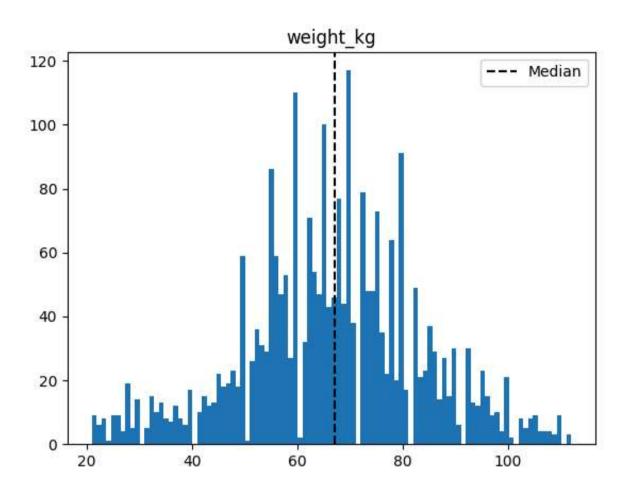
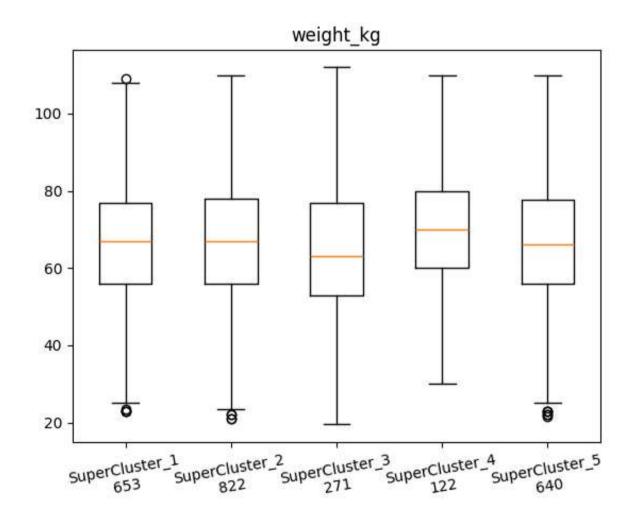
Weight_Kg

Clean data



Clean data



Kruskal-Wallis H-test - Significant

p-value = 0.023962060953509407

1 2 3 4 5 1 1.000000 1.000000 0.162333 0.397079 1.000000

2 1.000000 1.000000 0.101256 0.397079 1.000000

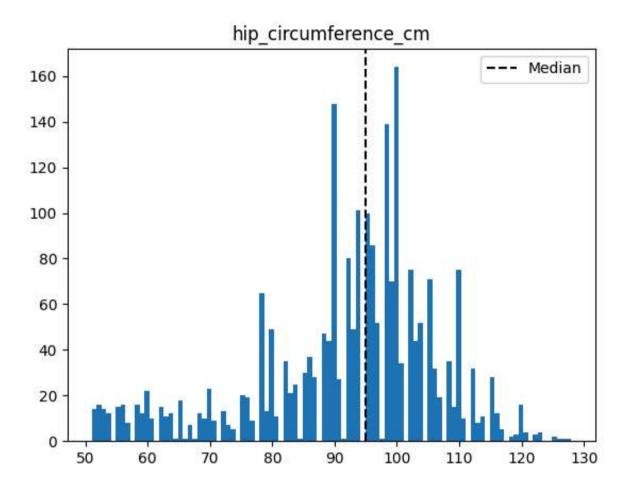
3 0.162333 0.101256 1.000000 0.017776 0.260971

4 0.397079 0.397079 **0.017776** 1.000000 0.328374

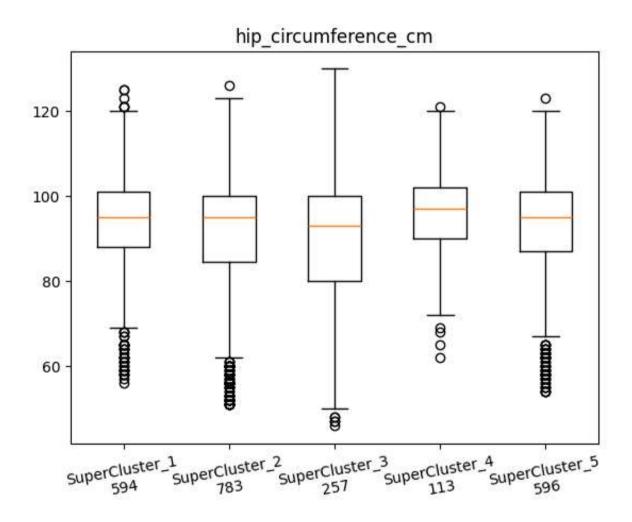
5 1.000000 1.000000 0.260971 0.328374 1.000000

