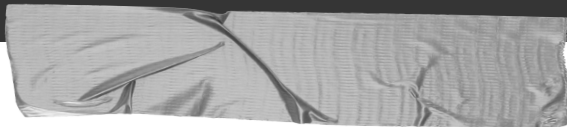

Istraživanje skupa podataka “Anuran Calls”

Stefanija Marković 306/2018

Osnovno o skupu “Anuran Calls”



- Kreiran segmentacijom 60 audio zapisa zvukova žaba.
- Žabe su iz četiri različite porodice, osam rodova i deset vrsti.
- Nakon segmentacije dobijeno je 7195 slogova.
- Za svaki slog izračunato je 22 kepstralnih koeficijenata mel skale (MFCC).

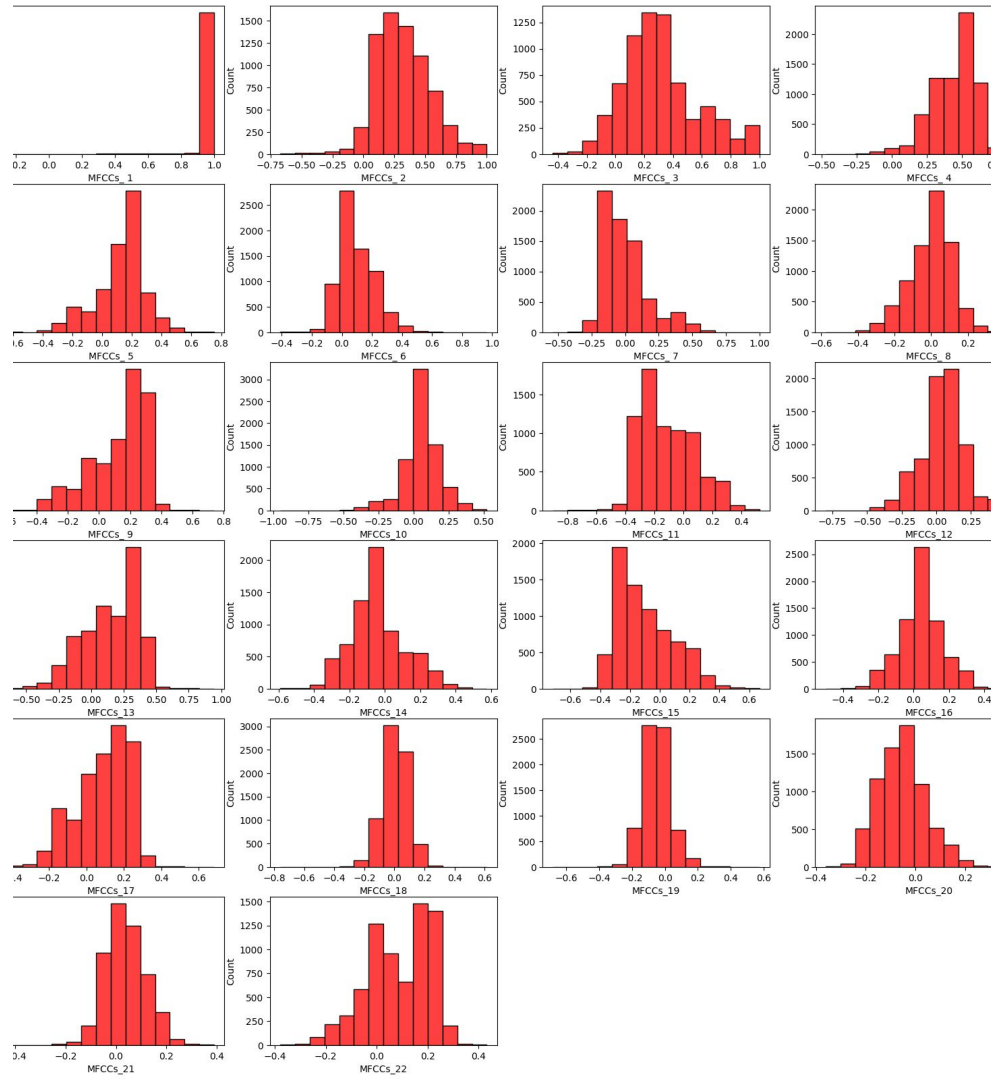
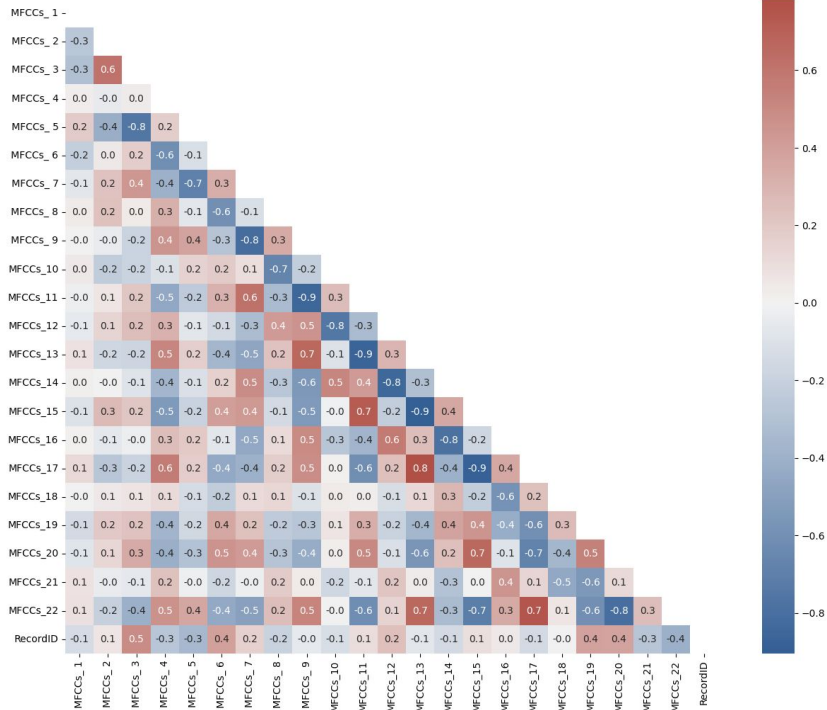
Eksplorativna analiza podataka

	MFCCs_ 1	MFCCs_ 2	MFCCs_ 3	MFCCs_ 4	MFCCs_ 5	MFCCs_ 6	MFCCs_ 7	MFCCs_ 8	MFCCs_ 9	MFCCs_10	...	MFCCs_17	MFCCs_18	MFCCs_19	MFCCs_20	MFCCs_21	MFCCs_22	Family	Genus	Species	RecordID
0	1.0	0.152936	-0.105586	0.200722	0.317201	0.260764	0.100945	-0.150063	-0.171128	0.124676	...	-0.108351	-0.077623	-0.009568	0.057684	0.118680	0.014038	Leptodactylidae	Adenomera	AdenomeraAndre	1
1	1.0	0.171534	-0.098975	0.268425	0.338672	0.268353	0.060835	-0.222475	-0.207693	0.170883	...	-0.090974	-0.056510	-0.035303	0.020140	0.082263	0.029056	Leptodactylidae	Adenomera	AdenomeraAndre	1
2	1.0	0.152317	-0.082973	0.287128	0.276014	0.189867	0.008714	-0.242234	-0.219153	0.232538	...	-0.050691	-0.023590	-0.066722	-0.025083	0.099108	0.077162	Leptodactylidae	Adenomera	AdenomeraAndre	1
3	1.0	0.224392	0.118985	0.329432	0.372088	0.361005	0.015501	-0.194347	-0.098181	0.270375	...	-0.136009	-0.177037	-0.130498	-0.054766	-0.018691	0.023954	Leptodactylidae	Adenomera	AdenomeraAndre	1
4	1.0	0.087817	-0.068345	0.306967	0.330923	0.249144	0.006884	-0.265423	-0.172700	0.266434	...	-0.048885	-0.053074	-0.088550	-0.031346	0.108610	0.079244	Leptodactylidae	Adenomera	AdenomeraAndre	1

5 rows × 26 columns

	MFCCs_ 1	MFCCs_ 2	MFCCs_ 3	MFCCs_ 4	MFCCs_ 5	MFCCs_ 6	MFCCs_ 7	MFCCs_ 8	MFCCs_ 9	MFCCs_10	...	MFCCs_14	MFCCs_15	MFCCs_16	MFCCs_17	MFCCs_18	MFCCs_19	MFCCs_20	MFCCs_21	MFCCs_22
count	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	...	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000	7195.000000
mean	0.989885	0.323584	0.311224	0.445997	0.127046	0.097939	-0.001397	-0.000370	0.128213	0.055998	...	-0.039244	-0.101748	0.042062	0.088680	0.007755	-0.049474	-0.053244	0.037313	0.087567
std	0.069016	0.218653	0.263527	0.160328	0.162722	0.120412	0.171404	0.116302	0.179008	0.127099	...	0.152515	0.187618	0.119915	0.138055	0.084733	0.082546	0.094181	0.079470	0.123442
min	-0.251179	-0.673025	-0.436028	-0.472676	-0.636012	-0.410417	-0.538982	-0.576506	-0.587313	-0.952266	...	-0.590380	-0.717156	-0.498675	-0.421480	-0.759322	-0.680745	-0.361649	-0.430812	-0.379304
25%	1.000000	0.165945	0.138445	0.336737	0.051717	0.012581	-0.125737	-0.063109	0.004648	-0.001132	...	-0.132980	-0.255929	-0.019549	-0.001764	-0.042122	-0.106079	-0.120971	-0.017620	0.000533
50%	1.000000	0.302184	0.274626	0.481463	0.161361	0.072079	-0.052630	0.013265	0.189317	0.063478	...	-0.050715	-0.143259	0.041081	0.112769	0.011820	-0.052626	-0.055180	0.031274	0.105373
75%	1.000000	0.466566	0.430695	0.559861	0.222592	0.175957	0.085580	0.075108	0.265395	0.117725	...	0.039157	0.017348	0.107046	0.201932	0.061889	0.006321	0.001342	0.089619	0.194819
max	1.000000	1.000000	1.000000	1.000000	0.752246	0.964240	1.000000	0.551762	0.738033	0.522768	...	0.575749	0.668924	0.670700	0.681157	0.614064	0.574209	0.467831	0.389797	0.432207

