

# PDAC gene expression analysis

Richard Moffitt

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## Setup

### load required libraries

```
## Setting options('download.file.method.GEOquery'='auto')

## Setting options('GEOquery.inmemory.gpl'=FALSE)

##

##

##

## Type 'citation("pROC")' for a citation.
```

```
##
## Attaching package: 'pROC'

## The following objects are masked from 'package:stats':
##
##   cov, smooth, var

##
## Attaching package: 'plyr'

## The following object is masked from 'package:ggpubr':
##
##   mutate
```

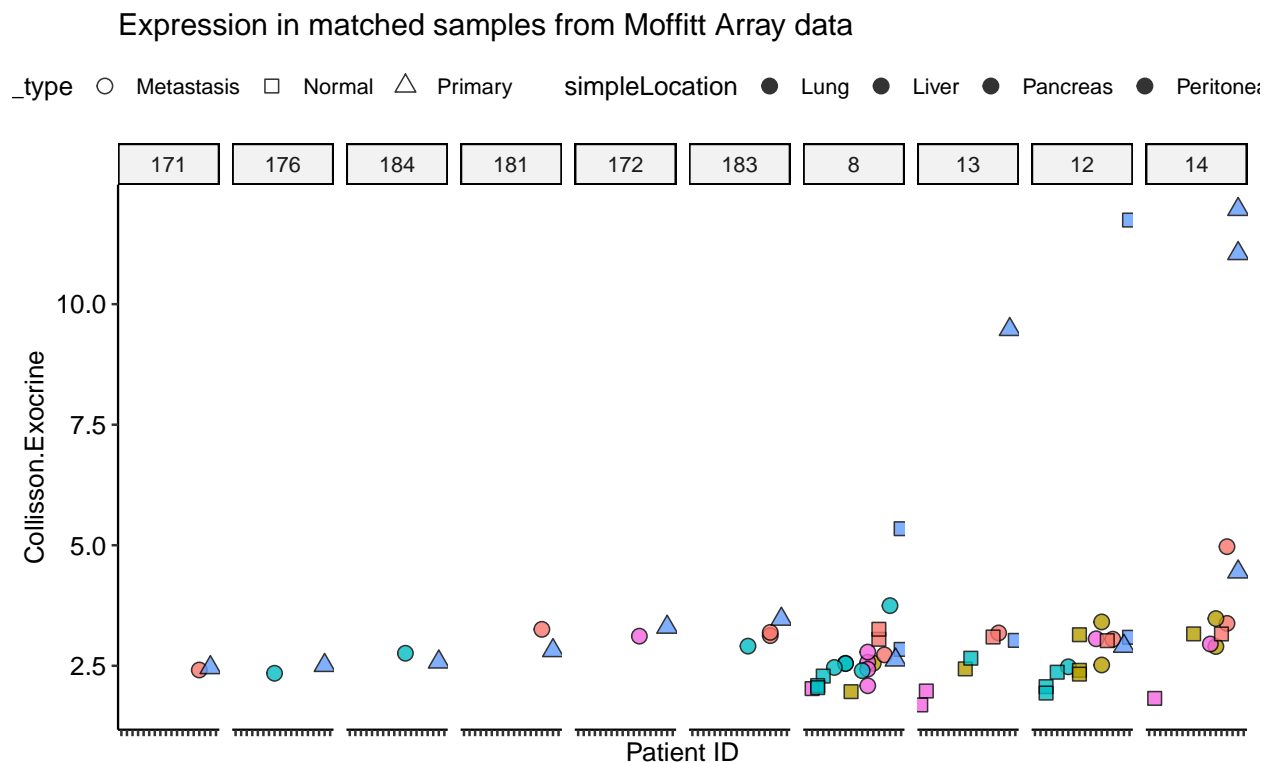
## Preprocessing of genesets

```
## Registered S3 method overwritten by 'GGally':
##   method from
##   +.gg      ggplot2
```

## Match samples from same source

## Results

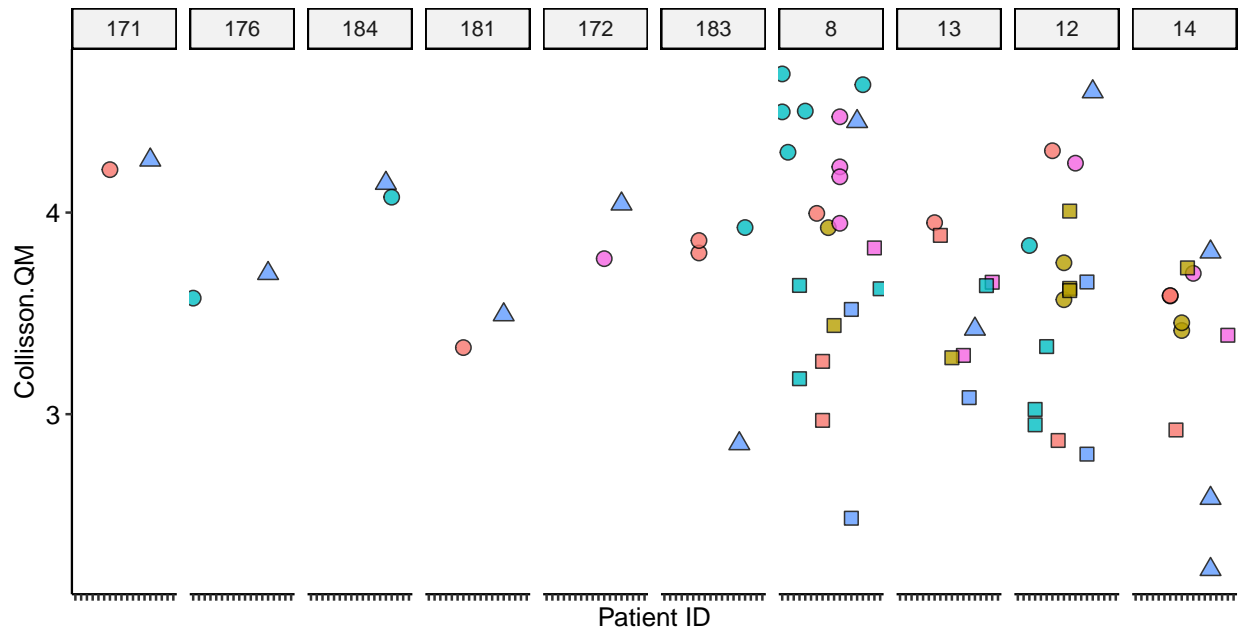
### Exocrine signature by matched samples