Hip_Circumference_Cm

Clean data



Clean data



Kruskal-Wallis H-test - Significant

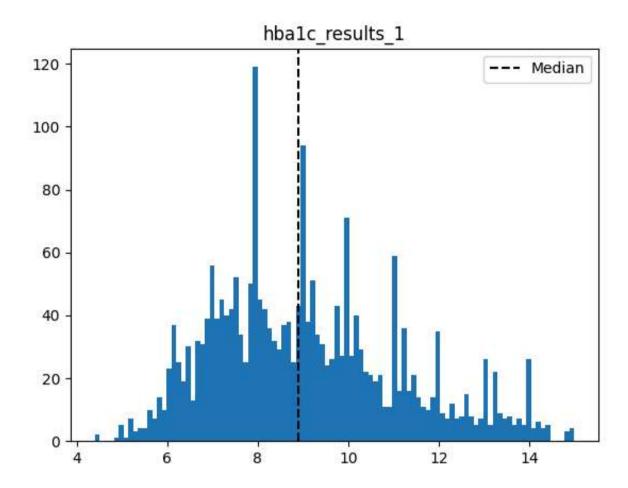
p-value = 0.043661904386337276

1 2 3 4 5 1 1.000000 0.788689 0.126628 0.788689 0.941246 2 0.788689 1.000000 0.788689 0.394075 0.941246 3 0.126628 0.788689 1.000000 0.077581 0.382819 4 0.788689 0.394075 0.077581 1.000000 0.788689

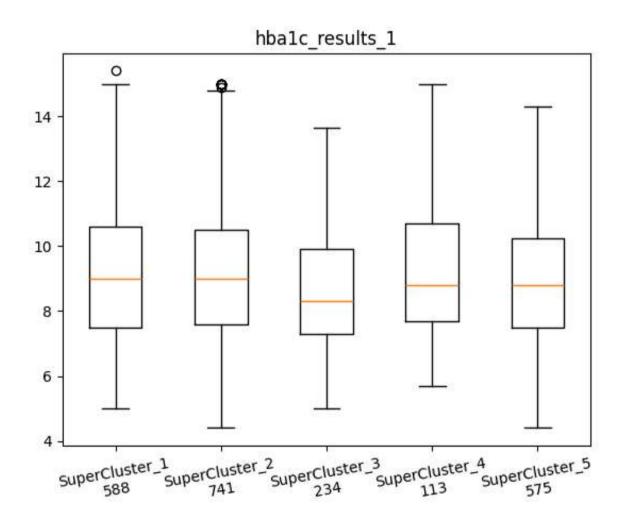
5 0.941246 0.941246 0.382819 0.788689 1.000000

Hba1C_Results_1

Clean data



Clean data



Kruskal-Wallis H-test - Significant

p-value = 0.017913429247121138

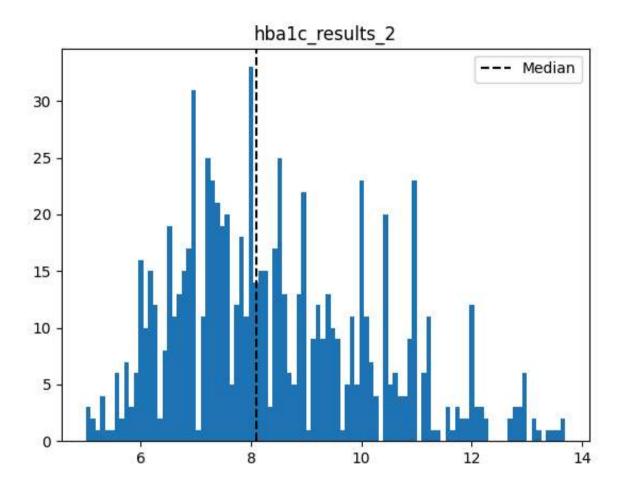
1 2 3 4 5 1 1.000000 1.000000 0.014380 1.000000 1.000000 2 1.000000 1.000000 0.019161 1.000000 1.000000 3 0.014380 0.019161 1.000000 0.390454 0.260709

