

Errata: Feature Selection in Taxonomies with Applications to Paleontology

Unfortunately the tables in the published version contain incorrect numbers for the baseline models FAMILY, GENUS, and SPECIES. The corrected tables can be found here.

Table 1: Results for the complete taxonomies with PALEO_10_10 and PALEO_5_5. *Columns:* “size” is the number of nodes of the discovered antichain; “calls” is the number of calls to the linear regressor in the model construction phase; “corr” corresponds to the the correlation coefficient of the linear regression when using the corresponding antichain (in the test data); “RMSE” reports the root mean squared error of the linear regression when using the corresponding antichain (in the test data). *Rows:* FAMILY, GENUS and SPECIES select a fix level of the taxonomy as the antichain. Random reports the mean values over 100 antichains.

	PALEO_10_10				PALEO_5_5			
	size	calls	corr	RMSE	size	calls	corr	RMSE
FAMILY	50	-	0.80	2.90	64	-	0.74	4.07
GENUS	238	-	0.59	5.67	374	-	0.47	9.32
SPECIES	428	-	0.88	2.38	868	-	0.75	4.93
Greedy	246	46534	0.67	4.89	464	140488	0.33	12.89
Top-down	145	1817	0.82	3.09	197	2619	0.79	3.91
Bottom-up	375	3737	0.86	2.65	794	10409	0.75	4.91
Tax Min-cut	201	726	0.51	6.62	306	1329	0.77	4.55
Random	325	-	0.83	3.04	615	-	0.1	150

Table 2: Results for the *Carnivora* and *Rodentia* subtrees of the taxonomy with PALEO_5_5 and PALEO_2_2. Random reports the mean values over 100 antichains.

	PALEO_5_5				PALEO_2_2			
	<i>Carnivora</i>		<i>Rodentia</i>		<i>Carnivora</i>		<i>Rodentia</i>	
	size	calls	corr	RMSE	size	calls	corr	RMSE
FAMILY	13	-	0.45	5.36	11	-	0.50	5.21
GENUS	62	-	0.44	5.42	122	-	0.78	3.86
SPECIES	113	-	0.31	6.72	345	-	0.03	102.2
Greedy	67	3238	0.45	5.42	163	14017	0.73	4.40
Top-down	25	99	0.46	5.34	151	1176	0.74	4.20
Bottom-up	48	578	0.45	5.38	232	2050	0.65	5.45
Tax Min-cut	33	188	0.47	5.30	106	476	0.78	3.77
Random	84	-	0.41	5.16	231	-	0.67	4.74
					size	calls	corr	RMSE
					190	2338	0.40	5.58
					500	4222	0.35	5.89
					234	861	0.42	5.55
					178	-	0.24	7.36
					178	-	0.52	5.34