Group Comparison on spatial DGMM segment means for file:

defaultname

properties	values
Number of m/z features	388
Range of m/z values	602.32 – 3181.63
Number of pixels	40
Range of x coordinates	2 – 74
Range of y coordinates	3 – 56
Range of intensities	0 – 8.14
Number of NA intensities	0
Number of Inf intensities	0
Number of duplicated coordinates	0

Distribution of the samples
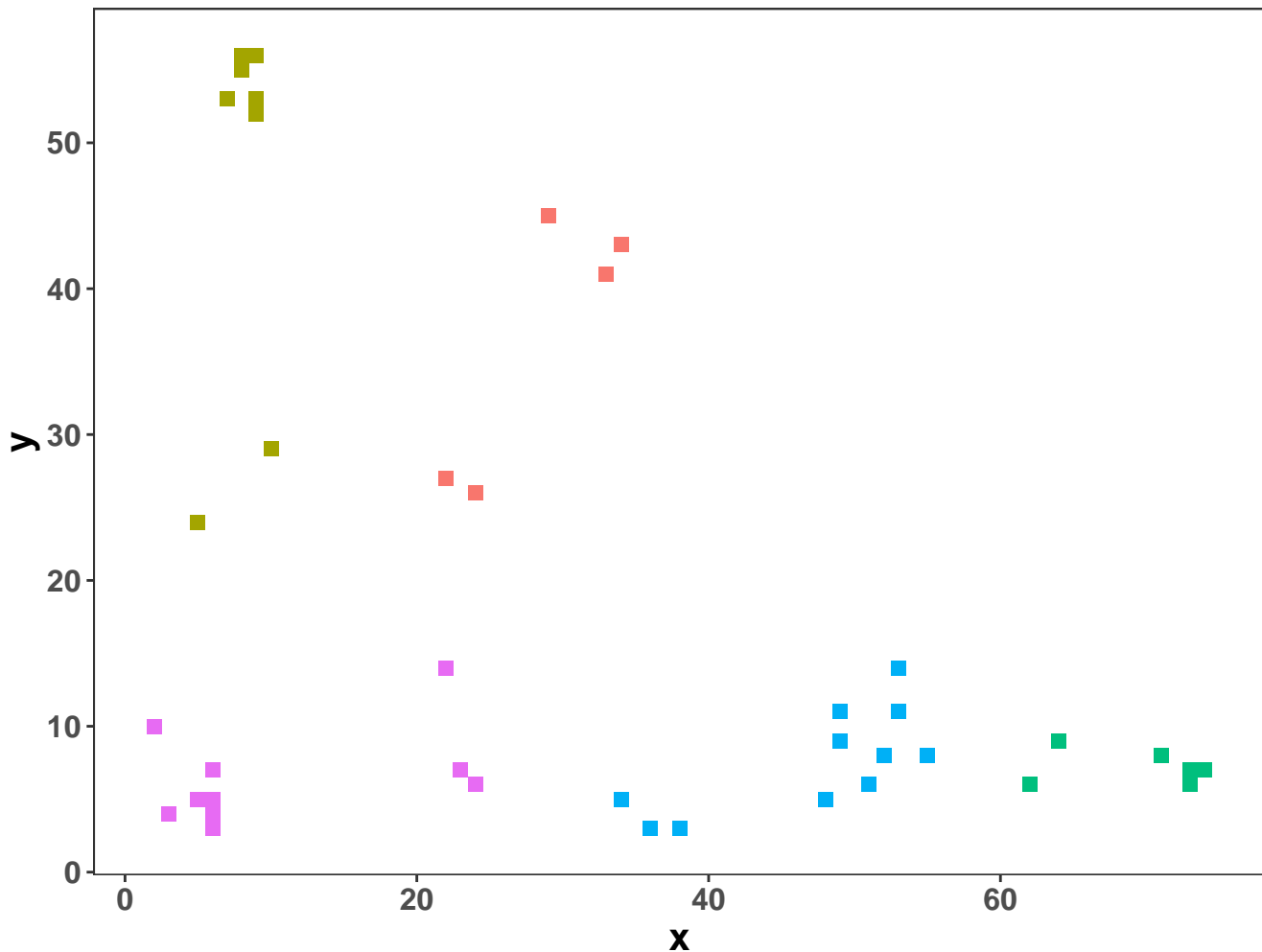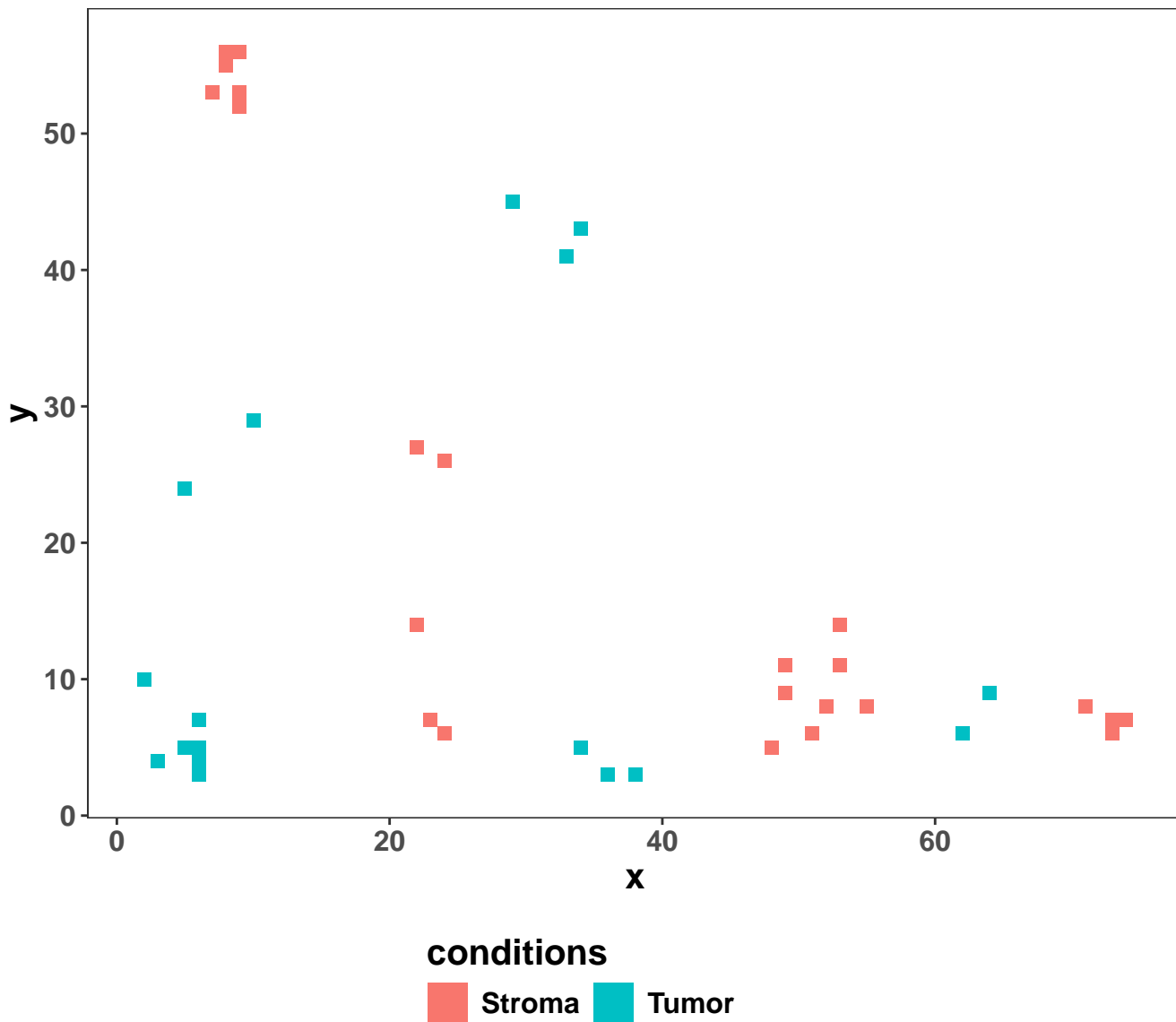
sample_id
- TMA_394
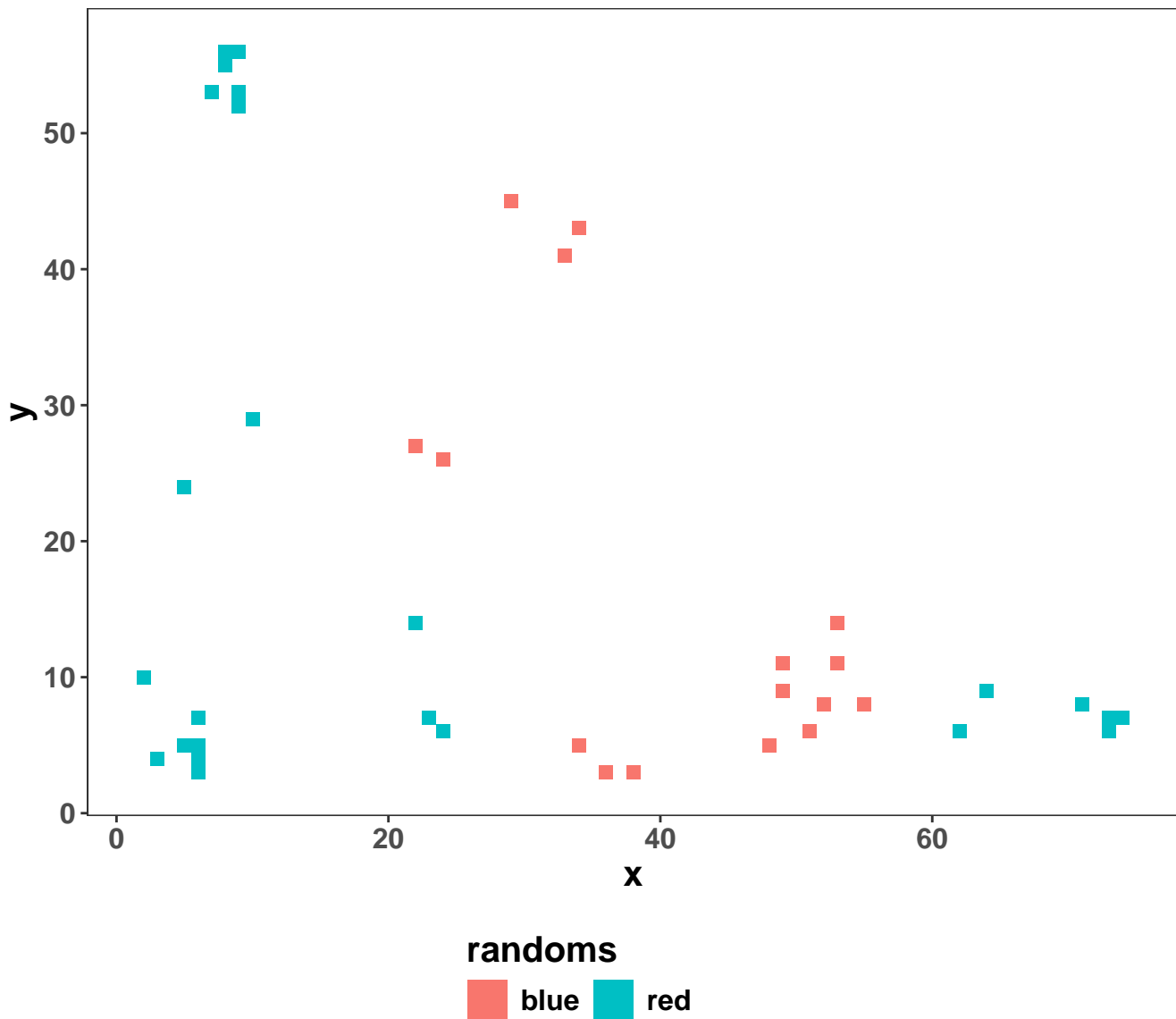- TMA_397
- TMA_398
- TMA_400
- TMA_402

Distribution of the Conditions

Distribution of the Random Effect

No m/z feature with adjusted p value below the fdr filter found.
Therefore, the selection of interesting m/z features for the boxplots is not possible.
vith 'Generate boxplots and results table for all m/z features?' set to 'YES' in order to generate
for all m/z features or repeat with less strict fdr value.