4 1.000000 1.000000 0.390454 1.000000 1.000000

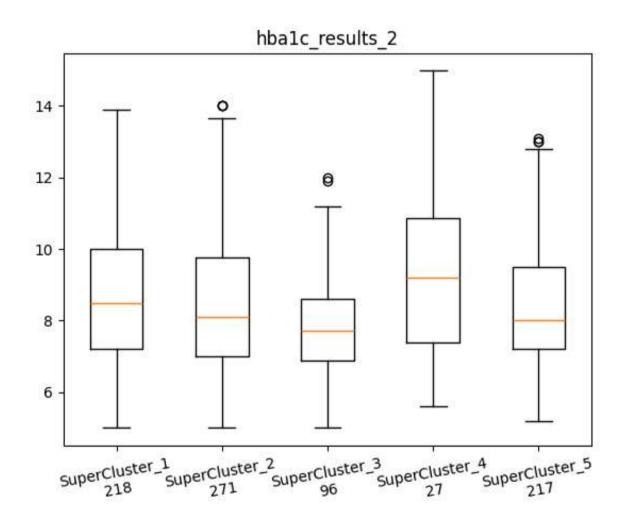
5 1.000000 1.000000 0.260709 1.000000 1.000000

Hba1C_Results_2

Clean data



Clean data



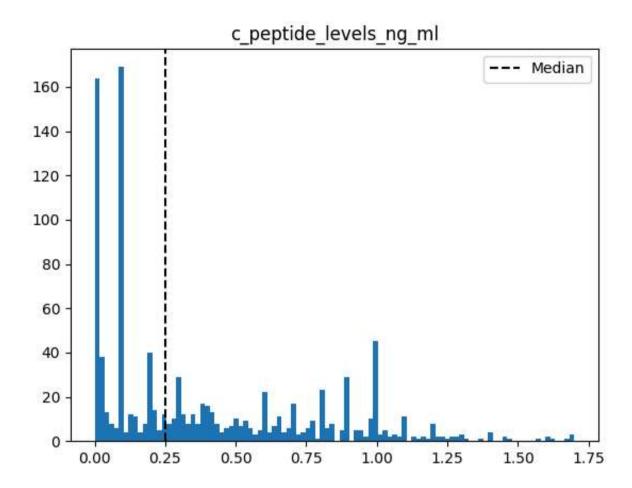
Kruskal-Wallis H-test - Significant

p-value = 0.009783854350719256

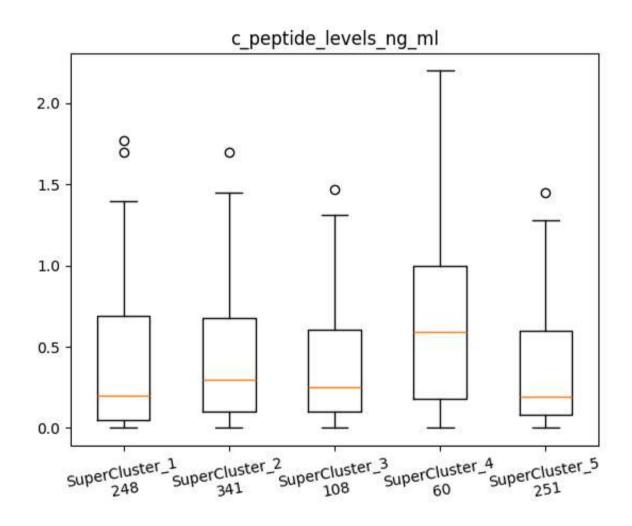
1 2 3 4 5 1 1.000000 0.526216 0.010626 0.809234 0.554319 2 0.526216 1.000000 0.270581 0.534411 0.809234 3 0.010626 0.270581 1.000000 0.078866 0.226782 4 0.809234 0.534411 0.078866 1.000000 0.554319 5 0.554319 0.809234 0.226782 0.554319 1.000000

