

File

Input data

Reference gens:	5	Show Table
Target genes:	7	Show Table
Line models:	26	Show List
Quantitative data:	Loaded!	Remove Show Table

Parameters

Normalization algorithm: Normfinder

Statistical model: Kruskal-Wallis/Kolmogorov-Smirnov, post: Dunn's test confidence: 0.05

Remove repetitions: 1

Single sample repetition: 3

Select best remove for model

Run calculations

Results summary

Coherence score list

B4GALT1	1.00
B4GALT2	1.00
B4GALT3	1.00
B4GALT4	0.90
B4GALT5	1.00
B4GALT6	0.90
B4GALT7	1.00
Average	0.97

Show best references

Show RQ values

Show p-values

Export results

Export graphs