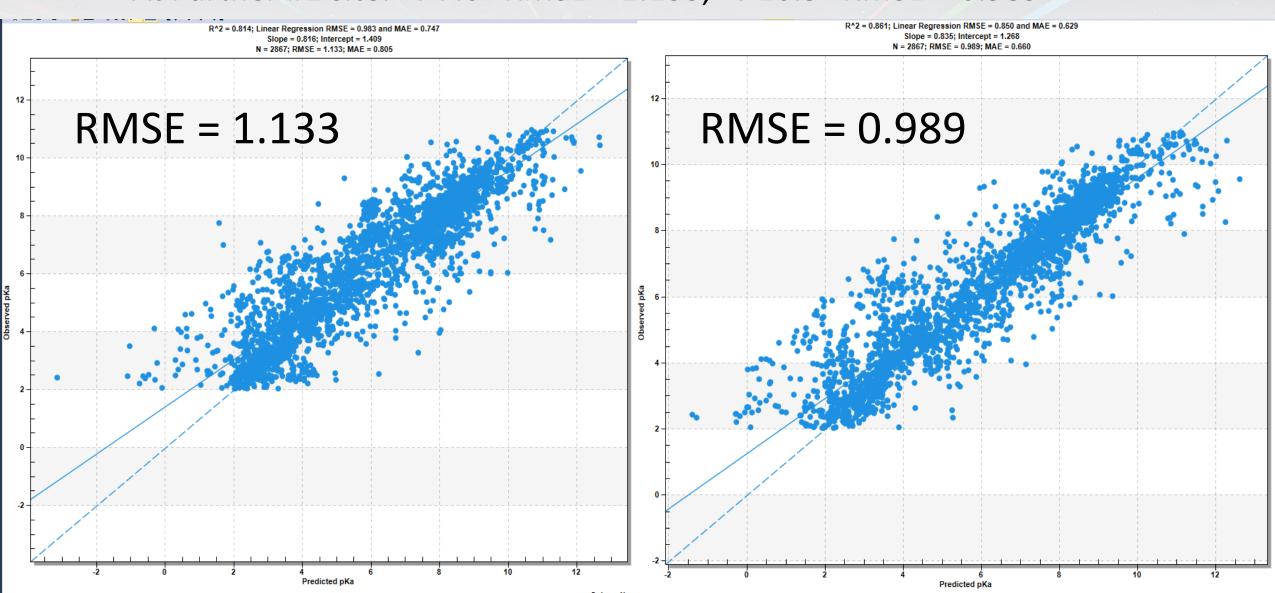
"v 7.0" and "v 10.5" models in external testing

• At Partner #1 site. "v 7.0" RMSE = 0.868, "v 10.5" RMSE = 0.665



"v 7.0" and "v 10.5" models in external testing

At Partner #2 site. "v 7.0" RMSE = 1.133, "v 10.5" RMSE = 0.989



Conclusions

- The chemical space covered by the new S+pKa model has been significantly expanded.
- Prediction accuracy has been improved.
- Partner #1 was very much forthcoming with a data set representative of their chemical space and reaped sizable rewards. Moreover, "changing input tautomer for some biggest outliers improved predictions".
- Partner #2 delivered much less data and the set's place in their chemical space remains uknown.
- From our side questions were raised regarding validity of some spectrophotometric measurements. We are awaiting answers from both partners.







And now a new message from the Little Bird:

In January 2023 we have received thousands of pK_a data points from a new industrial Partner #3.

Stay tuned...