C_Peptide_Levels_Ng_Ml

Clean data



Clean data



Kruskal-Wallis H-test - Significant

p-value = 0.0005656662277714814

1 2 3 4 5 1 1.000000 1.000000 1.000000 0.000561 1.000000 2 1.000000 1.000000 1.000000 0.005691 0.585994

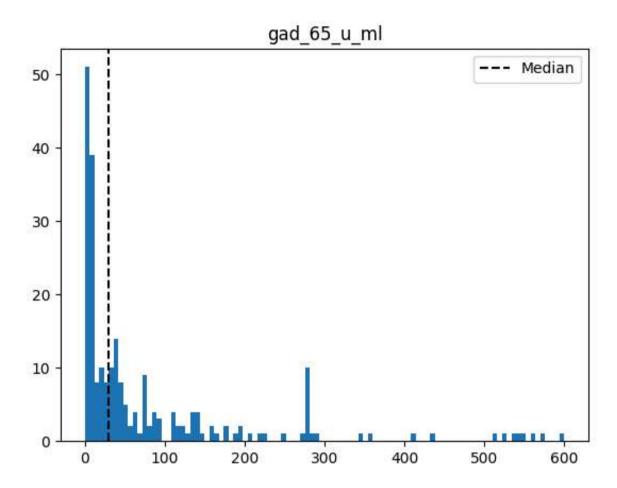
3 1.000000 1.000000 1.000000 **0.010295** 1.000000

4 0.000561 0.005691 0.010295 1.000000 0.000208

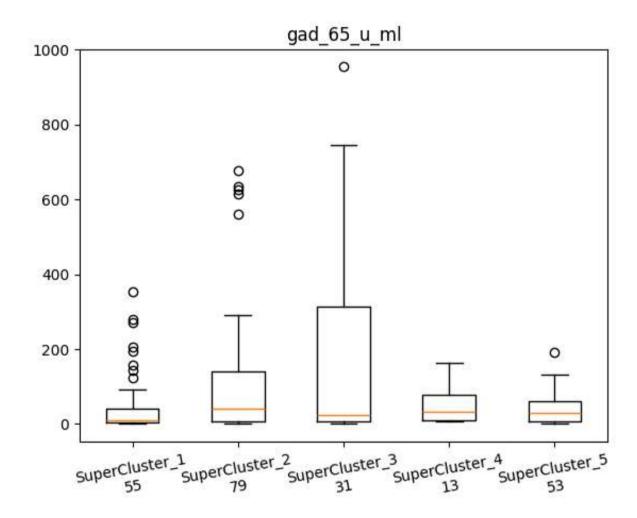
5 1.000000 0.585994 1.000000 **0.000208** 1.000000

