

	643311	4444760	687185	192795	164413	176468
RSL1	-1.1478018	4.4090263	-1.1478018	-1.1478018	-1.1478018	5.3772279
RSL3	-1.1994810	4.4317308	-1.1994810	-1.1994810	-1.1994810	5.6194430
RSL2	-1.1966517	4.0556217	3.5395467	-1.1966517	-1.1966517	5.3269106
RRD1	-2.3923418	3.4307041	-2.3923418	-2.3923418	-2.3923418	5.4147615
TPB3	-1.7247157	3.1194713	-1.7247157	-1.7247157	-1.7247157	3.1656334
NVB2	-1.3798340	4.3884869	-1.3798340	-1.3798340	4.4222843	5.3379706
DTL1	-1.8317339	-1.8317339	-1.8317339	3.5799122	-1.8317339	6.1895223
RRD2	-2.1666981	3.0150854	2.1373670	-2.1666981	-2.1666981	5.2746223
TPB2	-2.1883090	3.5924345	-2.1883090	-2.1883090	-2.1883090	4.4963027
RRD3	2.5201835	2.7484421	-2.5296725	-2.5296725	-2.5296725	5.2079438
NVB3	-1.3292235	3.7077291	-1.3292235	-1.3292235	4.6672286	5.7944493
DTL3	-1.5350387	-1.5350387	-1.5350387	-1.5350387	-1.5350387	5.8551427
DTL2	-1.7574065	-1.7574065	-1.7574065	-1.7574065	-1.7574065	6.2022188
TPB1	-2.1120912	3.5439006	-2.1120912	-2.1120912	-2.1120912	4.9645626
NVB1	-1.2249696	4.6912325	-1.2249696	-1.2249696	3.4289908	5.6986591
K6B2	-0.4585928	-0.4585928	-0.4585928	-0.4585928	4.2142361	-0.4585928
K6B1	-0.3850764	-0.3850764	-0.3850764	-0.3850764	-0.3850764	-0.3850764
L8B2	-0.4683688	-0.4683688	-0.4683688	-0.4683688	6.9389489	-0.4683688
L8B3	-0.4582567	-0.4582567	-0.4582567	-0.4582567	6.6005014	-0.4582567
TRS2	-0.6781239	-0.6781239	-0.6781239	-0.6781239	-0.6781239	-0.6781239
S4B1	-0.3564396	-0.3564396	-0.3564396	-0.3564396	-0.3564396	-0.3564396
R9B3	-0.4685848	-0.4685848	-0.4685848	-0.4685848	6.8683521	-0.4685848
TRS1	-0.6329046	-0.6329046	-0.6329046	-0.6329046	-0.6329046	4.3229225
TRS3	-0.5340613	-0.5340613	-0.5340613	-0.5340613	-0.5340613	-0.5340613
S4B3	-0.3228014	-0.3228014	-0.3228014	-0.3228014	-0.3228014	-0.3228014
C3B2	-0.3461644	-0.3461644	-0.3461644	-0.3461644	-0.3461644	-0.3461644
R9B1	-0.4067263	-0.4067263	-0.4067263	-0.4067263	6.5899552	-0.4067263
TB13	-1.1456869	-1.1456869	4.3096342	-1.1456869	-1.1456869	-1.1456869
K6B3	-0.4322439	-0.4322439	-0.4322439	-0.4322439	3.3519457	-0.4322439
S7B2	-0.3415675	-0.3415675	-0.3415675	-0.3415675	-0.3415675	-0.3415675

