Breast cancer, gene expression analysis in 'black or african-american' and 'white' cohorts, in PAM50 subtypes

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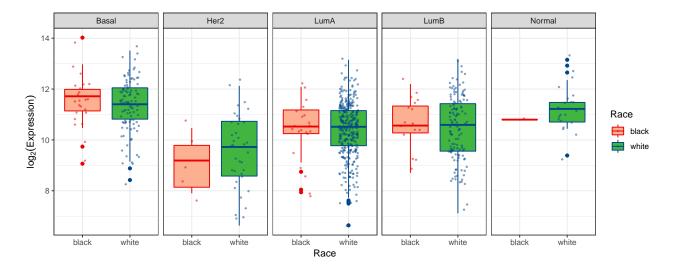
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selected_genes <- c("MYC")</pre>

Number of samples per Race and PAM50 subtypes

	Basal	Her2	LumA	LumB	Normal
black	23	5	21	16	1
\mathbf{white}	101	37	344	128	21

Expression analysis in 'black or african-american' and 'white' cohorts, in PAM50 subtypes



T-test, differences between races in individual subtypes

For each PAM50 subtypes, compare expression between EA and AA cohorts. Only p-values are shown

In Basal

[1] 0.4303541

In Her2

[1] 0.4020254

In LumA

[1] 0.7885612

In LumB

[1] 0.5600073

In Normal

Not enough observations

Two-way Anova

Race and PAM50 effects across all subtypes.

Across all subtypes, "Race" differences are not significant

Across all subtypes, "PAM50" differences are significant

Dunnett post-hoc test, differences between PAM50 subtypes separately in European-American and African-American cohorts

In EA cohort ("white" in TCGA classification)

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Dunnett's test for comparing several treatments with a control : 95\% family-wise confidence level
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\$Basal

```
diff lwr.ci upr.ci pval
Her2-Basal -1.78266220 -2.3094421 -1.2558823 2.2e-15 ***
LumA-Basal -0.95144106 -1.2616764 -0.6412057 3.5e-13 ***
LumB-Basal -0.87883631 -1.2436770 -0.5139956 1.7e-08 ***
Normal-Basal -0.06088636 -0.7183335 0.5965607 0.9983
```

Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1

In AA cohort ("black or african american" in TCGA classification)

Dunnett's test for comparing several treatments with a control : 95% family-wise confidence level

\$Basal

 diff
 lwr.ci
 upr.ci
 pval

 Her2-Basal
 -2.4586488
 -3.797190
 -1.12010779
 6.8e-05

 LumA-Basal
 -1.2096290
 -2.028385
 -0.39087331
 0.0015
 **

 LumB-Basal
 -0.9059245
 -1.789025
 -0.02282427
 0.0422
 *

 Normal-Basal
 -0.7568016
 -3.527845
 2.01424154
 0.9196

Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1