Gad_65_U_Ml

Clean data



Clean data



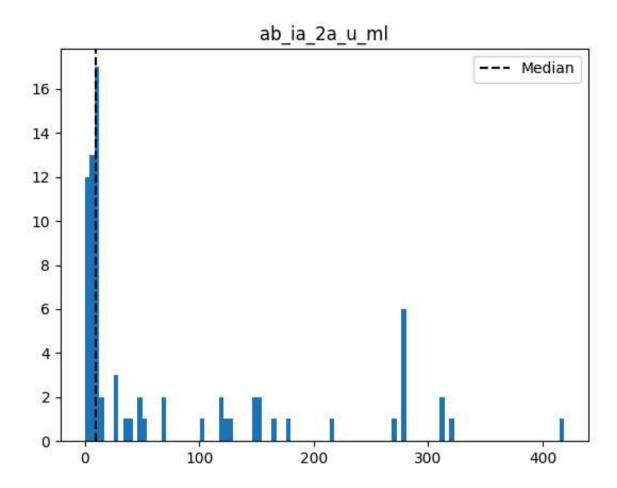
Kruskal-Wallis H-test - Significant

p-value = 0.03707873106994417

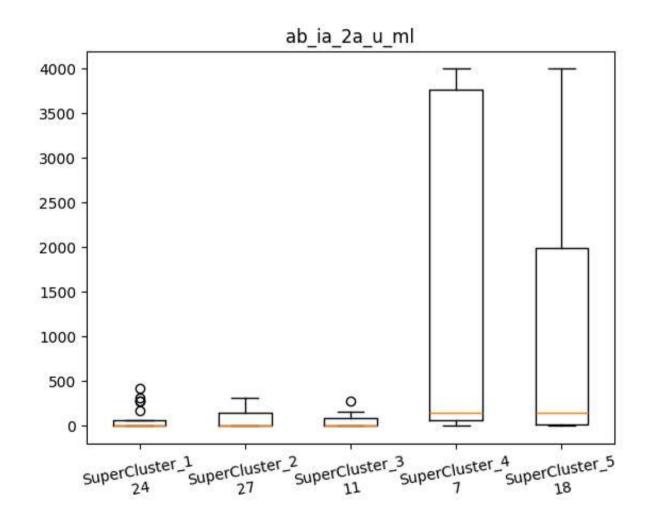
1 2 3 4 5 1 1.000000 0.062413 0.163075 0.871036 1.000000 2 0.062413 1.000000 1.000000 1.000000 0.755657 3 0.163075 1.000000 1.000000 1.000000 0.871036 4 0.871036 1.000000 1.000000 1.000000 1.000000 5 1.000000 0.755657 0.871036 1.000000 1.000000

Ab_Ia_2A_U_Ml

Clean data



Clean data



Kruskal-Wallis H-test - Significant

p-value = 0.04472133739513693

1 2 3 4 5 1 1.000000 1.000000 1.000000 0.488425 0.192237

2 1.000000 1.000000 1.000000 0.583209 0.268032

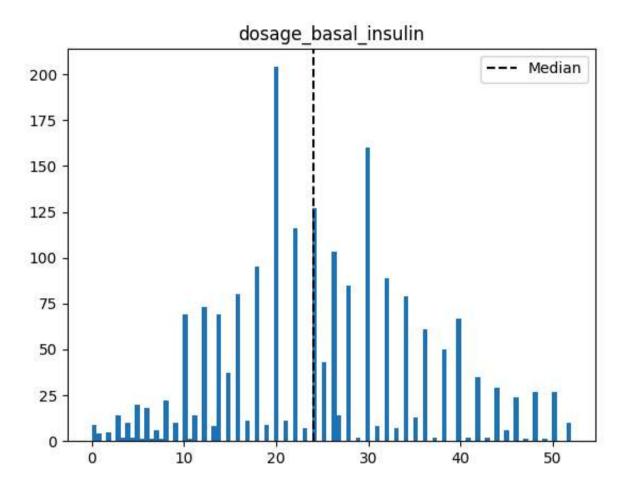
3 1.000000 1.000000 1.000000 0.455534 0.202386

4 0.488425 0.583209 0.455534 1.000000 1.000000

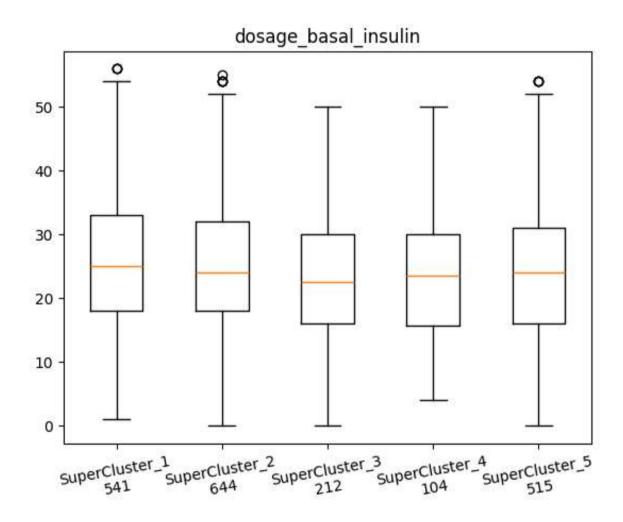
5 0.192237 0.268032 0.202386 1.000000 1.000000