C3B3	-0.2672858	-0.2672858	-0.2672858	-0.2672858	-0.2672858	-0.2672858
R9B2	-0.4176403	-0.4176403	-0.4176403	-0.4176403	6.5744562	-0.4176403
TB22	-1.0574762	-1.0574762	3.6787223	-1.0574762	-1.0574762	-1.0574762
TB23	-1.0876274	-1.0876274	3.9230079	-1.0876274	3.4556674	-1.0876274
T5B2	-0.4862838	-0.4862838	-0.4862838	-0.4862838	6.8655160	-0.4862838
T5B1	-0.5194111	-0.5194111	-0.5194111	-0.5194111	7.1486826	-0.5194111
T5B3	-0.5091447	-0.5091447	-0.5091447	-0.5091447	6.5817651	-0.5091447
TB21	-0.9675910	-0.9675910	2.9836527	-0.9675910	-0.9675910	-0.9675910
L8B1	-0.3614232	-0.3614232	-0.3614232	-0.3614232	5.4131284	-0.3614232
TB11	-1.0916114	-1.0916114	3.4830996	-1.0916114	-1.0916114	-1.0916114
TB12	-0.8011263	-0.8011263	3.1108967	-0.8011263	-0.8011263	-0.8011263
S7B3	-0.3484924	-0.3484924	-0.3484924	-0.3484924	-0.3484924	-0.3484924
C3B1	-0.2899390	-0.2899390	-0.2899390	-0.2899390	-0.2899390	-0.2899390
S7B1	-0.2623762	-0.2623762	-0.2623762	-0.2623762	-0.2623762	-0.2623762
S4B2	-0.3394032	-0.3394032	-0.3394032	-0.3394032	-0.3394032	-0.3394032
	255657	181589	219151	4425571		
RSL1	-1.1478018	5.8077908	-1.1478018	-1.1478018		
RSL3	-1.1994810	6.0701357	-1.1994810	-1.1994810		
RSL2	-1.1966517	5.5065364	-1.1966517	-1.1966517		
RRD1	-2.3923418	4.2409767	2.7781422	-2.3923418		
TPB3	-1.7247157	2.9757646	3.9277734	1.9388459		
NVB2	-1.3798340	3.6953398	3.0628172	-1.3798340		
DTL1	-1.8317339	5.8131855	-1.8317339	-1.8317339		
RRD2	-2.1666981	4.5191628	2.5337823	-2.1666981		
TPB2	-2.1883090	3.3088592	4.1467452	2.4751301		
RRD3	-2.5296725	4.3760808	2.8030463	-2.5296725		
NVB3	-1.3292235	4.0824226	-1.3292235	-1.3292235		
DTL3	-1.5350387	5.4224587	-1.5350387	-1.5350387		
DTL2	-1.7574065	5.8369747	-1.7574065	-1.7574065		
TPB1	-2.1120912	3.5983358	4.6081289	2.9938542		
NVB1	-1.2249696	4.1731931	3.7095044	-1.2249696		
K6B2	-0.4585928	4.2863394	-0.4585928	-0.4585928		
K6B1	-0.3850764	4.4824581	-0.3850764	-0.3850764		
L8B2	-0.4683688	-0.4683688	-0.4683688	-0.4683688		
L8B3	-0.4582567	-0.4582567	-0.4582567	-0.4582567		
TRS2	4.7109478	-0.6781239	-0.6781239	-0.6781239		
S4B1	-0.3564396	-0.3564396	-0.3564396	-0.3564396		
R9B3	-0.4685848	-0.4685848	-0.4685848	-0.4685848		
TRS1	4.5375794	-0.6329046	-0.6329046	-0.6329046		
TRS3	4.7337969	-0.5340613	-0.5340613	-0.5340613		
S4B3	-0.3228014	-0.3228014	-0.3228014	-0.3228014		
C3B2	-0.3461644	-0.3461644	-0.3461644	-0.3461644		
R9B1	-0.4067263	-0.4067263	-0.4067263	-0.4067263		
TB13	-1.1456869	5.1587619	-1.1456869	-1.1456869		
K6B3	-0.4322439	3.3289562	-0.4322439	-0.4322439		
S7B2	-0.3415675	-0.3415675	-0.3415675	-0.3415675		
C3B3	-0.2672858	-0.2672858	-0.2672858	-0.2672858		
R9B2	-0.4176403	-0.4176403	-0.4176403	-0.4176403		
TB22	-1.0574762	4.8800600	-1.0574762	-1.0574762		

```
TB23 -1.0876274 5.0370560 -1.0876274 -1.0876274
T5B2 -0.4862838 -0.4862838 -0.4862838 -0.4862838
T5B1 -0.5194111 -0.5194111 -0.5194111 -0.5194111
T5B3 -0.5091447 -0.5091447 -0.5091447 -0.5091447
TB21 -0.9675910 4.4350864 -0.9675910 -0.9675910
L8B1 -0.3614232 -0.3614232 -0.3614232 -0.3614232
TB11 -1.0916114 4.5178604 -1.0916114 -1.0916114
TB12 -0.8011263 -0.8011263 -0.8011263 -0.8011263
S7B3 -0.3484924 -0.3484924 -0.3484924 -0.3484924
C3B1 -0.2899390 4.6445349 -0.2899390 -0.2899390
S7B1 -0.2623762 -0.2623762 -0.2623762 -0.2623762
S4B2 -0.3394032 -0.3394032 -0.3394032 -0.3394032
```

```
# load necessary packages
# install.packages("glasso")
library(glasso)
# install.packages("qgraph")
library(qgraph)

# calculate the covariance matrix
cov_matrix <- cov(data_subset)
cov_matrix
```