## QM signature by matched samples

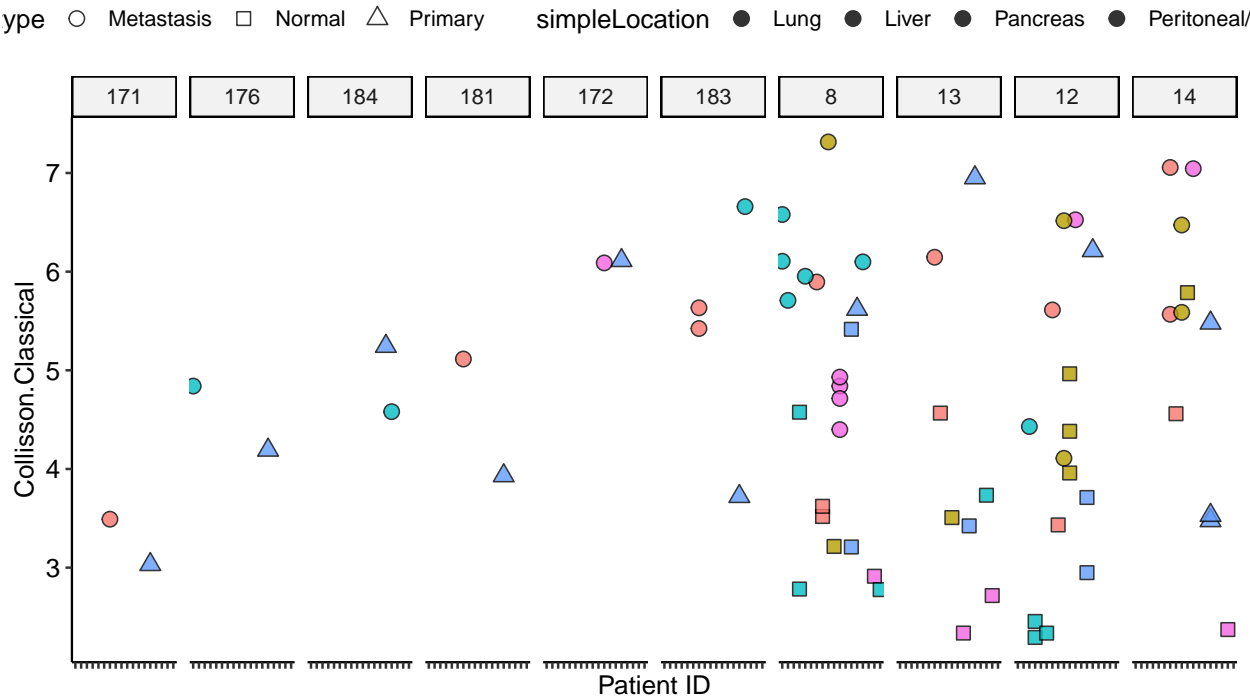
Expression in matched samples from Moffitt Array data

type ○ Metastasis □ Normal △ Primary simpleLocation ● Lung ● Liver ● Pancreas ● Peritoneal/



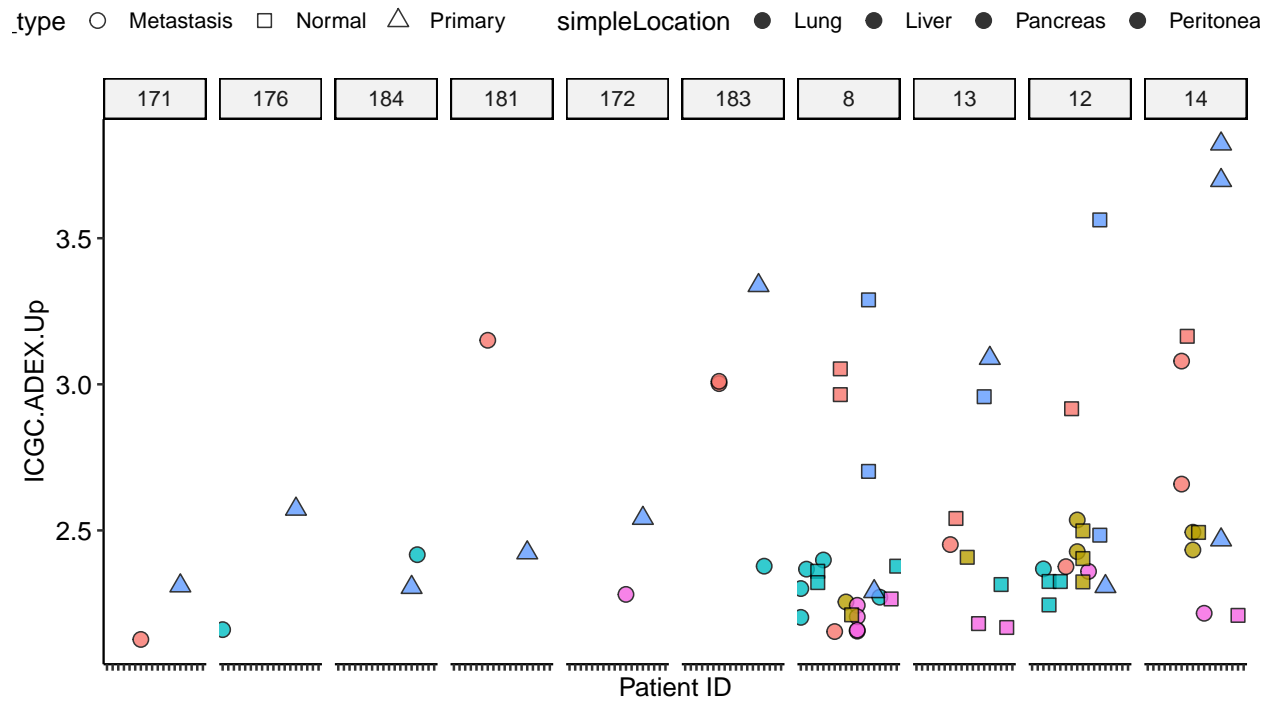
Collisson Classical signature by matched samples