Dosage_Basal_Insulin

Clean data



Clean data



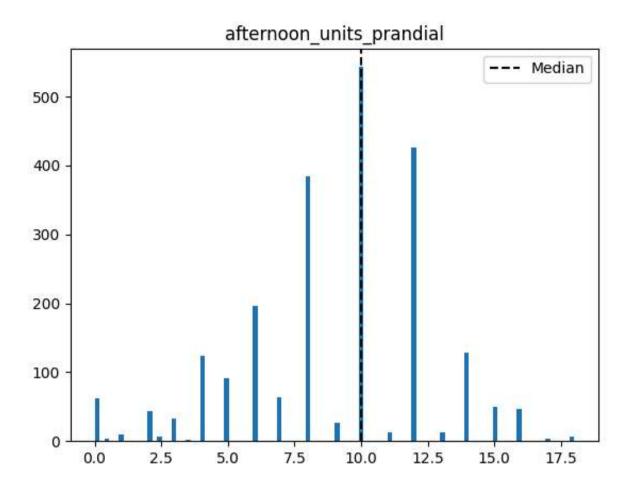
Kruskal-Wallis H-test - Significant

p-value = 0.0472281038968443

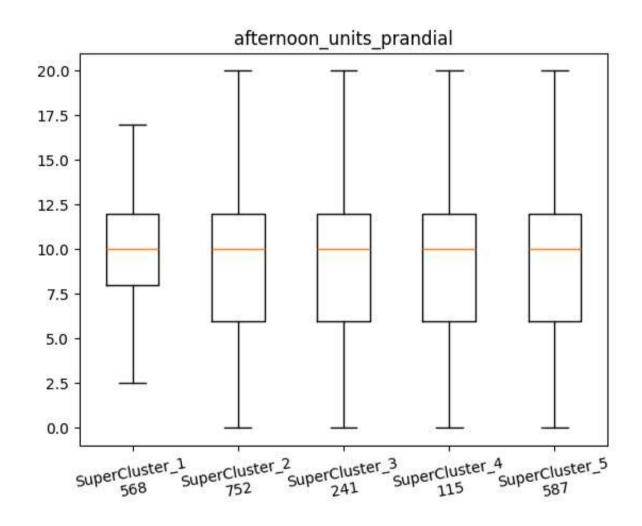
1 2 3 4 5 1 1.000000 1.000000 0.211129 0.316020 0.424907 2 1.000000 1.000000 0.514565 0.526215 1.000000 3 0.211129 0.514565 1.000000 1.000000 1.000000 4 0.316020 0.526215 1.000000 1.000000 1.000000 5 0.424907 1.000000 1.000000 1.000000 1.000000