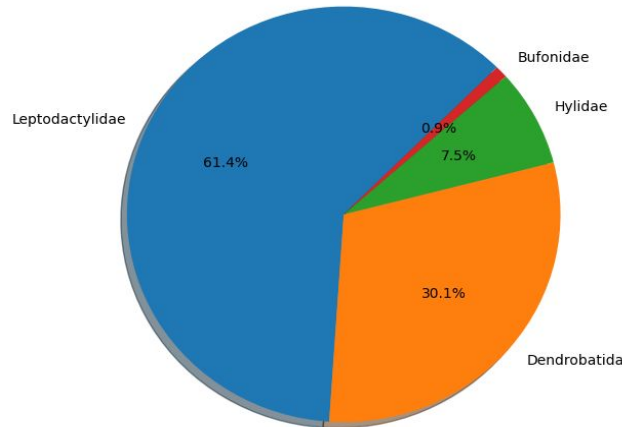
Eksplorativna analiza podataka



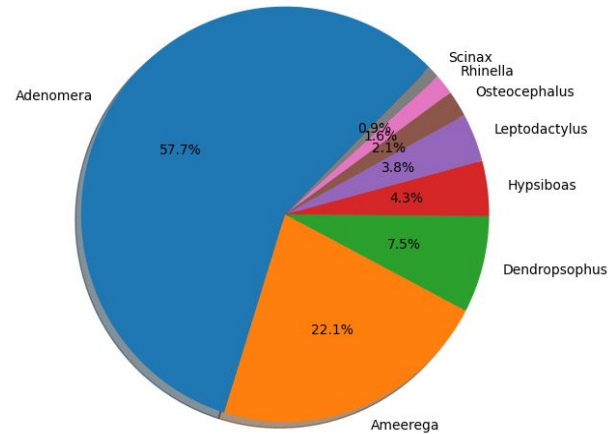
Eksplorativna analiza skupa podataka

Raspoređenost jedinki

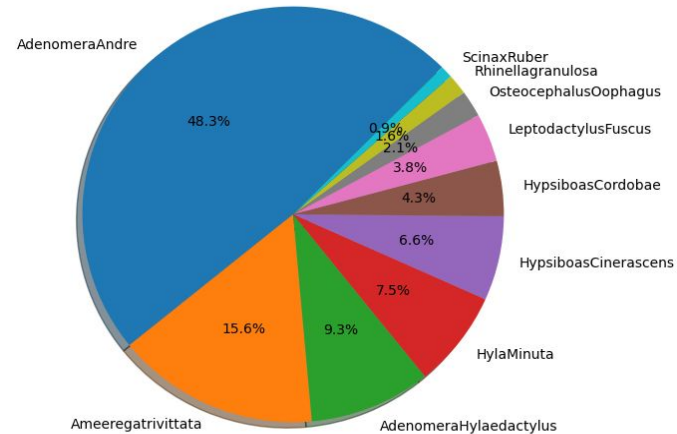
Family Distribution



Genus Distribution

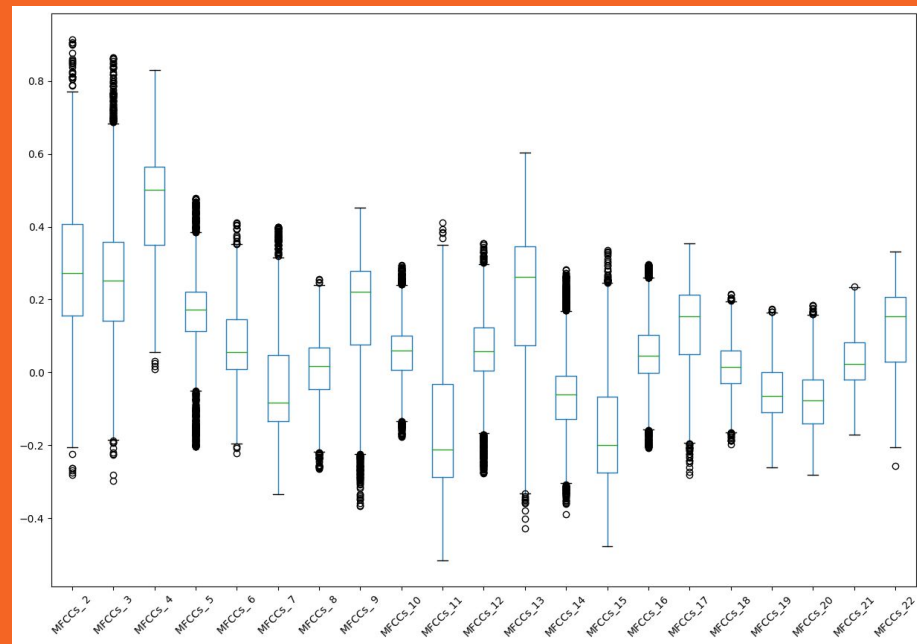
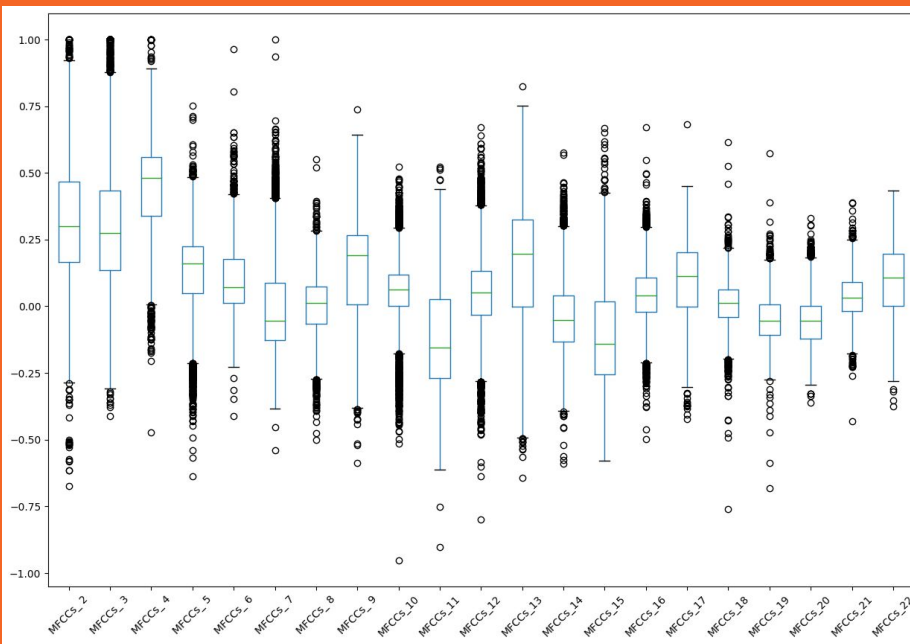


Species Distribution

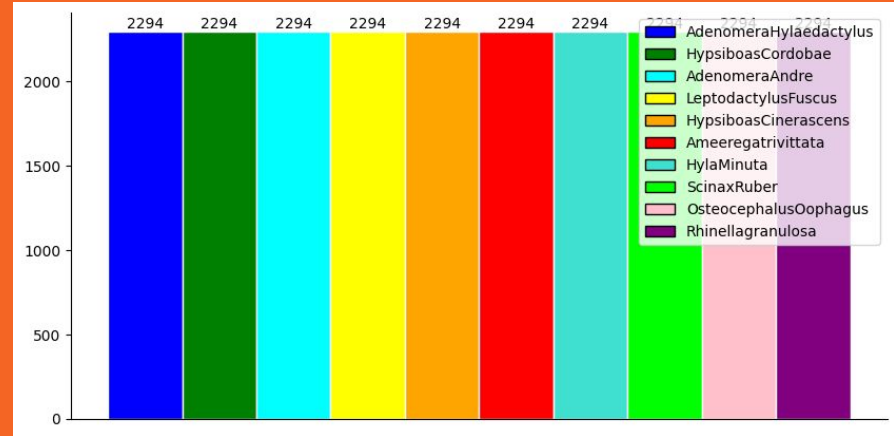
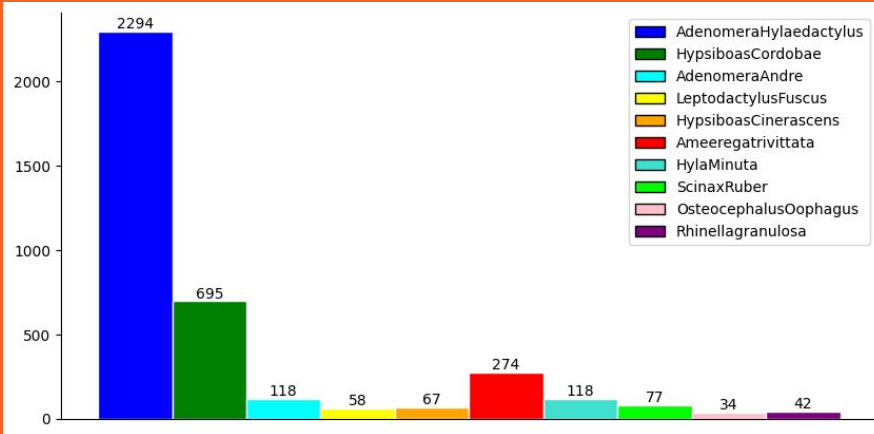
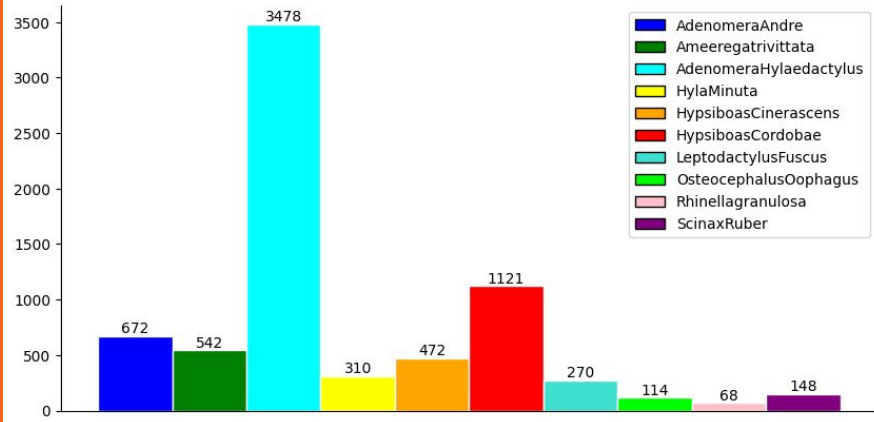