Expression in matched samples from Moffitt Array data



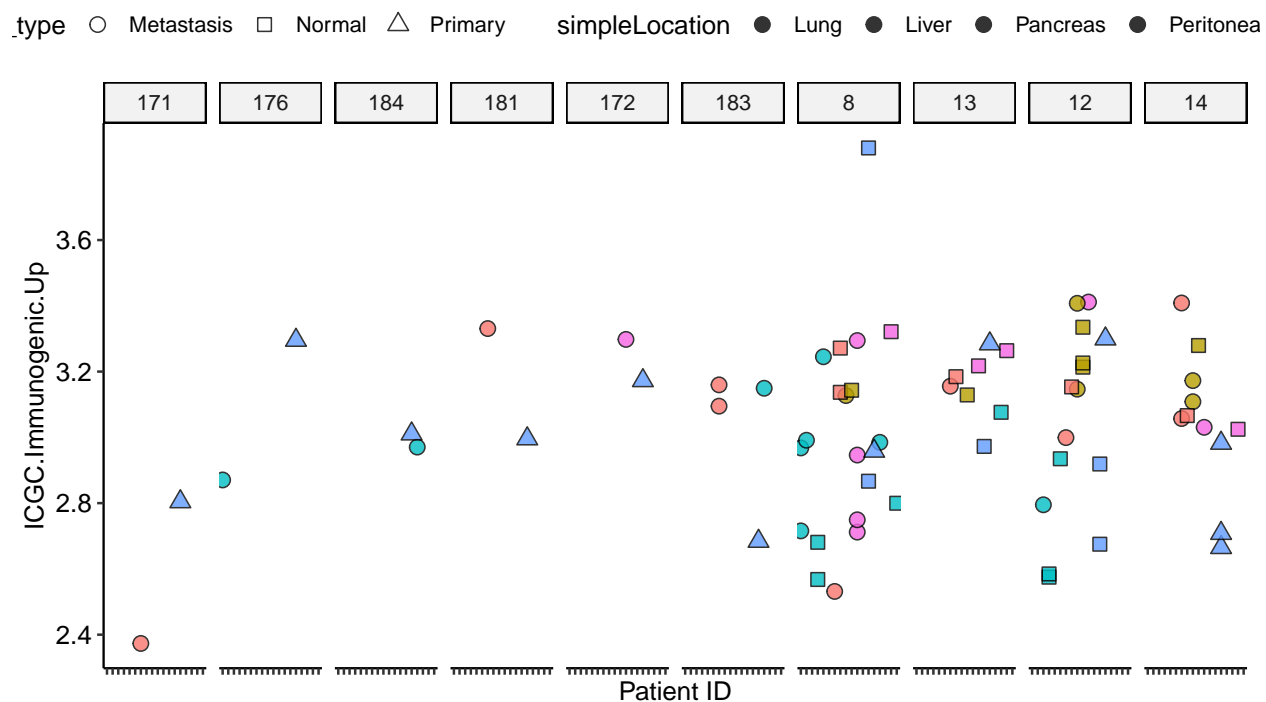
### ADEX signature by matched samples

### Expression in matched samples from Moffitt Array data

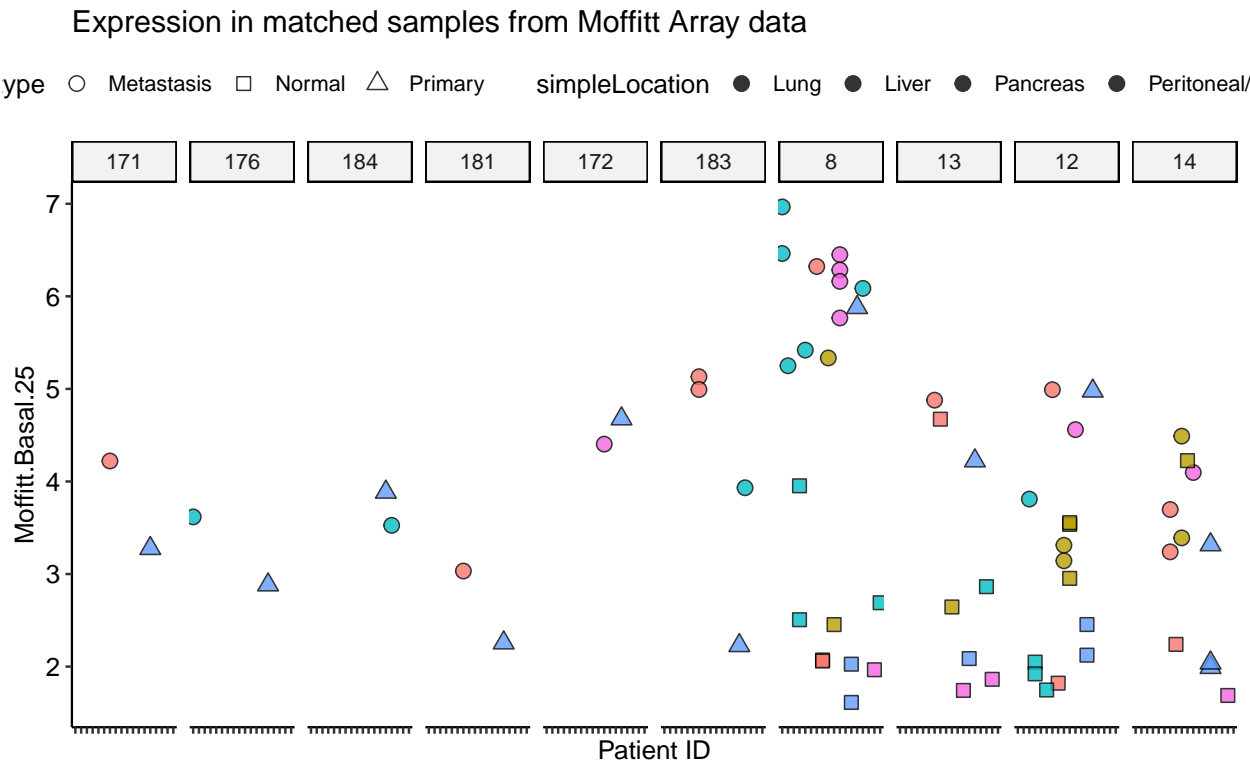


## Immunogenic signature by matched samples

Expression in matched samples from Moffitt Array data



