

Pairwise comparisons using t tests with non-pooled SD

data: DRM_number and Level_hierarchy

	0	1	2
0			
1	0.09	-	-
2	0.09	0.30	-
3	0.09	0.20	0.45

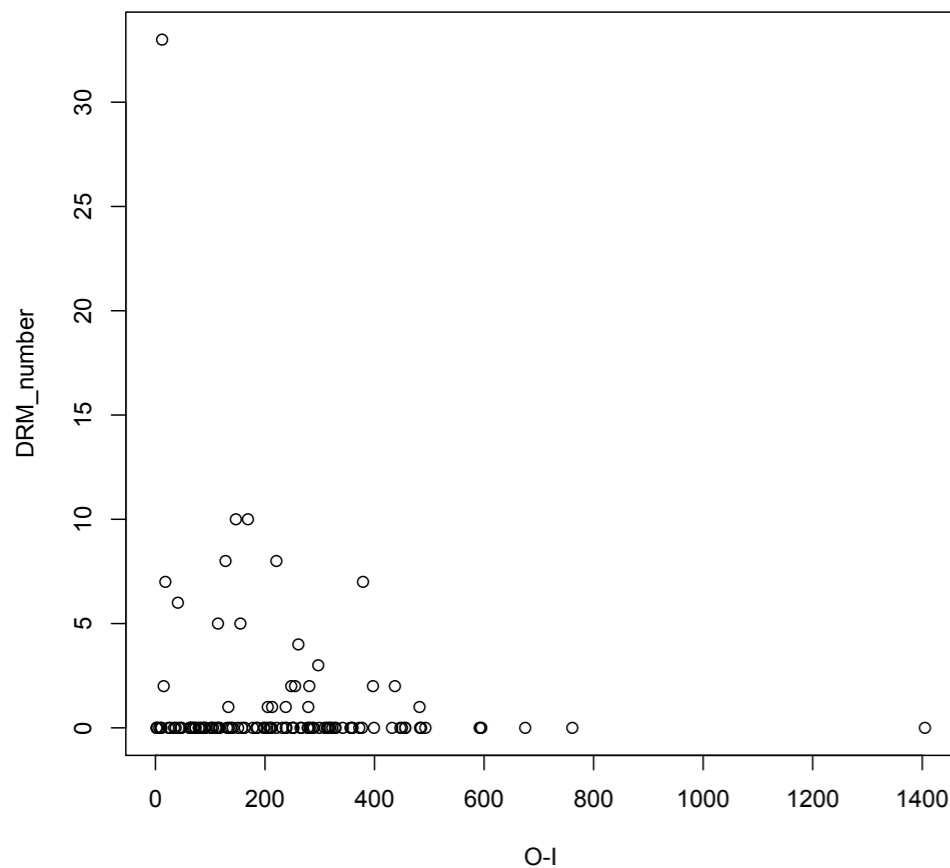
P value adjustment method: BH

Pairwise comparisons using Wilcoxon rank sum test

data: DRM_number and Level_hierarchy

	0	1	2
0			
1	0.25	-	-
2	0.25	0.46	-
3	0.35	0.25	0.46

P value adjustment method: BH



Call:

lm(formula = DRM_number ~ OI, model = T)

Residuals:

	Min	1Q	Median	3Q	Max
Residuals	-1.6493	-1.3735	-0.9278	-0.3363	31.3768

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.654551	0.515690	3.208	0.00172 **
OI	-0.002614	0.001699	-1.539	0.12655

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

Residual standard error: 3.576 on 117 degrees of freedom

Multiple R-squared: 0.01984, Adjusted R-squared: 0.01146

F-statistic: 2.368 on 1 and 117 DF, p-value: 0.1266