Klasifikacija

Preprocesiranje - rad sa elementima van granica



Balansiranje



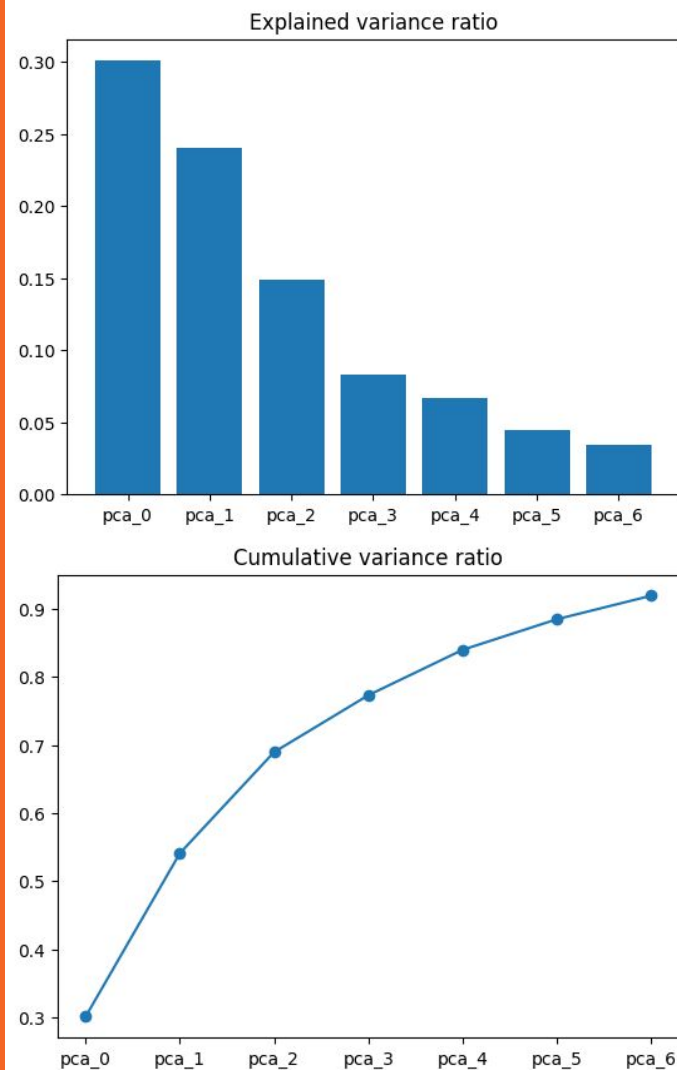


Redukcija dimenzionalnosti

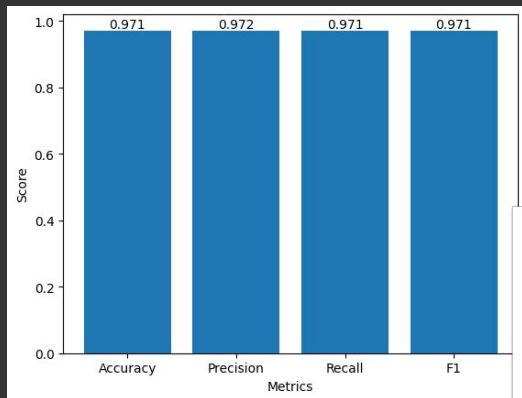
Eliminirani su atributi ***Species***, ***Family***, ***Genus***, **RecordID** i **MFCC_1**.

Primenjena je tehnika **PCA** (Principal component analysis).

Na prvom grafiku je predstavljeno koliko varijanse svaki atribut opisuje, dok je na drugom predstavljena njihova kumulativna suma.

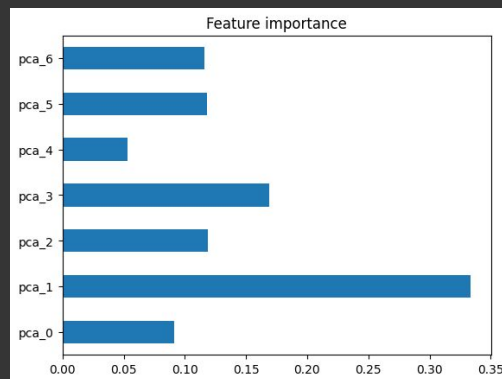
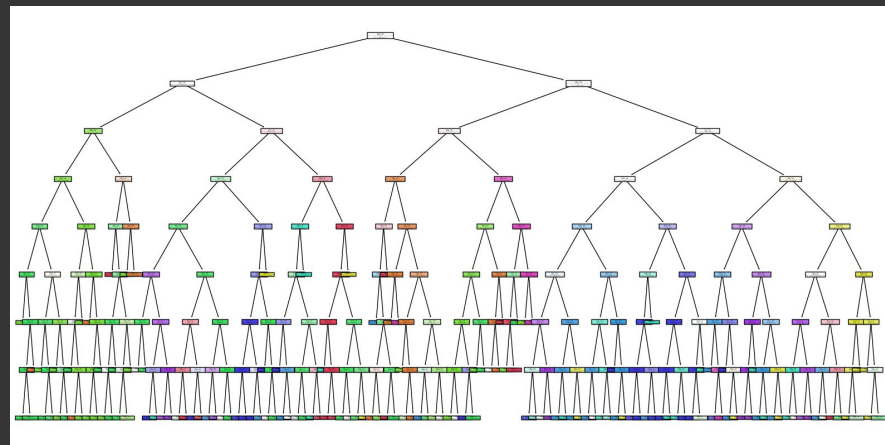
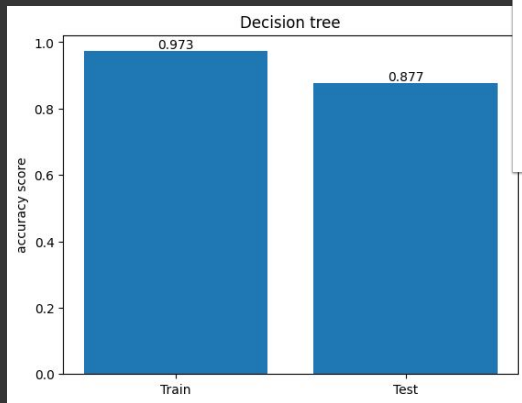


Stablo odlučivanja



Gornja slika
pokazuje rezultate
različnih **mera**.

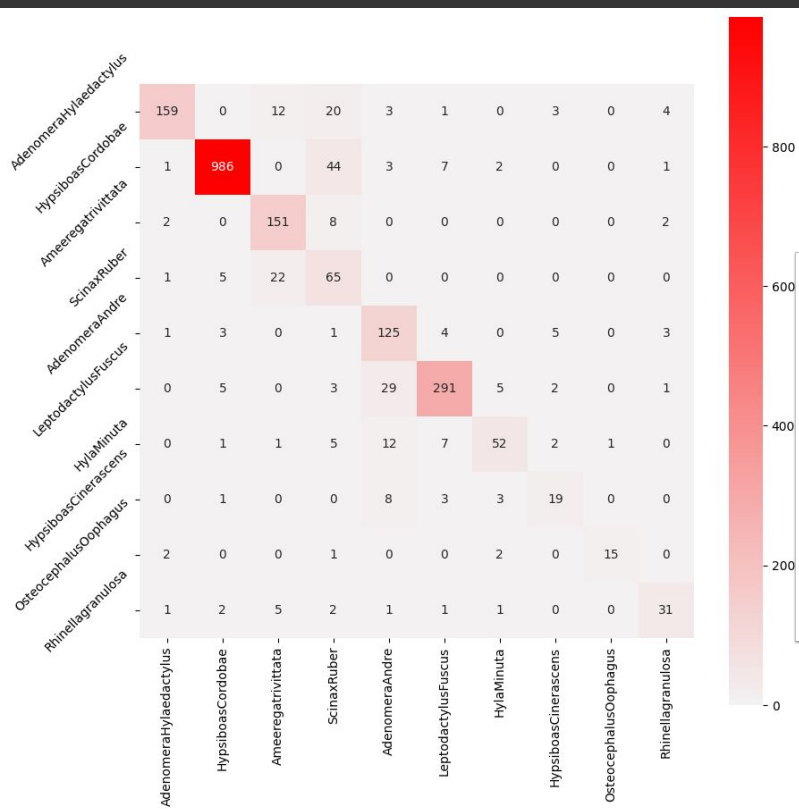
Donja slika
pokazuje **tačnost**
na trening i test
skupu.