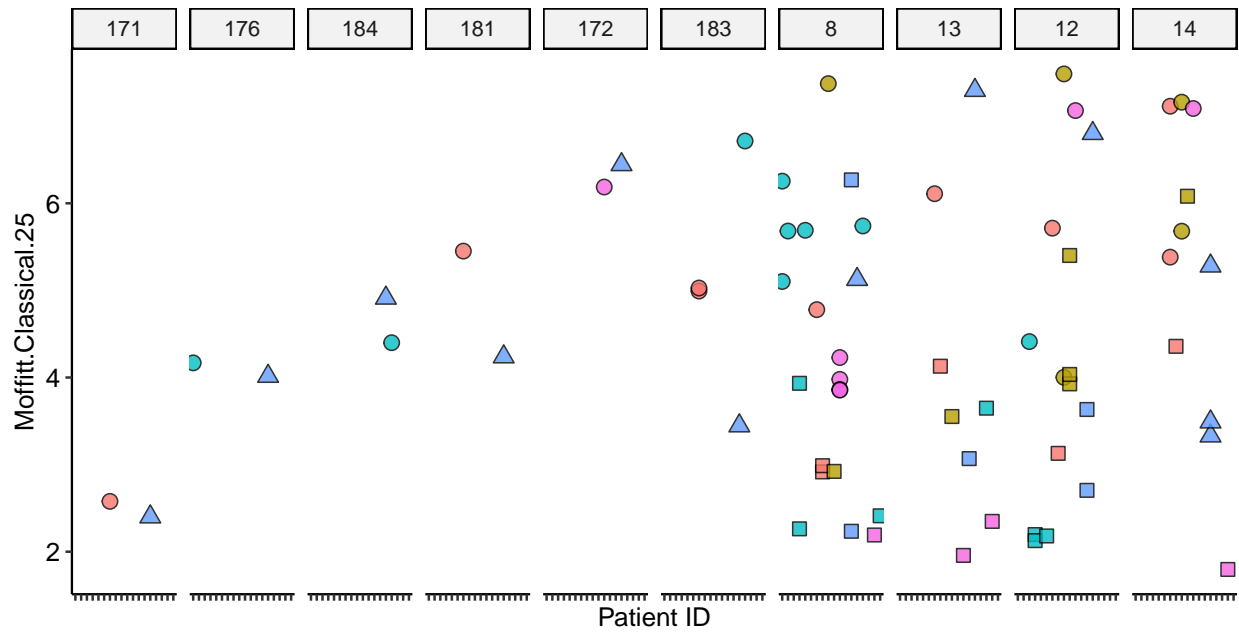
Basal-like signature by matched samples



## Classical signature by matched samples

### Expression in matched samples from Moffitt Array data

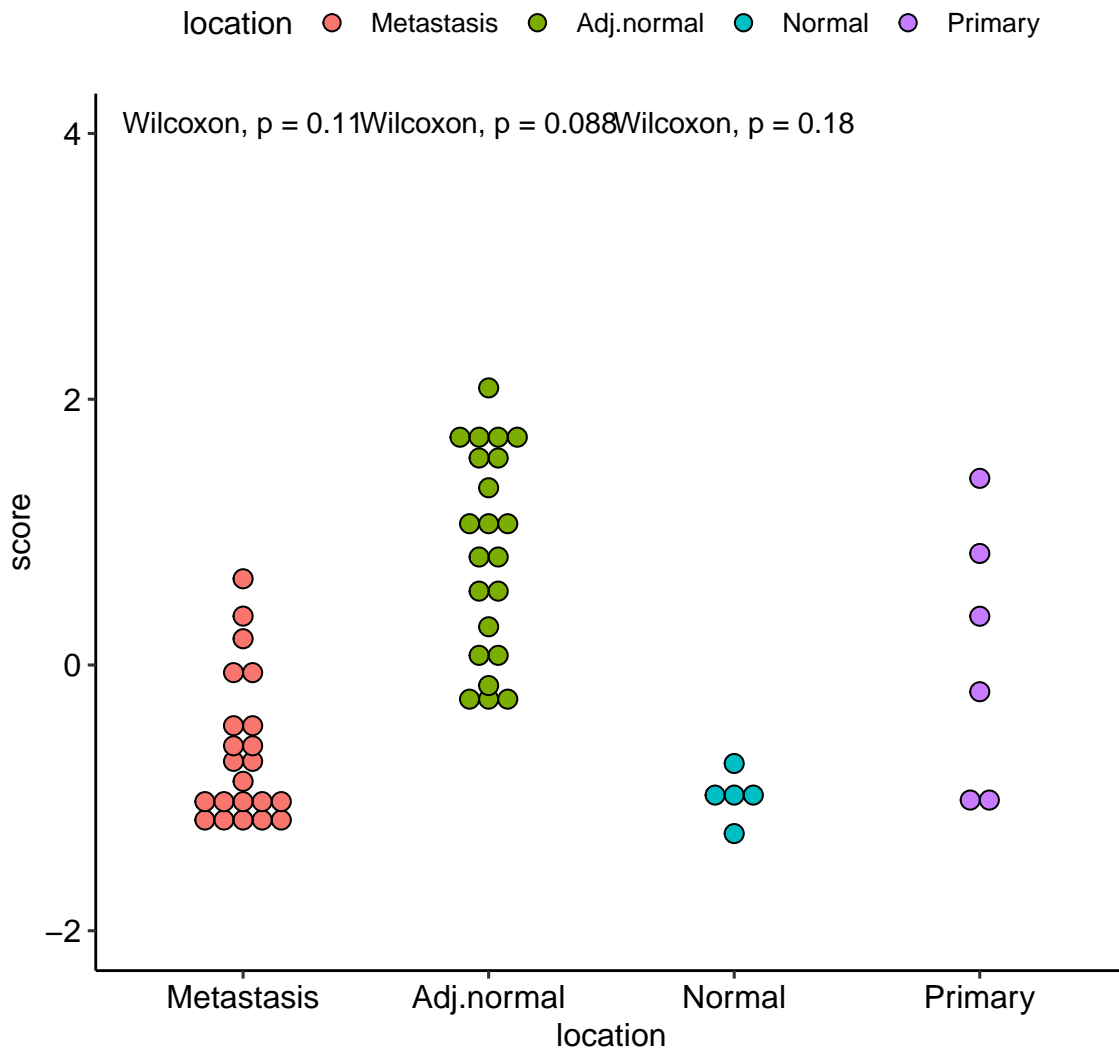
type ○ Metastasis □ Normal △ Primary      simpleLocation ● Lung ● Liver ● Pancreas ● Peritoneal/



