

Entrée [1]: `import pandas
import os`

Entrée [10]: `th = pandas.read_csv("1335_vdh_tab" sep='\t' lineterminator='\n')`

Entrée [11]: `hev_shane`

Out[11]: (4106, 16)

Entrée [23]: `#split column containing taxonomy levels into 6 columns
new = th["OTUConTaxonomy"].str.split(".", n=7, expand=True)`

Entrée [24]: `new.head`

Out[24]: <bound method NDFrame.head of 0

	1	2 \	0
0	Bacteria(100)	Actinobacteria(100)	Actinobacteria(100)
1	Bacteria(100)	Proteobacteria(100)	Gammaproteobacteria(100)
2	Bacteria(100)	Actinobacteria(100)	Actinobacteria(100)
3	Bacteria(100)	Actinobacteria(100)	Actinobacteria(100)
4	Bacteria(100)	Actinobacteria(100)	Actinobacteria(100)
...
4101	Bacteria(100)	Proteobacteria(100)	Alphaproteobacteria(100)
4102	Bacteria(100)	Firmicutes(100)	unclassified(100)
4103	Bacteria(100)	Acidobacteria(100)	Acidobacteria_Gp6(100)
4104	Bacteria(100)	Bacteroidetes(100)	Sphingobacteriia(100)
4105	NaN	NaN	NaN

	3	4 \
0	Actinomycetales(100)	Propionibacteriaceae(100)
1	Pseudomonadales(100)	Moraxellaceae(100)
2	Actinomycetales(100)	Corynebacteriaceae(100)
3	Actinomycetales(100)	Micrococcaceae(99)
4	Actinomycetales(100)	Corynebacteriaceae(100)
...
4101	Rhizobiales(100)	unclassified(100)
4102	unclassified(100)	unclassified(100)
4103	Gp6(100)	unclassified(100)
4104	Sphingobacteriales(100)	Chitinophagaceae(100)
4105	NaN	NaN

	5	6
0	Propionibacterium(100)	
1	Enhydrobacter(100)	
2	Corynebacterium(100)	
3	Arthrobacter(86)	
4	Corynebacterium(100)	
...
4101	unclassified(100)	
4102	unclassified(100)	
4103	unclassified(100)	
4104	unclassified(100)	
4105	NaN	NaN

[4106 rows x 7 columns]>

Entrée [25]: `new_shane`

Out[25]: (4106, 7)

Entrée [28]: `head(new["1"])`

```
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NameError                                Traceback (most recent c  
all last)  
<ipython-input-28-bf21fccffc3d> in <module>  
----> 1 head(new["1"])
```

NameError: name 'head' is not defined

Entrée []: