

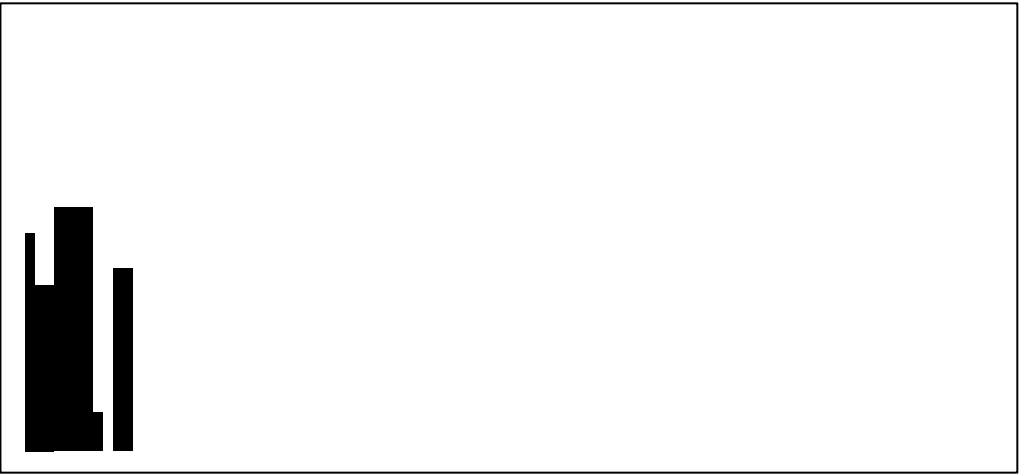
OTU_442;size=1158;.csv

Providencia stuartii

Enterobacteriaceae

Enterobacteriales

90.24



Providencia stuartii strain ATCC 33672, complete g	CP008920	90.24	18.39	0.00167958	Enterobacteriales	Enterobacteriaceae	Providencia stuartii
Providencia stuartii strain FDAARGOS_145, complete	CP014024	87.8	18.39	0.0204614	Enterobacteriales	Enterobacteriaceae	Providencia stuartii
Providencia stuartii MRSN 2154, complete genome	CP003488	87.8	18.39	0.0204614	Enterobacteriales	Enterobacteriaceae	Providencia stuartii MRSN 2154
PREDICTED: Monodelphis domestica intraflagellar tr	XM_001366349	91.43	15.25	3.03675	Didelphimorphia	Didelphidae	Monodelphis domestica
PREDICTED: Monodelphis domestica intraflagellar tr	XM_007495202	91.43	15.25	3.03675	Didelphimorphia	Didelphidae	Monodelphis domestica
PREDICTED: Monodelphis domestica intraflagellar tr	XM_016421847	91.43	15.25	3.03675	Didelphimorphia	Didelphidae	Monodelphis domestica
PREDICTED: Monodelphis domestica intraflagellar tr	XM_007495201	91.43	15.25	3.03675	Didelphimorphia	Didelphidae	Monodelphis domestica
Schistosoma margrebowiei genome assembly S_margreb	LL882280	81.82	19.73	3.03675	Strigeidida	Schistosomatidae	Schistosoma margrebowiei
Strongyloides ratti genome assembly S_ratti_ED321	LN609398	70.41	43.5	3.03675	Rhabditida	Strongyloididae	Strongyloides ratti
Bathycoccus prasinus CMP-N-acetylneuraminate-poly-	XM_007513887	88.57	15.7	3.03675	Mamiellales	Bathycoccaceae	Bathycoccus prasinus
Bathycoccus prasinus genomic : Chromosome_3	FO082276	88.57	15.7	3.03675	Mamiellales	Bathycoccaceae	Bathycoccus prasinus
Sus scrofa BAC clone KNP_442E8	FN673743	73.56	37.67	3.03675		Suidae	Sus scrofa
Sus scrofa BAC clone KNP_47E7	FN673711	73.56	37.67	3.03675		Suidae	Sus scrofa