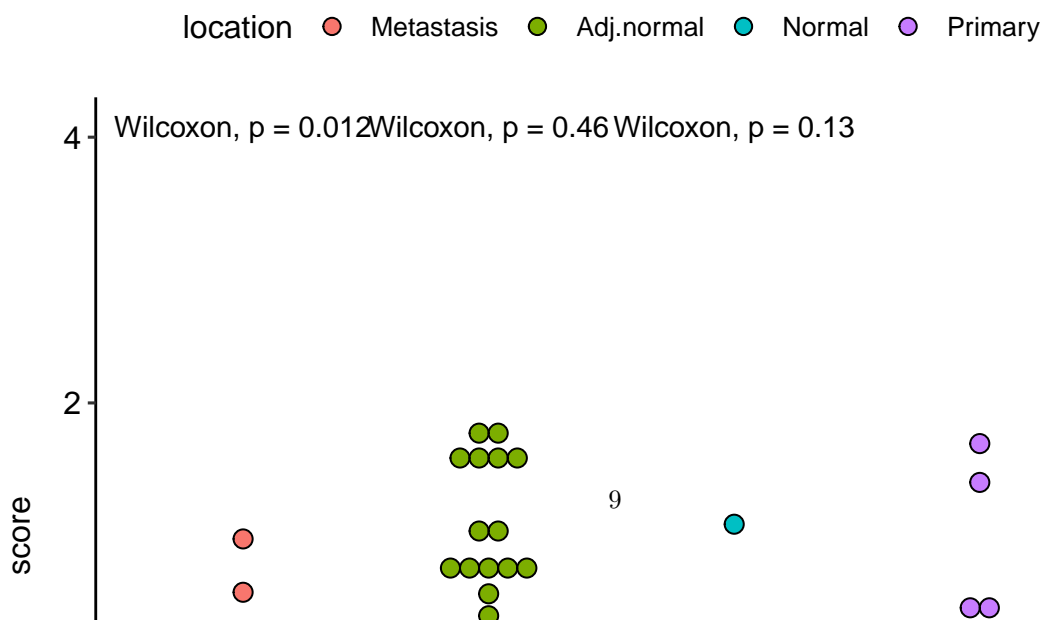


### Simplified patient matched sample plots

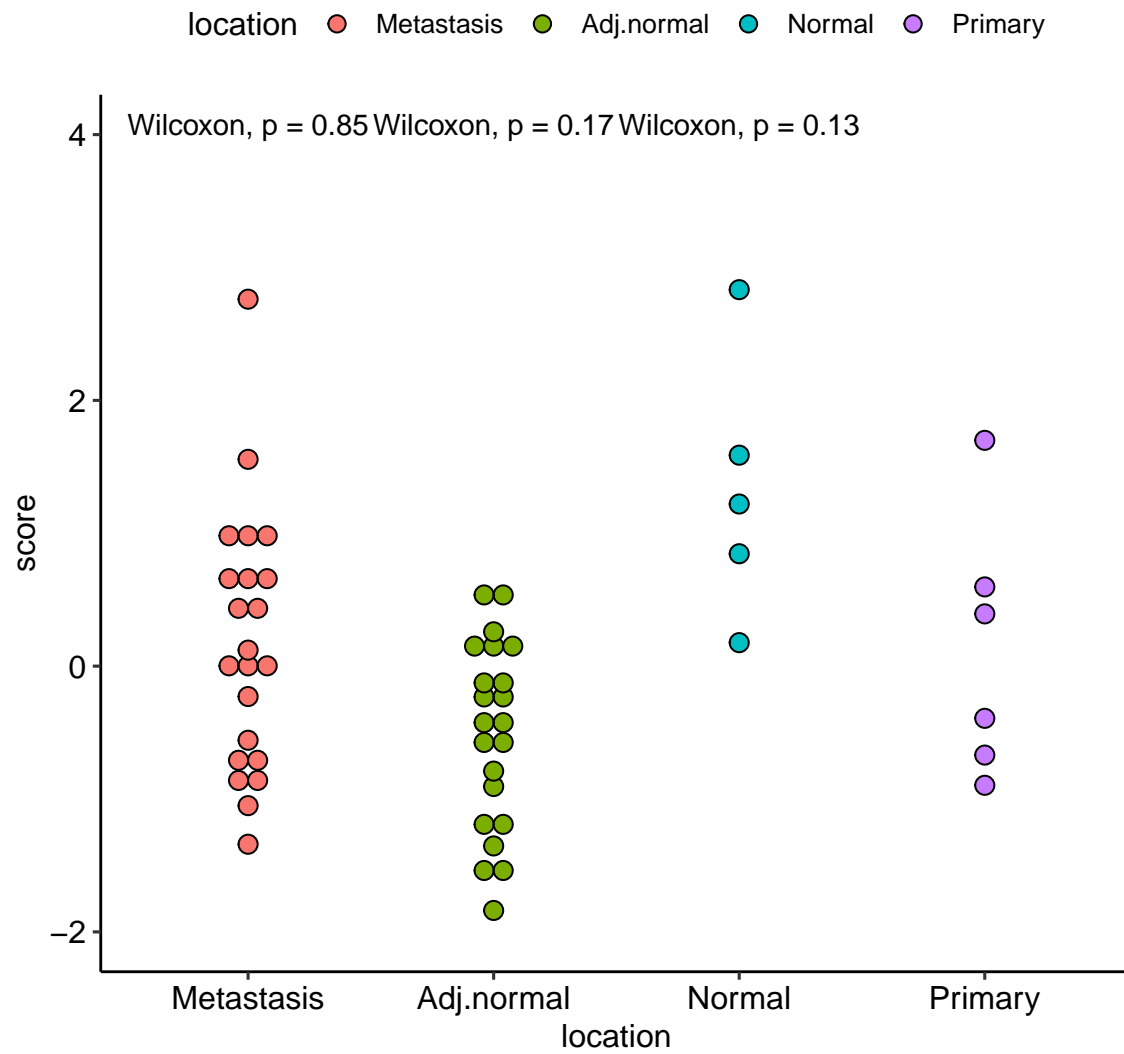
## Moffitt.Basal.25 score