Gore je prikazano
kreirano **stablo**
odlučivanja.

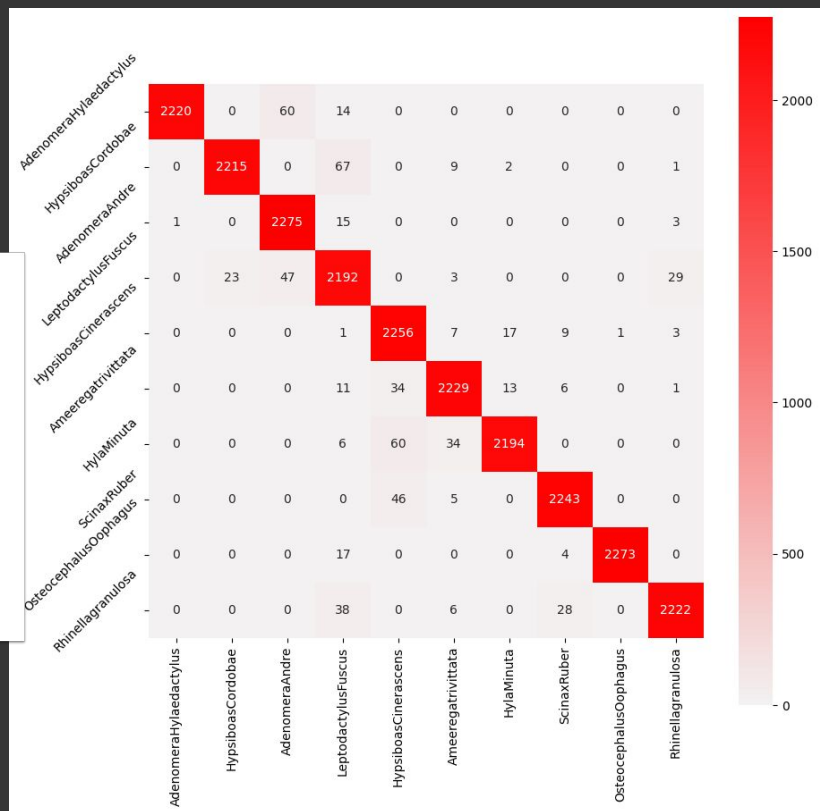
Levo vidimo
važnost atributa
pri odlučivanju.

Stablo odlučivanja

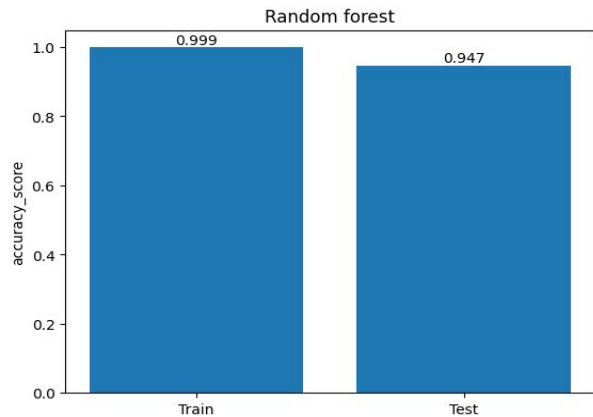


Matrica konfuzije

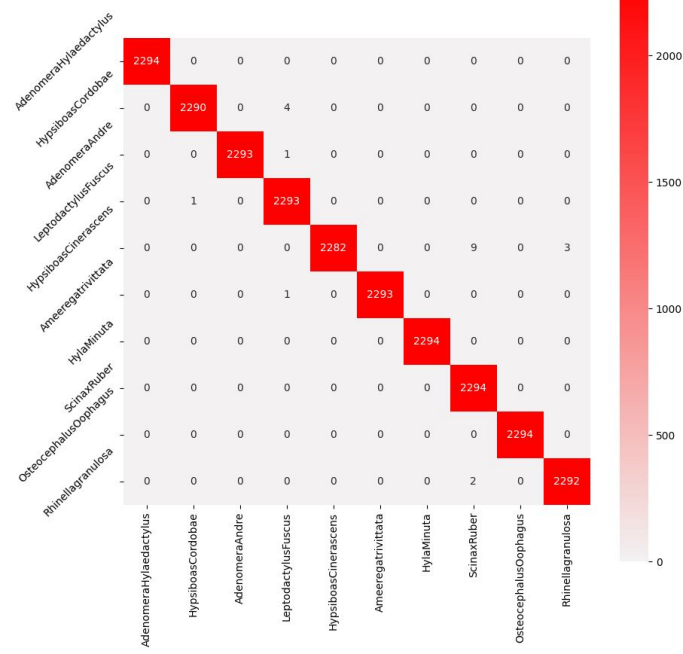
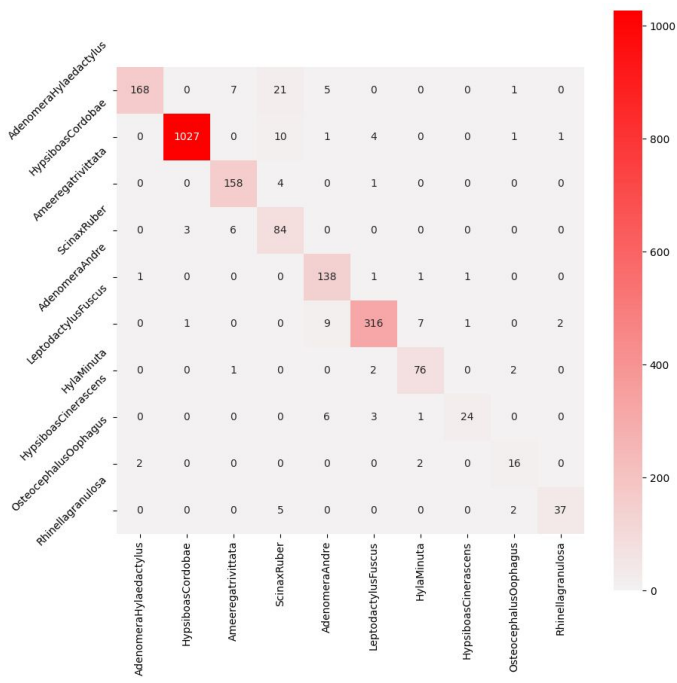
Leva se odnosi na **test** skup, a desna na **trening** skup.



Slučajna šuma



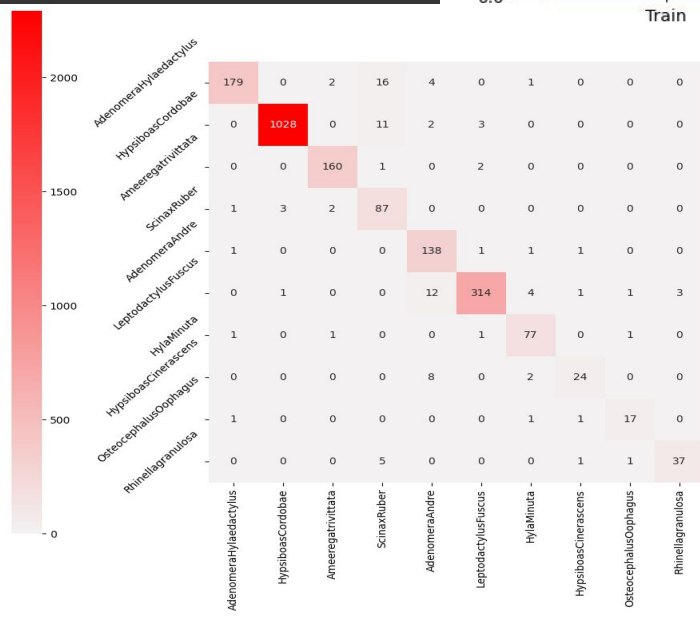
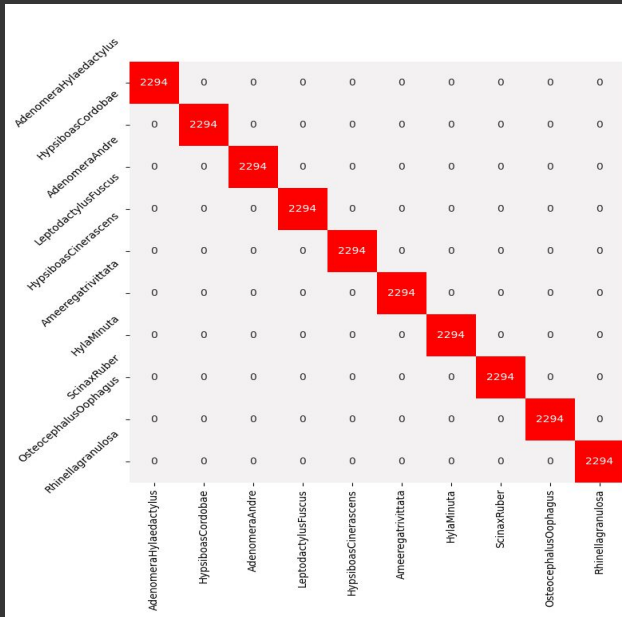
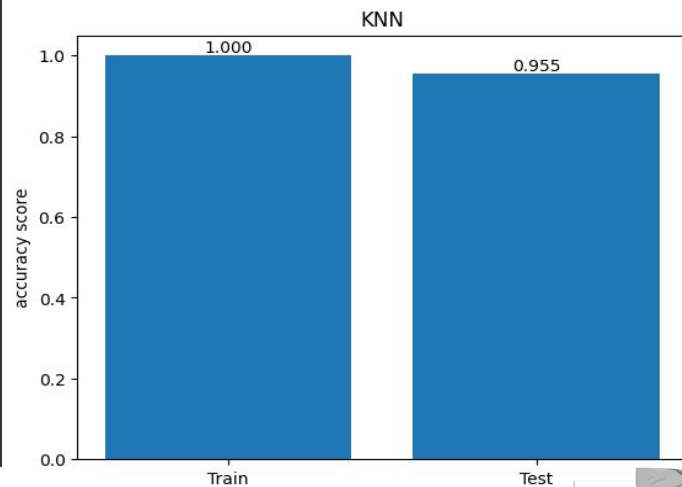
Tačnost
slučajne šume
na trening i
test skupu.