Afternoon_Units_Prandial

Clean data



Clean data



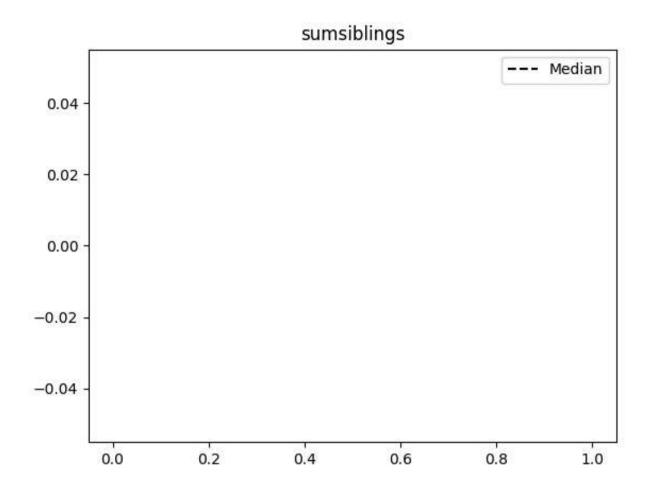
Kruskal-Wallis H-test - Significant

p-value = 0.04200175637275408

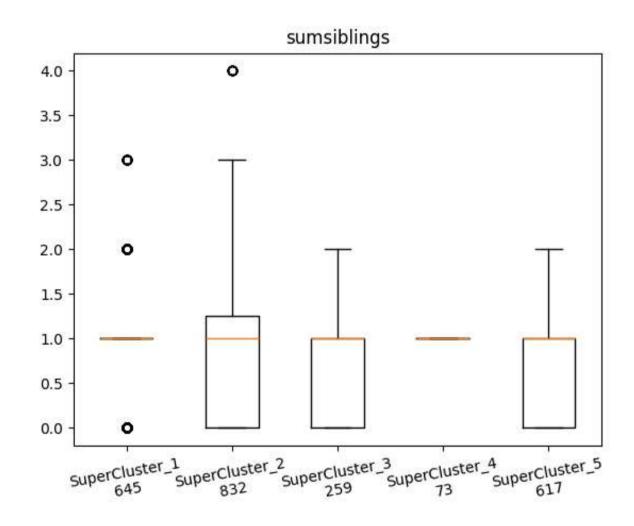
1 2 3 4 5 1 1.000000 0.183496 0.108832 0.941841 0.108832 2 0.183496 1.000000 1.000000 1.000000 1.000000 3 0.108832 1.000000 1.000000 1.000000 1.000000 4 0.941841 1.000000 1.000000 1.000000 1.000000 5 0.108832 1.000000 1.000000 1.000000 1.000000

Sumsiblings

Clean data



Clean data



Kruskal-Wallis H-test - Significant

p-value = 4.79135848360545e-05

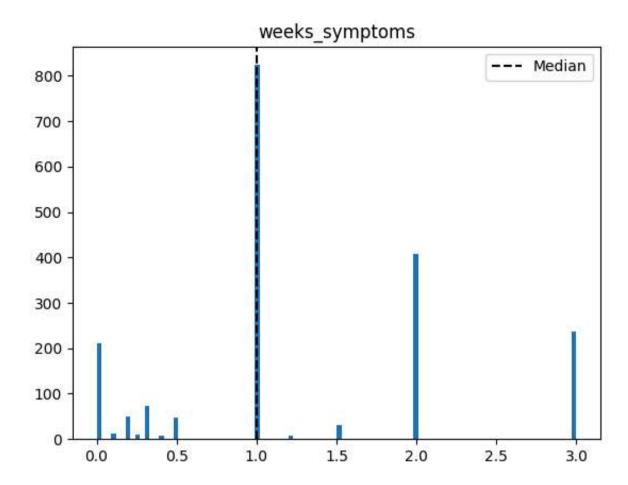
1 2 3 4 5 1 1.000000 1.000000 0.105327 1.000000 0.000535 2 1.000000 1.000000 0.105327 1.000000 0.000375 3 0.105327 0.105327 1.000000 0.672707 1.000000

4 1.000000 1.000000 0.672707 1.000000 0.303302

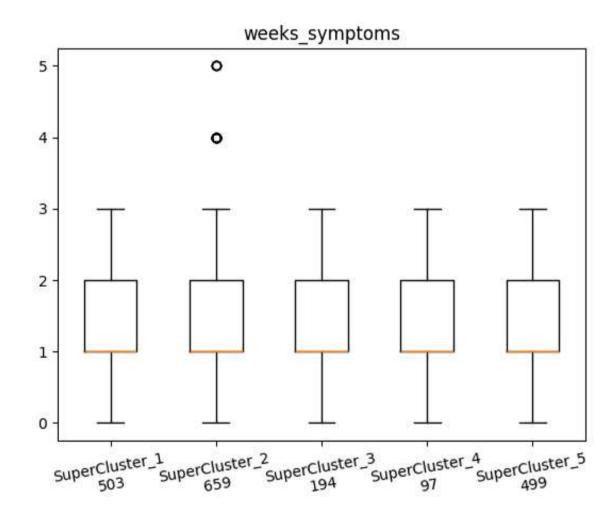
5 0.000535 0.000375 1.000000 0.303302 1.000000

Weeks_Symptoms

Clean data



Clean data



Kruskal-Wallis H-test - Significant

```
p-value = 0.001976686603427168

1 2 3 4 5

1 1.000000 0.004021 1.000000 1.000000 1.000000
2 0.004021 1.000000 1.000000 1.000000 0.006198
3 1.000000 1.000000 1.000000 1.000000 1.000000
4 1.000000 1.000000 1.000000 1.000000 1.000000
5 1.000000 0.006198 1.000000 1.000000 1.000000
```