	643311	4444760	687185	192795	164413
176468					
643311	0.63111786	-0.4845084	-0.060607931	0.123333259	0.63009539
4444760	-0.48450841	4.1201626	-1.099488832	-1.029576948	-1.13802002
687185	-0.06060793	-1.0994888	3.187831382	0.005397083	-0.37176143
192795	0.12333326	-1.0295769	0.005397083	0.860640701	0.68578581
164413	0.63009539	-1.1380200	-0.371761434	0.685785813	10.94770958
176468	-1.03532388	3.9670235	-2.291533489	-0.884317116	-3.45106979
255657	0.32140967	-1.1506849	0.037081388	0.391224984	0.42051766
181589	-0.89283599	2.0376490	0.563570325	-0.698532855	-3.79846100
219151	-0.28839638	2.2278574	-1.163651040	-0.835632050	-0.91516449
4425571	-0.12248916	0.1511638	-0.368565769	-0.052236320	0.08313653
	255657	181589	219151	4425571	
643311	0.32140967	-0.8928360	-0.2883964	-0.12248916	
4444760	-1.15068490	2.0376490	2.2278574	0.15116382	
687185	0.03708139	0.5635703	-1.1636510	-0.36856577	
192795	0.39122498	-0.6985329	-0.8356321	-0.05223632	

```

164413    0.42051766 -3.7984610 -0.9151645  0.08313653
176468   -1.64565695  4.3816342  1.8627034 -0.60740639
255657    2.43093182 -2.1655117 -0.8387100  0.10174094
181589   -2.16551169  6.8549149  0.4177501 -0.80612232
219151   -0.83870998  0.4177501  2.8317758  0.68555385
4425571   0.10174094 -0.8061223  0.6855539  1.05979881

```

```
# fit the Gaussian graphical model using graphical lasso
```

```
glasso_fit <- glasso(cov_matrix, rho = 0.1)
```

```
glasso_fit
```

```
$w
```

	[,1]	[,2]	[,3]	[,4]	[,5]
[,6]					
[1,]	0.73111786	-0.46181735	0.03939238	0.12682034	0.5621217
[2,]	-0.46181735	4.22016257	-1.19949104	-0.92957157	-1.2380190
[3,]	0.03939238	-1.19949104	3.28783138	0.10539706	-0.2717551
[4,]	0.12682034	-0.92957157	0.10539706	0.96064070	0.5857832
[5,]	0.56212168	-1.23801900	-0.27175508	0.58578317	11.0477096
[6,]	-0.93532293	3.86703274	-2.19153004	-0.84971972	-3.3510890
[7,]	0.26531610	-1.05067877	0.11718924	0.31464344	0.5205122
[8,]	-0.79283806	1.93765864	0.46357282	-0.59853442	-3.6984700
[9,]	-0.23126306	2.12785357	-1.06365224	-0.73563118	-0.8151735
[10,]	-0.02248898	0.08779529	-0.26856577	-0.07242801	0.1831340
	[,7]	[,8]	[,9]	[,10]	
[1,]	0.26531610	-0.7928381	-0.2312631	-0.02248898	
[2,]	-1.05067877	1.9376586	2.1278536	0.08779529	
[3,]	0.11718924	0.4635728	-1.0636522	-0.26856577	
[4,]	0.31464344	-0.5985344	-0.7356312	-0.07242801	
[5,]	0.52051217	-3.6984700	-0.8151735	0.18313404	
[6,]	-1.56601488	4.2816294	1.8576004	-0.50738917	
[7,]	2.53093182	-2.0655103	-0.7387166	0.09034248	
[8,]	-2.06551026	6.9549149	0.5177495	-0.70612297	
[9,]	-0.73871662	0.5177495	2.9317758	0.58555585	
[10,]	0.09034248	-0.7061230	0.5855558	1.15979881	

```
$wi
```