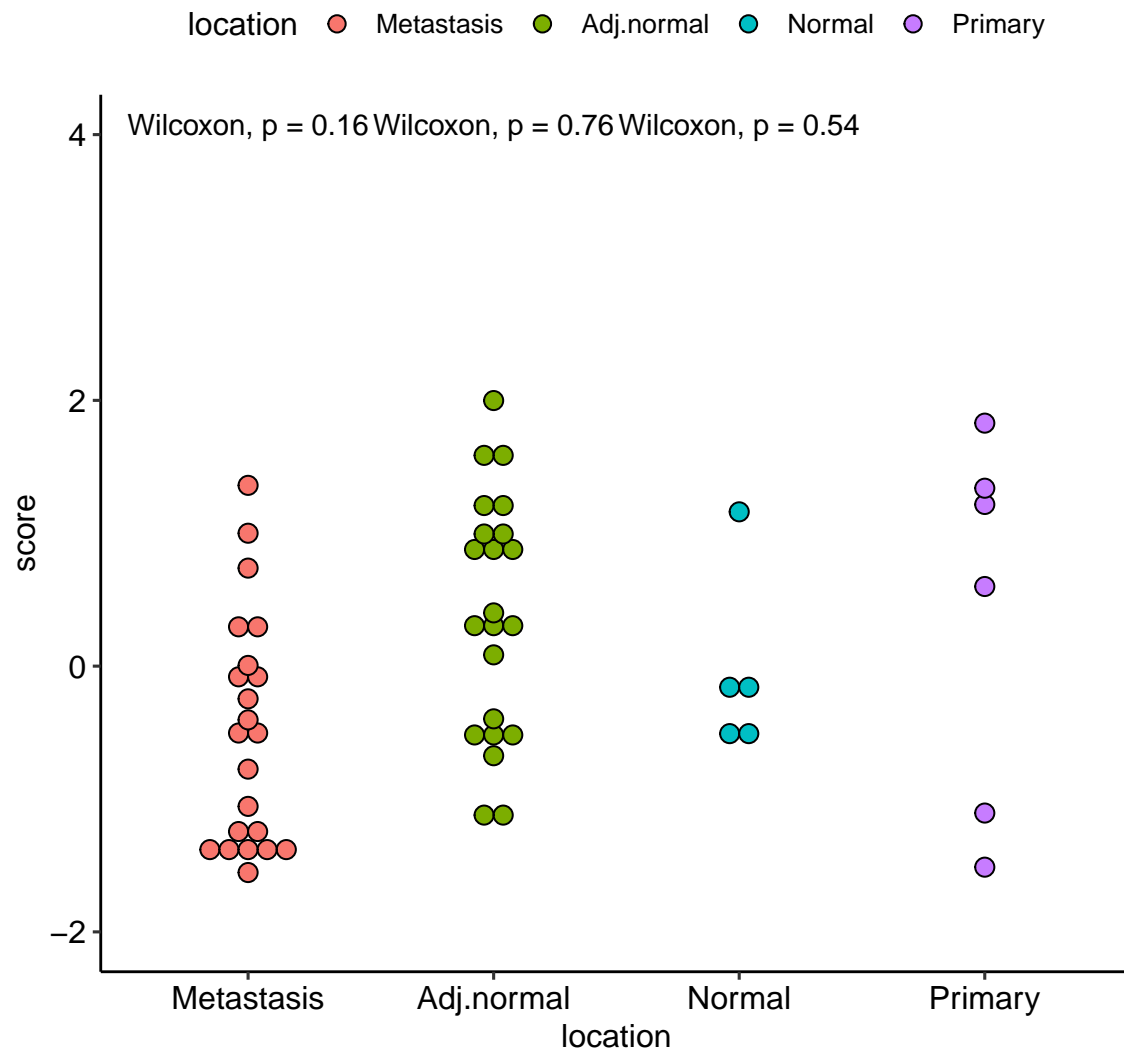
## Moffitt.Classical.25 score



## Moffitt.Normal.25 score

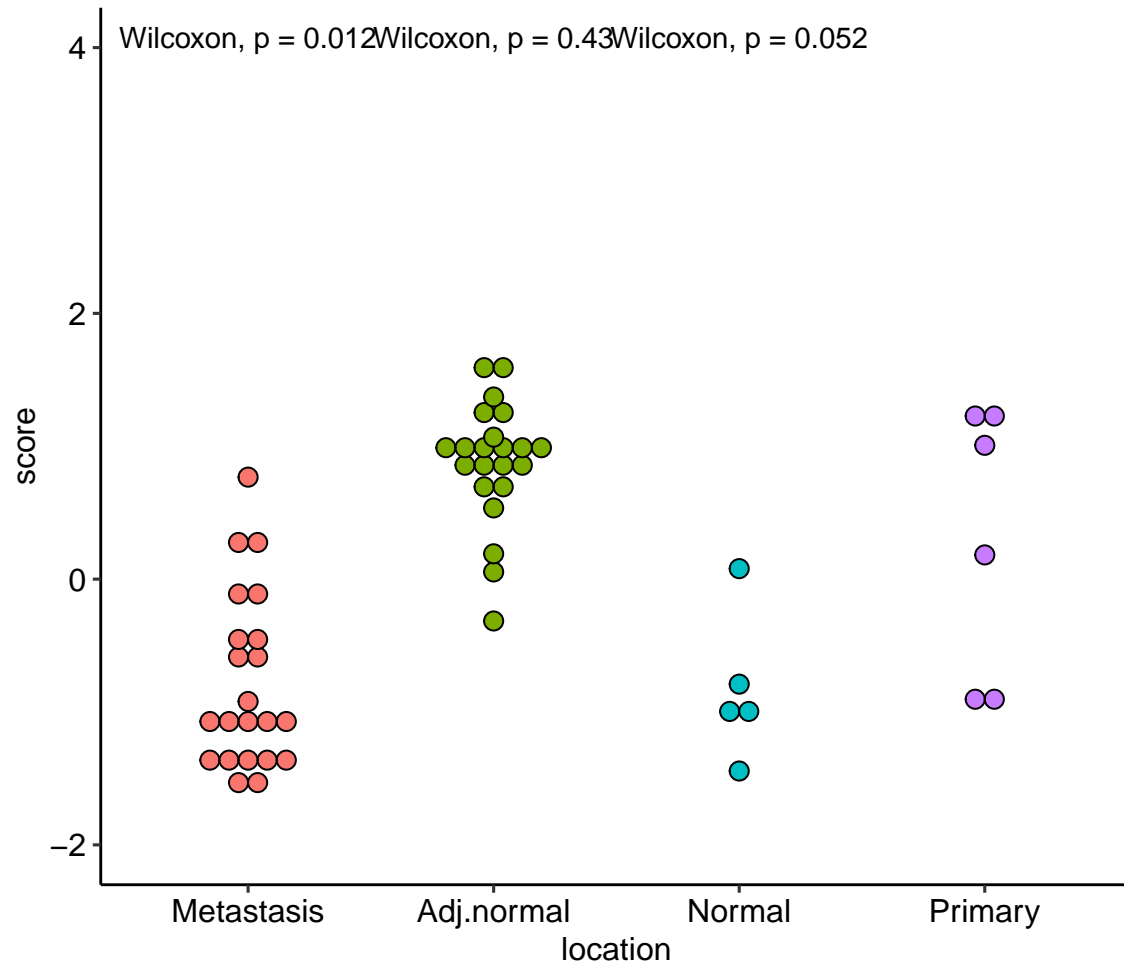


## Moffitt.Activated.25 score