Matrica konfuzije

Leva se odnosi na
test skup, a desna
na **trening** skup.

K najbližih suseda



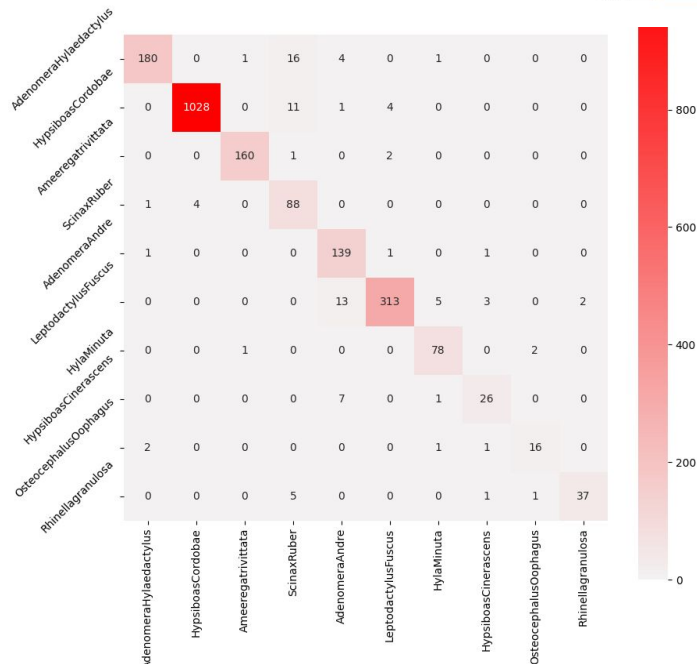
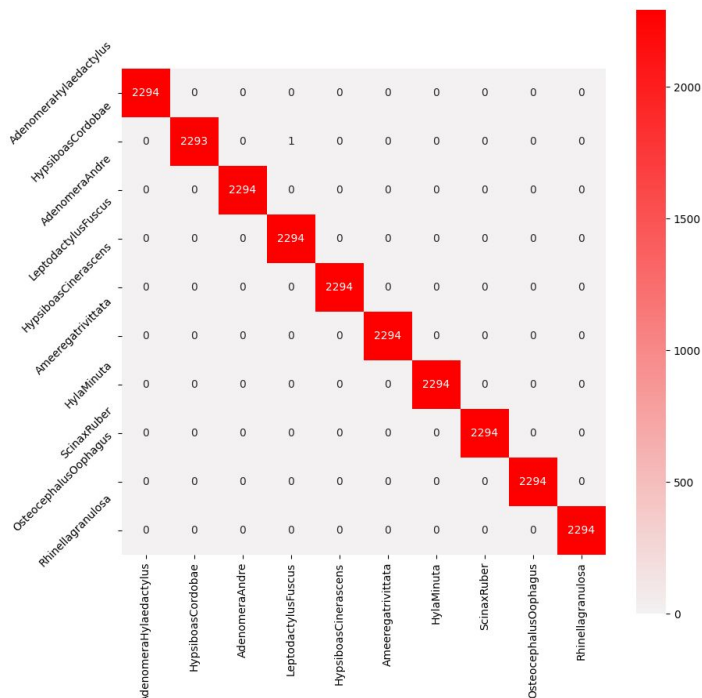
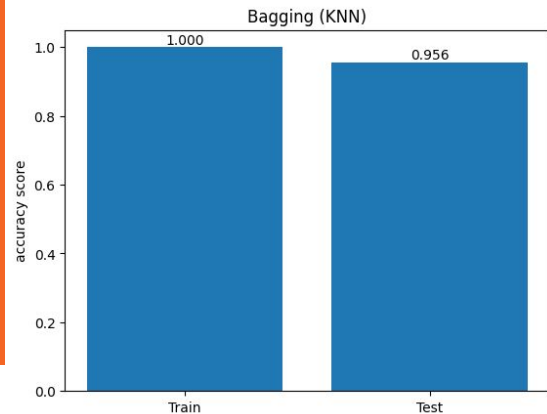
Gornja slika predstavlja **tačnost** na trening i test skupovima.

Matrica konfuzije

Leva se odnosi na **trening** skup, a desna na **test** skup.

Pakovanje

Tačnost
ansambla na
trening i test
skupu.

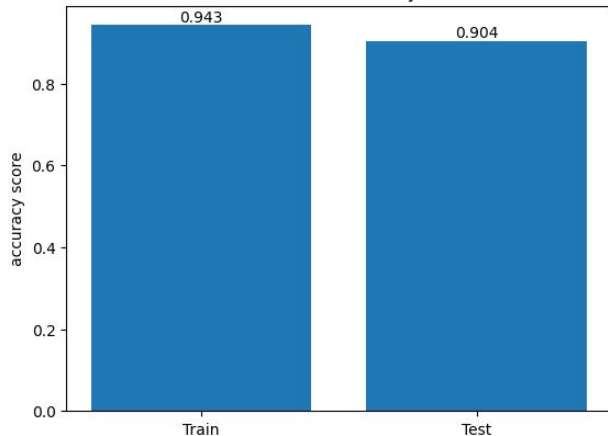


Matrica konfuzije

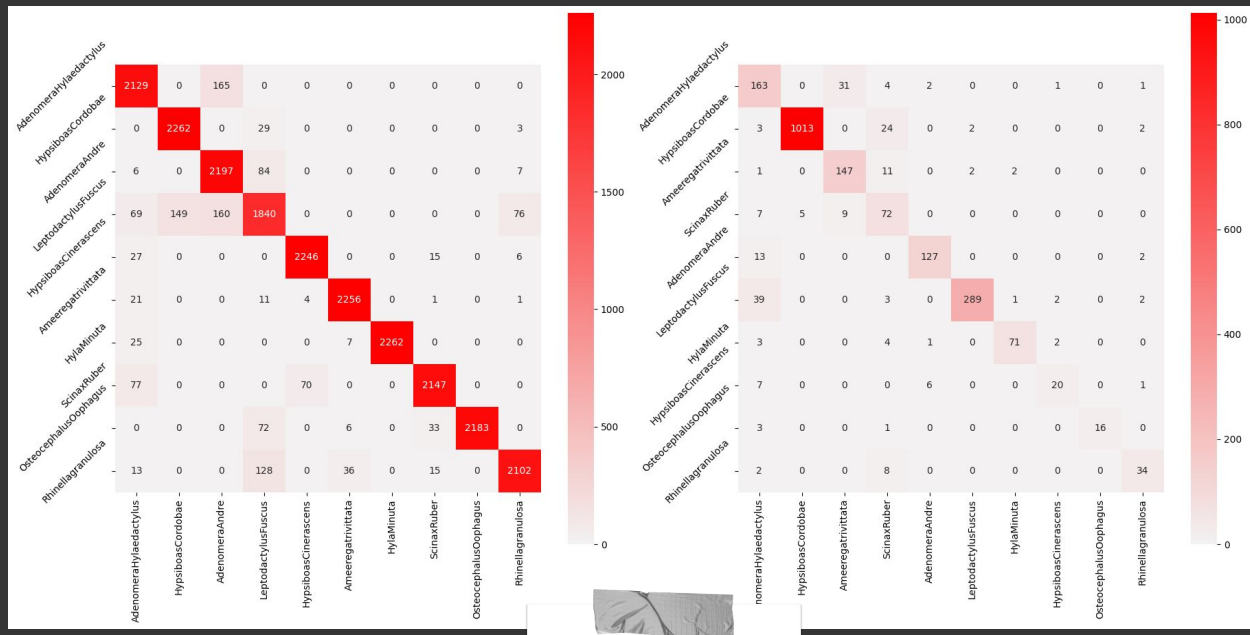
Leva se odnosi na
trening skup, a
desna na **test**
skup.

Naivni Bajes

Gaussian naive Bayes



Gornja slika predstavlja **tačnost** na trening i test skupovima.