	[,1]	[,2]	[,3]	[,4]	[,5]
[,6]					
[1,]	1.70646426	0.00000000	0.10619439	0.00000000	0.00000000

```

0.18569604
[2,] 0.00000000 0.60208630 -0.01112989 0.20617035 -0.03146500 -
0.20848204
[3,] 0.10620161 -0.01115004 0.50643817 0.09938869 0.03107650
0.22757833
[4,] 0.00000000 0.20620921 0.09938915 1.44405510 -0.02346325
0.00000000
[5,] 0.00000000 -0.03146834 0.03107706 -0.02346322 0.12045880
0.04400977
[6,] 0.18570671 -0.20851458 0.22757608 0.00000000 0.04400672
0.40384552
[7,] 0.00000000 0.01529398 0.00000000 0.00000000 0.04356024
0.00000000
[8,] 0.09297855 -0.01374790 -0.13104525 0.03009555 0.05460553 -
0.14557999
[9,] 0.00000000 -0.25965514 0.09398462 0.23691995 0.03062859
0.00000000
[10,] 0.19553382 0.00000000 0.09379811 0.00000000 0.02273115
0.15317650

```

```

      [,7]      [,8]      [,9]     [,10]
[1,] 0.00000000 0.09298378 0.00000000 0.19552876
[2,] 0.01527024 -0.01376973 -0.25964480 0.00000000
[3,] 0.00000000 -0.13104520 0.09398372 0.09379879
[4,] 0.00000000 0.03009558 0.23692082 0.00000000
[5,] 0.04356436 0.05460625 0.03062606 0.02273604
[6,] 0.00000000 -0.14557885 0.00000000 0.15317156
[7,] 0.57305413 0.18065049 0.11352134 0.00000000
[8,] 0.18065040 0.34483449 0.05540745 0.06997103
[9,] 0.11351761 0.05540500 0.70641800 -0.28038463
[10,] 0.00000000 0.06996858 -0.28038699 1.13531005

```

```
$loglik
```

```
[1] NA
```

```
$errflag
```

```
[1] 0
```

```
$approx
```

```
[1] FALSE
```

```
$del
```

```
[1] 9.554498e-05
```

```
$niter
```

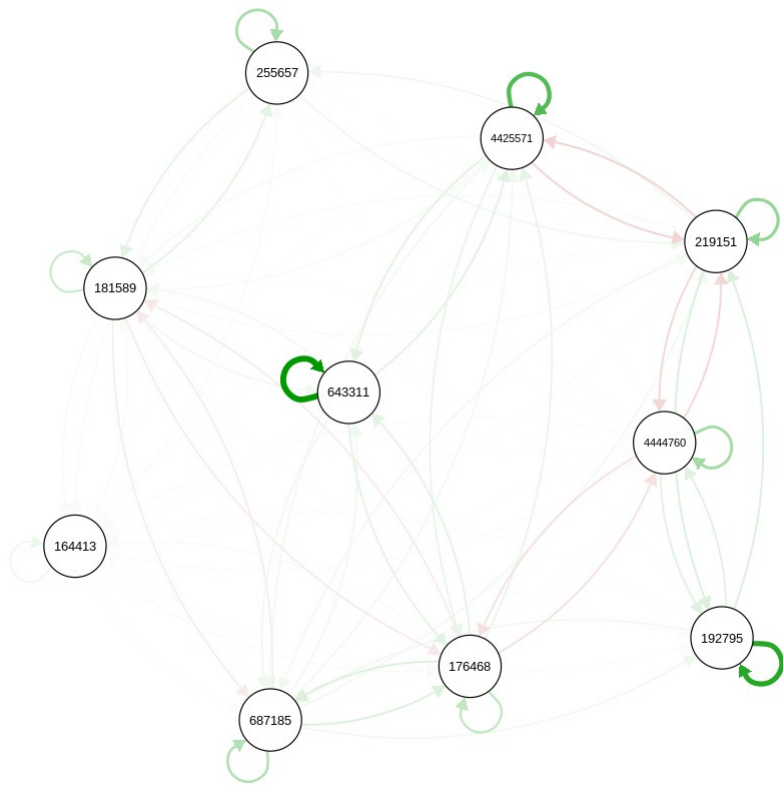
```
[1] 3
```

```
# visualize the network
```

```

qgraph(glasso_fit$wi, labels = colnames(data_subset), layout =
"spring",
      vsize = 7, esize = 5)

```



Interpretation:

Strong Associations:

- Microbes identified by OTUs 643311, 192795, and 442571 have thicker green loops, indicating strong positive self-associations.
- There is a strong positive association between 643311 and 442571, indicated by the thick green edge connecting them.

Weak Associations:

- Several nodes are connected by thin green and red edges, indicating weaker positive and negative associations, respectively.

Negative Associations:

- Some red edges indicate negative associations, suggesting that as the abundance of one microbe increases, the abundance of the connected microbe tends to decrease.