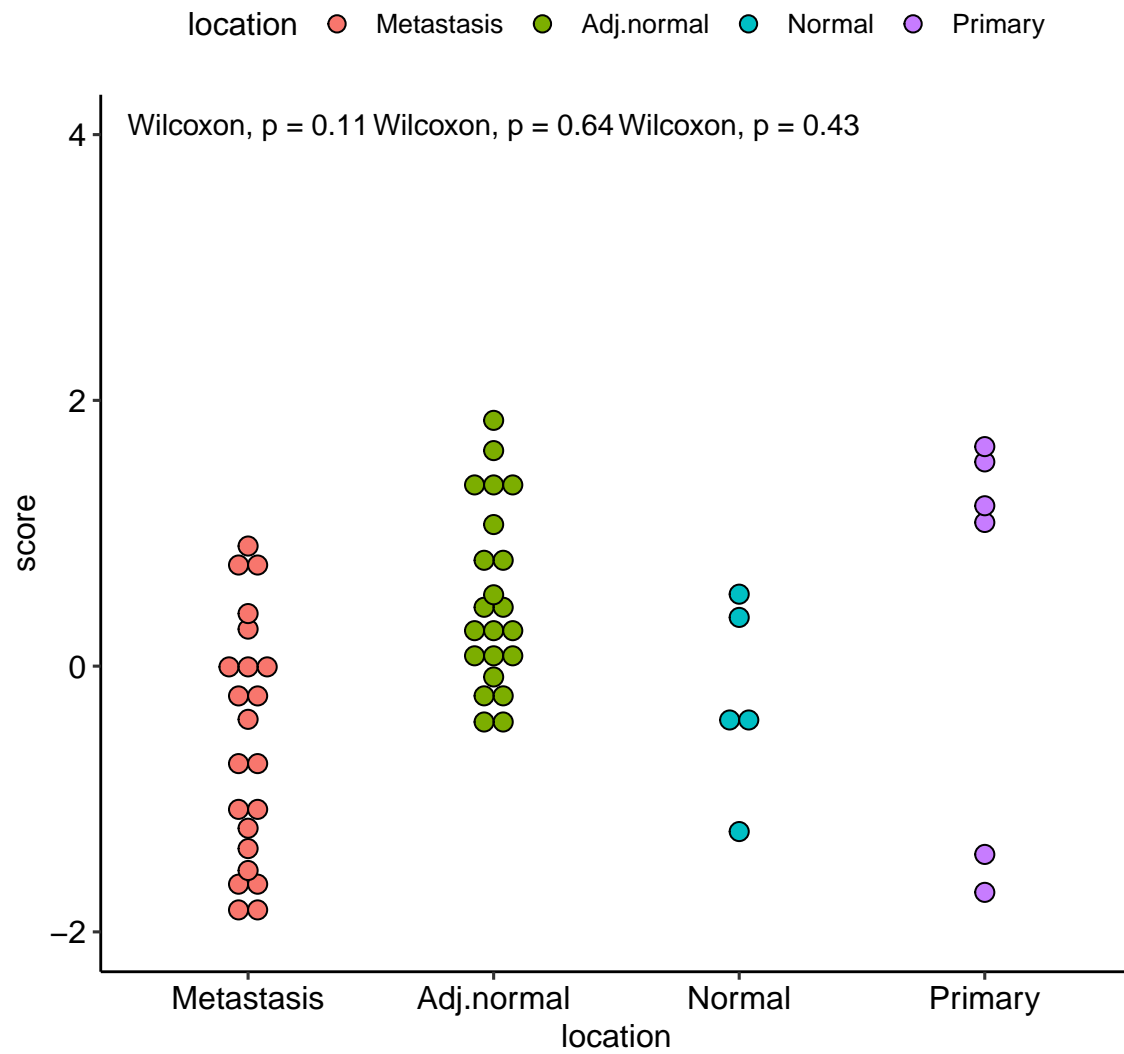


## Moffitt.Tumor score

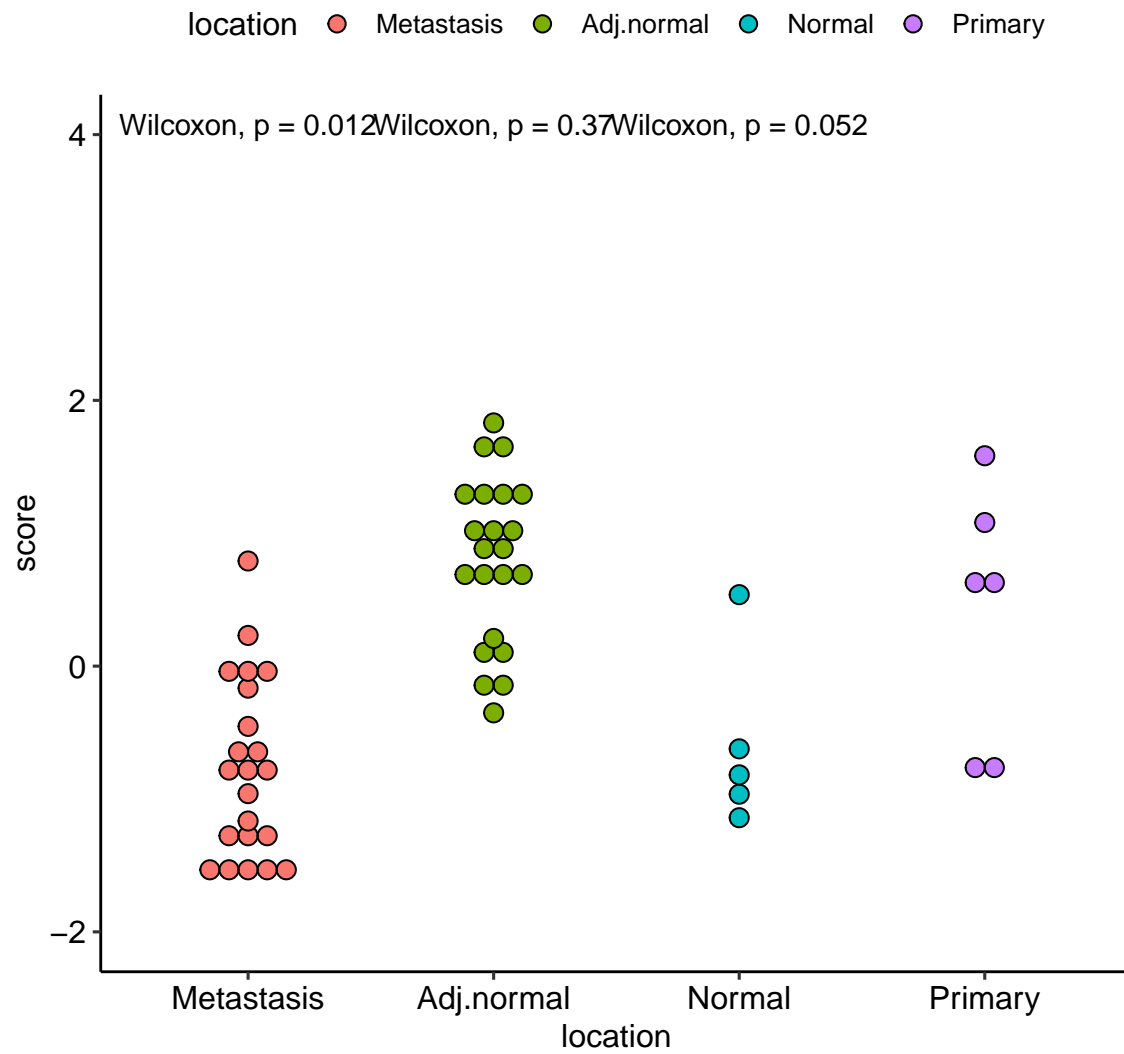
location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