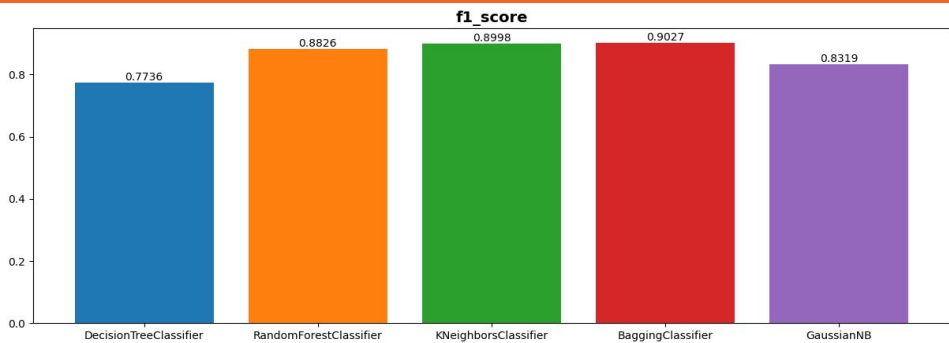
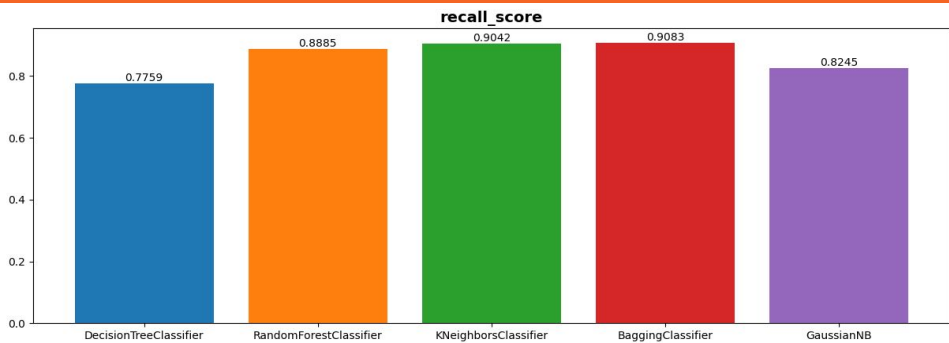
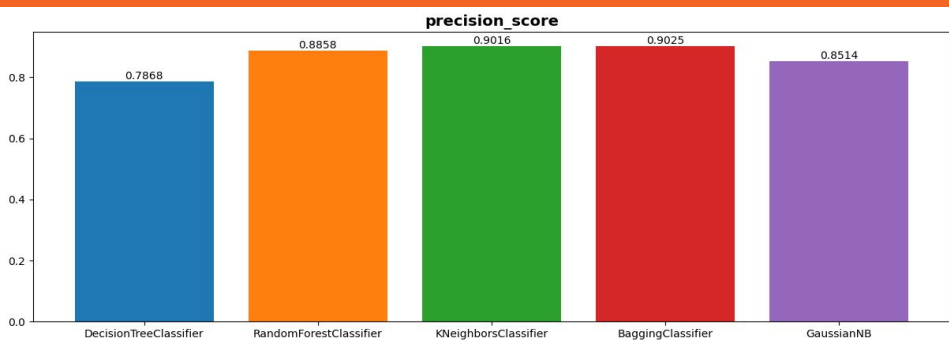
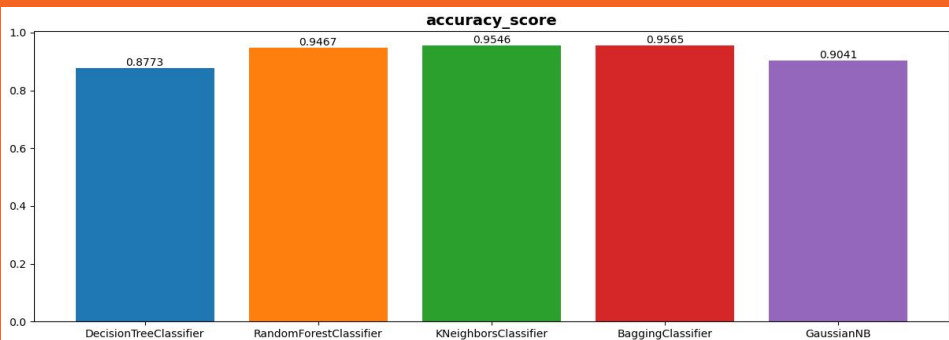


Matrica konfuzije

Leva se odnosi na **trening** skup, a desna na **test** skup.

Klasifikacija

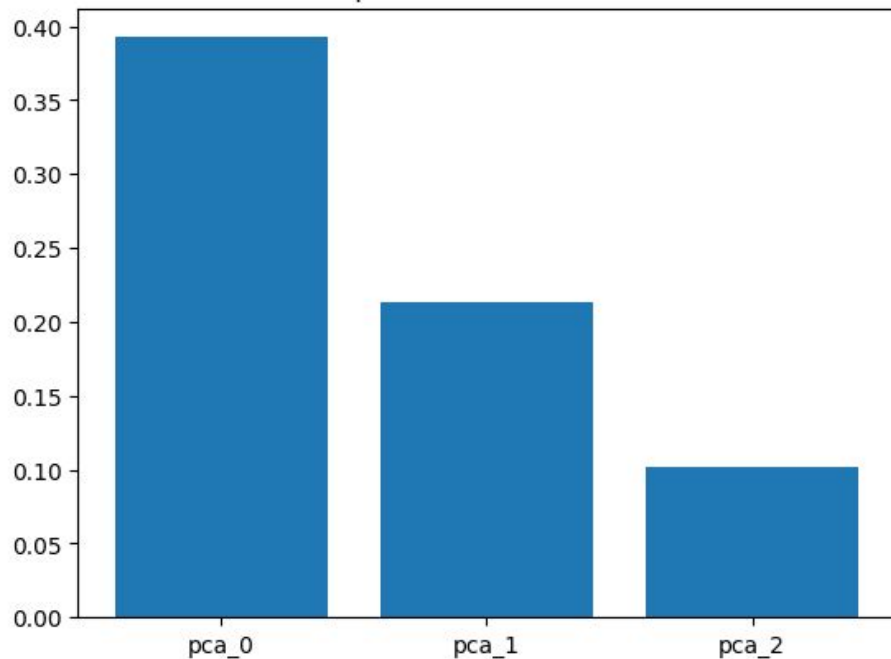
Poređenje modela



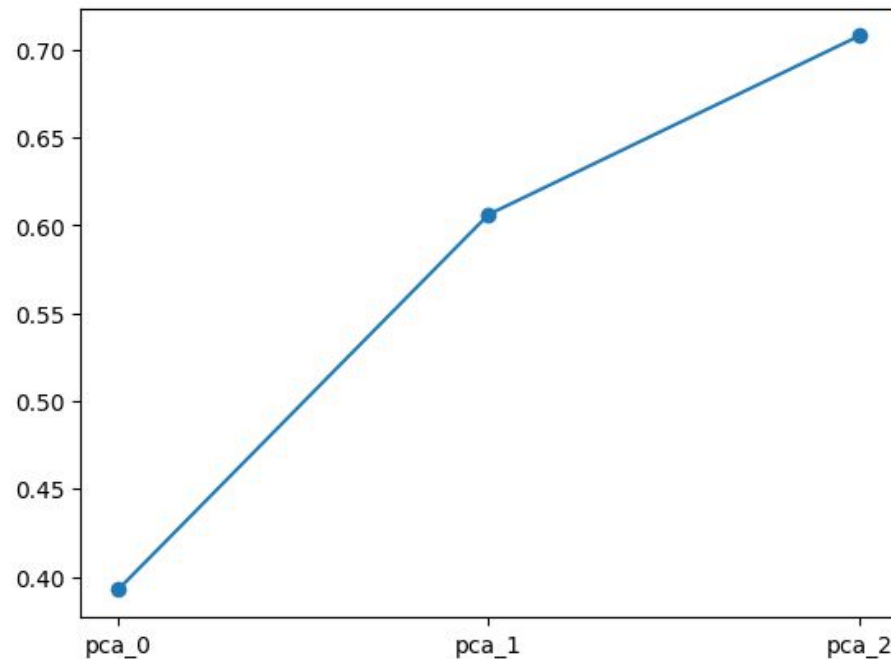
Klasterovanje

Pretprocesiranje

Explained variance ratio

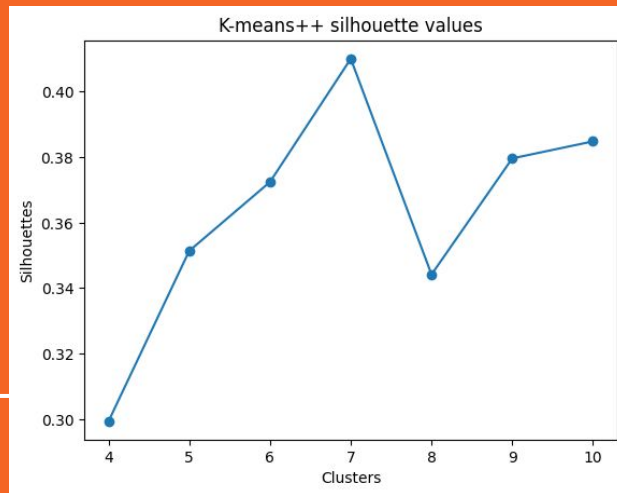
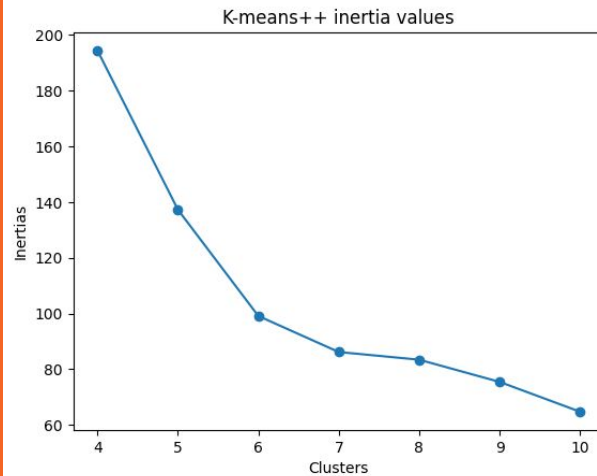
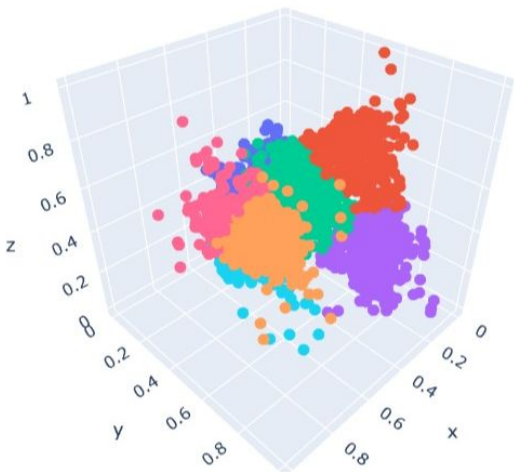
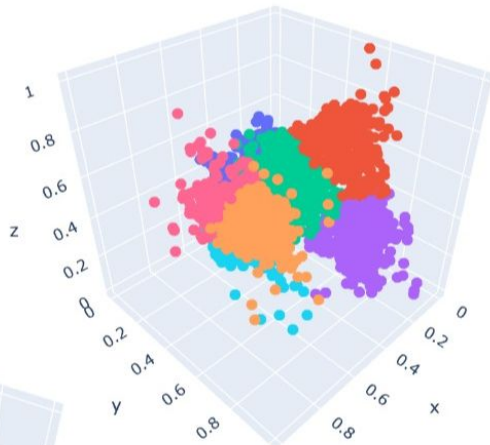


Cumulative variance ratio

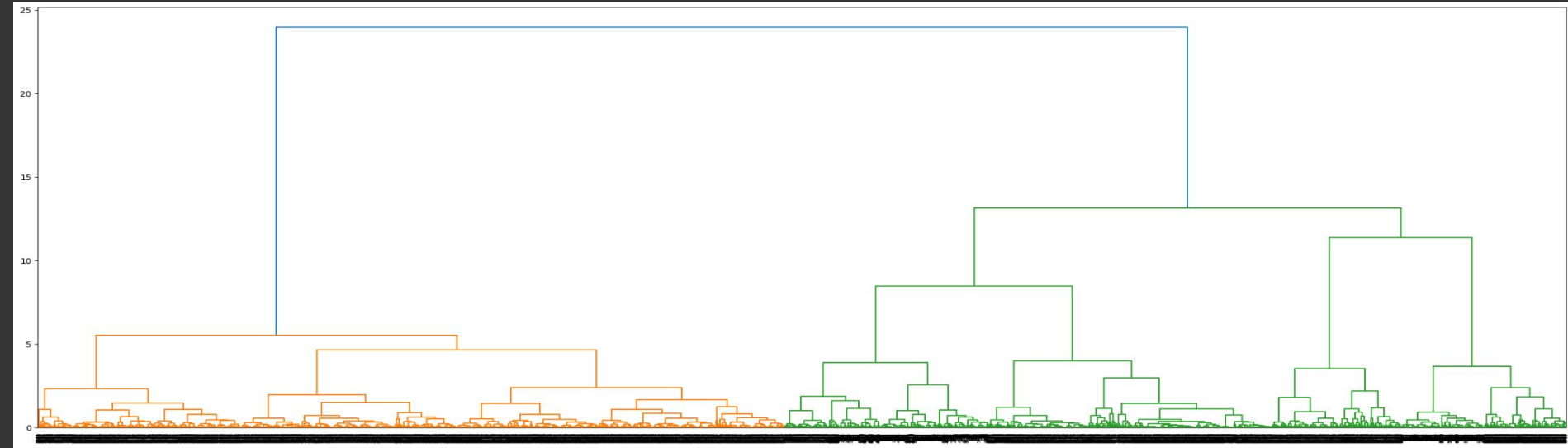
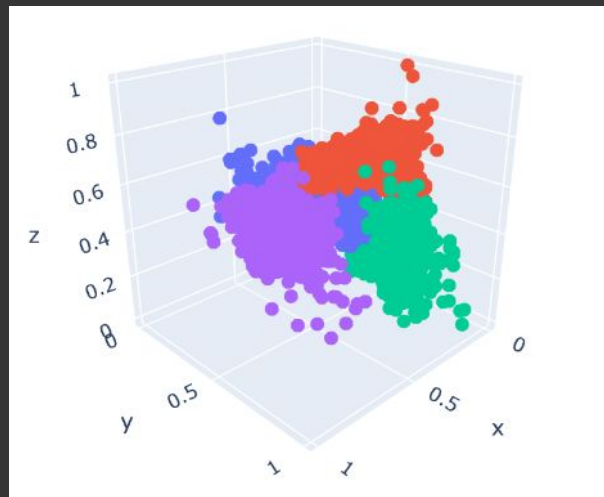


K sredina

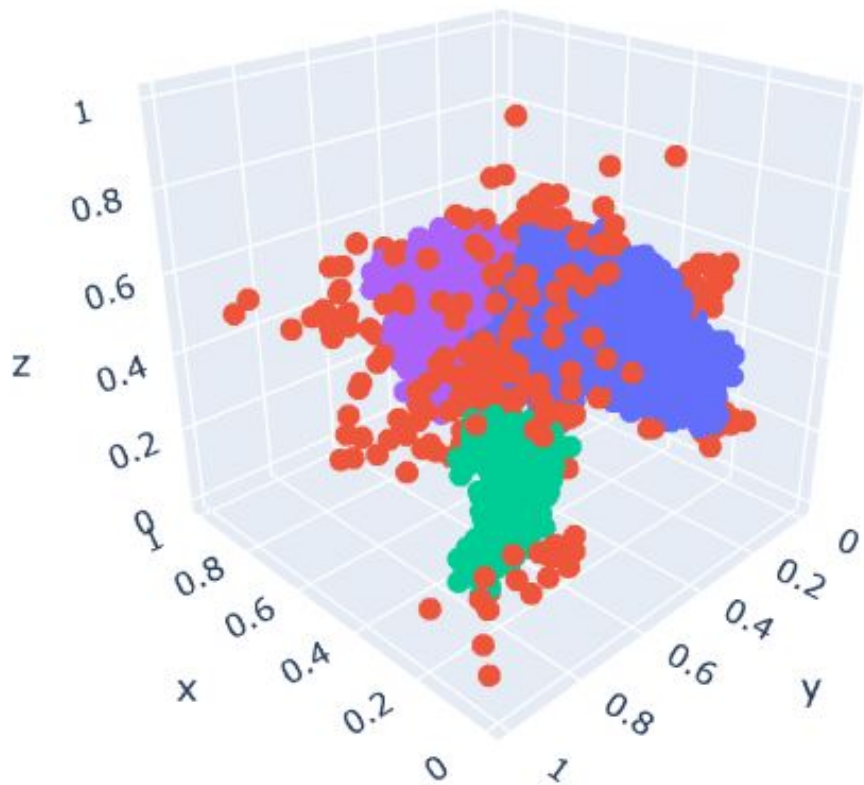
Prikaz sedam klastera koristeći *k-means++* i *random* načine za odabir centroida.



Hijerarhijsko klasterovanje



DBSCAN

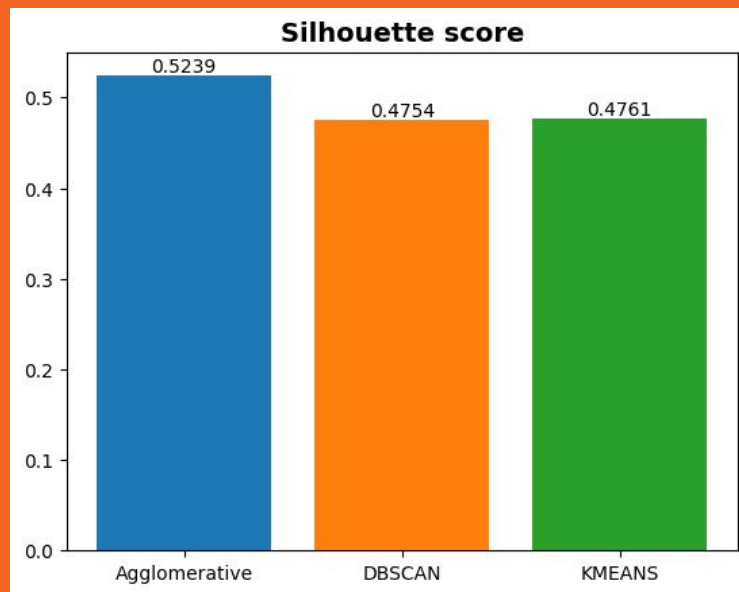
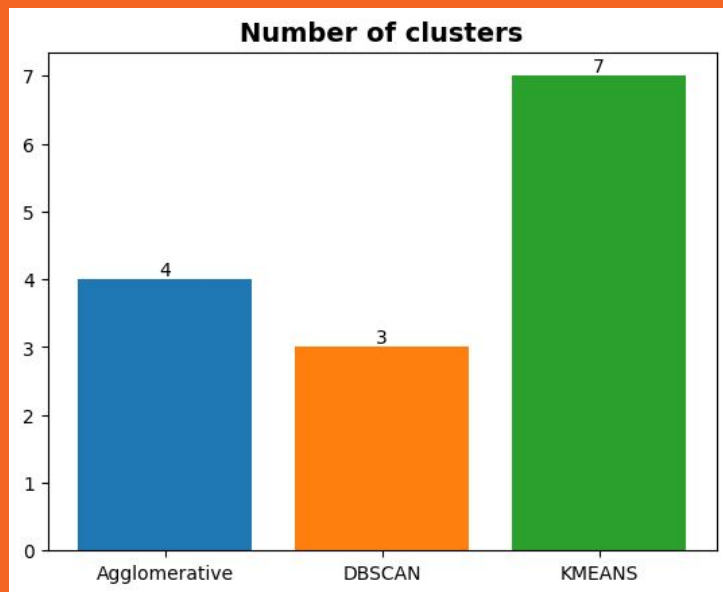


Klasteri

Algoritam definiše tri klastera, obojeni zelenom, plavom i ljubičastom bojom, i elemente van granica obojene crvenom bojom.

Klasterovanje

Poređenje modela

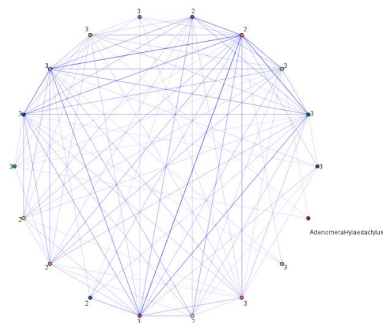


Pravila pridruživanja

Apriori

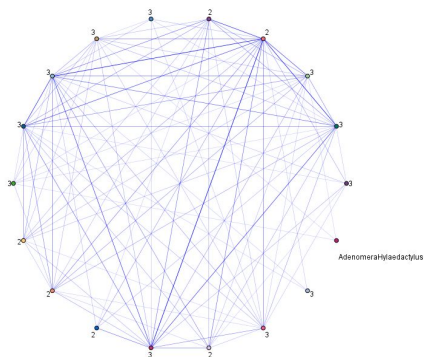
Consequent	Antecedent	Support %	Confidence %	Lift
Species = AdenomeraHylaedactylus	MFCCs_9_BIN = 3 MFCCs_10_BIN = 3	58.277	80.014	1.655
Species = AdenomeraHylaedactylus	MFCCs_9_BIN = 3 MFCCs_12_BIN = 3	56.511	82.612	1.709
Species = AdenomeraHylaedactylus	MFCCs_15_BIN = 2 MFCCs_5_BIN = 3 MFCCs_12_BIN = 3	55.942	80.547	1.666
Species = AdenomeraHylaedactylus	MFCCs_15_BIN = 2 MFCCs_5_BIN = 3 MFCCs_12_BIN = 3 MFCCs_10_BIN = 3	54.454	80.807	1.672
Species = AdenomeraHylaedactylus	MFCCs_9_BIN = 3 MFCCs_10_BIN = 3 MFCCs_6_BIN = 2	54.149	82.161	1.7

Consequent	Antecedent	Support %	Confidence %	Lift
Species = AdenomeraHylaedactylus	MFCCs_20_BIN = 1 MFCCs_11_BIN = 2 MFCCs_17_BIN = 3 MFCCs_3_BIN = 2	12.203	100.0	2.069
Species = AdenomeraHylaedactylus	MFCCs_20_BIN = 1 MFCCs_11_BIN = 2 MFCCs_17_BIN = 3 MFCCs_19_BIN = 2	12.064	100.0	2.069
Species = AdenomeraHylaedactylus	MFCCs_20_BIN = 1 MFCCs_11_BIN = 2 MFCCs_17_BIN = 3 MFCCs_13_BIN = 3	13.468	100.0	2.069
Species = AdenomeraHylaedactylus	MFCCs_20_BIN = 1 MFCCs_11_BIN = 2 MFCCs_17_BIN = 3 MFCCs_14_BIN = 2	11.383	100.0	2.069
Species = AdenomeraHylaedactylus	MFCCs_20_BIN = 1 MFCCs_11_BIN = 2 MFCCs_17_BIN = 3 MFCCs_16_BIN = 2	11.744	100.0	2.069



Consequent	Antecedent	Support %	Confidence %	Lift
Species = HypsiobasCordobae	MFCCs_3_BIN = 3 MFCCs_8_BIN = 2 MFCCs_11_BIN = 3 MFCCs_14_BIN = 2 MFCCs_12_BIN = 3	11.049	99.371	6.378
Species = HypsiobasCordobae	MFCCs_3_BIN = 3 MFCCs_8_BIN = 2 MFCCs_13_BIN = 2 MFCCs_14_BIN = 2 MFCCs_12_BIN = 3	10.716	99.351	6.377
Species = HypsiobasCordobae	MFCCs_3_BIN = 3 MFCCs_8_BIN = 2 MFCCs_13_BIN = 2 MFCCs_17_BIN = 2 MFCCs_14_BIN = 2	10.021	99.307	6.374
Species = HypsiobasCordobae	MFCCs_3_BIN = 3 MFCCs_8_BIN = 2 MFCCs_11_BIN = 3 MFCCs_17_BIN = 2 MFCCs_14_BIN = 2	10.354	99.195	6.367
Species = HypsiobasCordobae	MFCCs_3_BIN = 3 MFCCs_8_BIN = 2 MFCCs_13_BIN = 2 MFCCs_14_BIN = 2 MFCCs_20_BIN = 2	10.688	99.09	6.36

Pravila pridruživanja FP-Growth



MFCCs_10_BIN MFCCs_11_BIN MFCCs_12_BIN MFCCs_13_BIN MFCCs_14_BIN MFCCs_15_BIN MFCCs_16_BIN MFCCs_17_BIN MFCCs_18_BIN MFCCs_19_BIN MFCCs_2_BIN
MFCCs_20_BIN MFCCs_21_BIN MFCCs_22_BIN MFCCs_3_BIN MFCCs_4_BIN MFCCs_5_BIN MFCCs_6_BIN MFCCs_7_BIN MFCCs_8_BIN MFCCs_9_BIN Species

Most Interesting Rules by Rule Support

Rank	Rule ID	Condition	Prediction	Sorted By Rule Support(%)	Other Evaluation Statistics			
					Condition Support (%)	Confidence (%)	Lift	Deployability (%)
1	984	MFCCs_16_BIN = 2 MFCCs_12_BIN = 2	MFCCs_14_BIN = 3	15.82	18.28	86.54	2.83	2.46
2	990	MFCCs_4_BIN = 3 MFCCs_12_BIN = 2	MFCCs_14_BIN = 3	14.64	17.08	85.68	2.80	2.45
3	970	MFCCs_18_BIN = 3 MFCCs_12_BIN = 2	MFCCs_14_BIN = 3	14.50	16.50	87.87	2.88	2.00
4	952	MFCCs_17_BIN = 2 MFCCs_12_BIN = 2	MFCCs_14_BIN = 3	14.32	15.98	89.57	2.93	1.67
5	954	MFCCs_18_BIN = 3 MFCCs_16_BIN = 2 MFCCs_12_BIN = 2	MFCCs_14_BIN = 3	13.50	15.08	89.49	2.93	1.58

Most Interesting Rules by Confidence

Rank	Rule ID	Condition	Prediction	Sorted By Confidence(%)	Other Evaluation Statistics			
					Condition Support (%)	Rule Support (%)	Lift	Deployability (%)
1	1	MFCCs_12_BIN = 3 MFCCs_14_BIN = 2 MFCCs_11_BIN = 3 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	99.37	11.05	10.98	6.38	0.07
2	2	MFCCs_12_BIN = 3 MFCCs_14_BIN = 2 MFCCs_13_BIN = 2 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	99.35	10.72	10.65	6.38	0.07
3	3	MFCCs_14_BIN = 2 MFCCs_17_BIN = 2 MFCCs_11_BIN = 3 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	99.19	10.35	10.27	6.37	0.08
4	4	MFCCs_20_BIN = 2 MFCCs_14_BIN = 2 MFCCs_13_BIN = 2 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	99.09	10.69	10.59	6.36	0.10
5	5	MFCCs_20_BIN = 2 MFCCs_14_BIN = 2 MFCCs_11_BIN = 3 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	98.99	11.04	10.92	6.35	0.11

Most Interesting Rules by Lift

Rank	Rule ID	Condition	Prediction	Sorted By Lift	Condition Support (%)	Other Evaluation Statistics		
						Confidence (%)	Rule Support (%)	Deployability (%)
1	1	MFCCs_12_BIN = 3 MFCCs_14_BIN = 2 MFCCs_11_BIN = 3 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	6.38	11.05	99.37	10.98	0.07
2	2	MFCCs_12_BIN = 3 MFCCs_14_BIN = 2 MFCCs_13_BIN = 2 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	6.38	10.72	99.35	10.65	0.07
3	3	MFCCs_14_BIN = 2 MFCCs_17_BIN = 2 MFCCs_11_BIN = 3 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	6.37	10.35	99.19	10.27	0.08
4	4	MFCCs_20_BIN = 2 MFCCs_14_BIN = 2 MFCCs_13_BIN = 2 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	6.36	10.69	99.09	10.59	0.10
5	5	MFCCs_20_BIN = 2 MFCCs_14_BIN = 2 MFCCs_11_BIN = 3 MFCCs_8_BIN = 2 MFCCs_3_BIN = 3	Species = HypsiboasCordobae	6.35	11.04	98.99	10.92	0.11