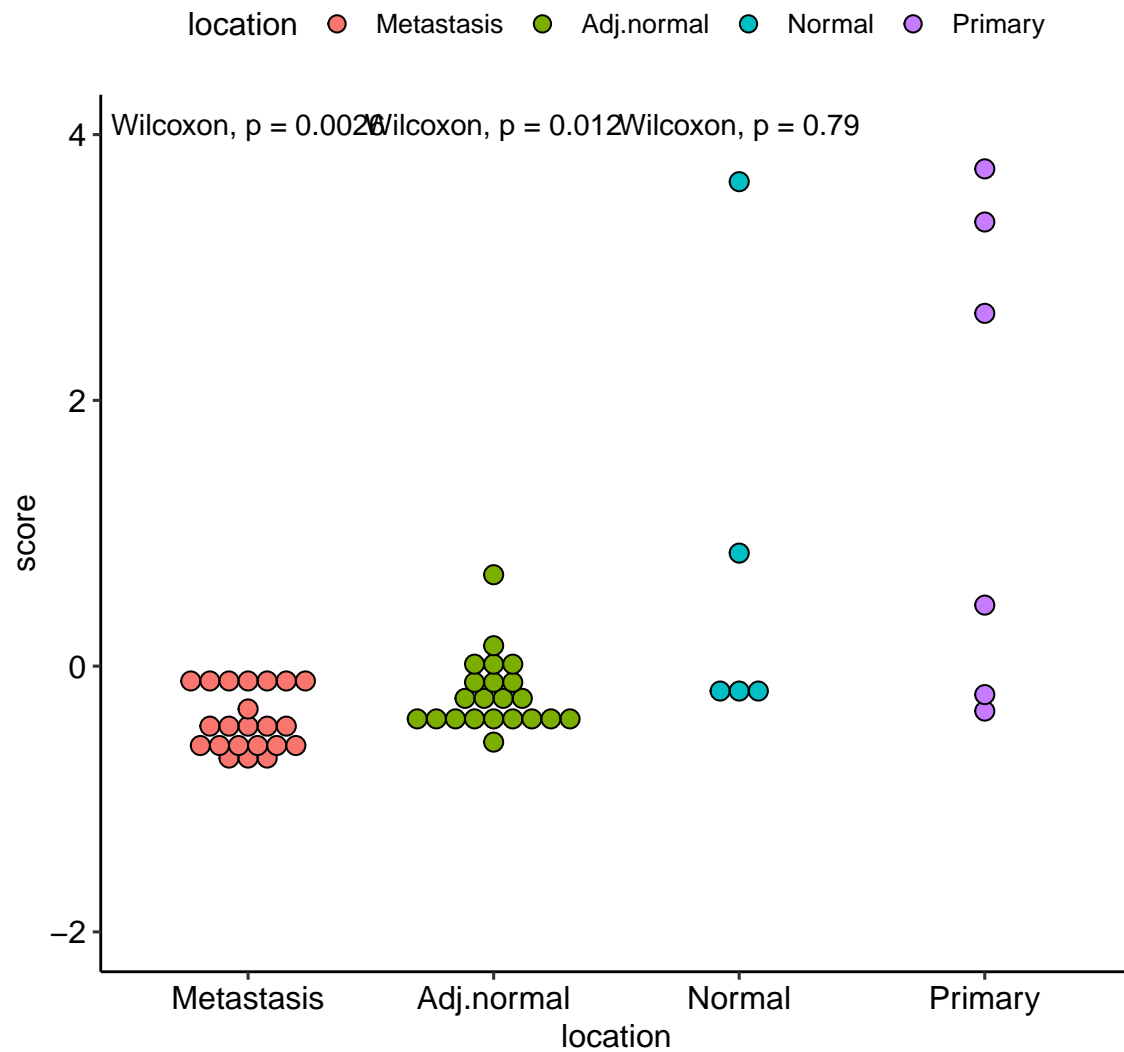
### Puleo.Centroids score



## Collisson.Classical score



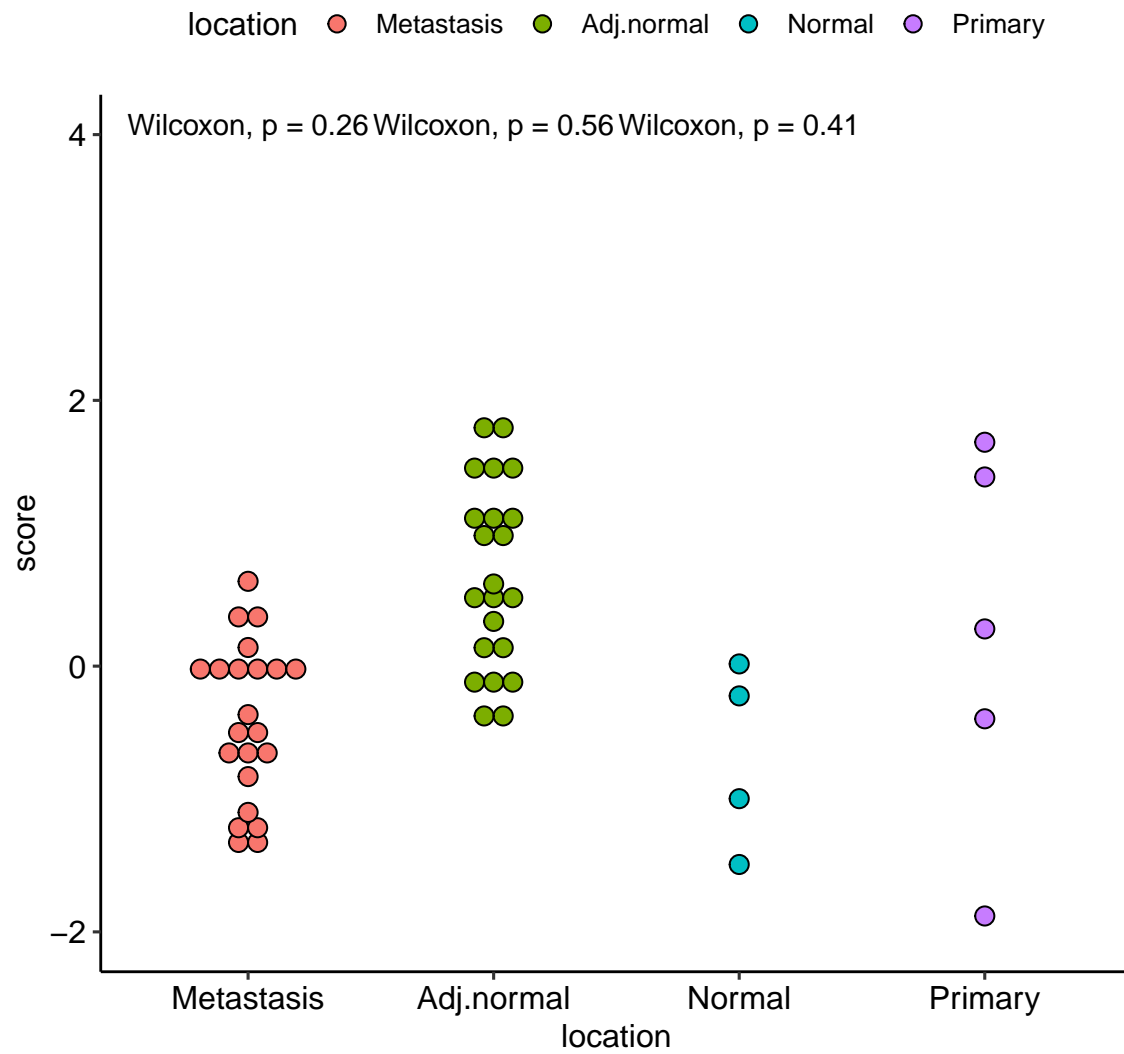
## Collisson.Exocrine score



```
## Warning: Removed 2 rows containing non-finite values (stat_bindot).
```

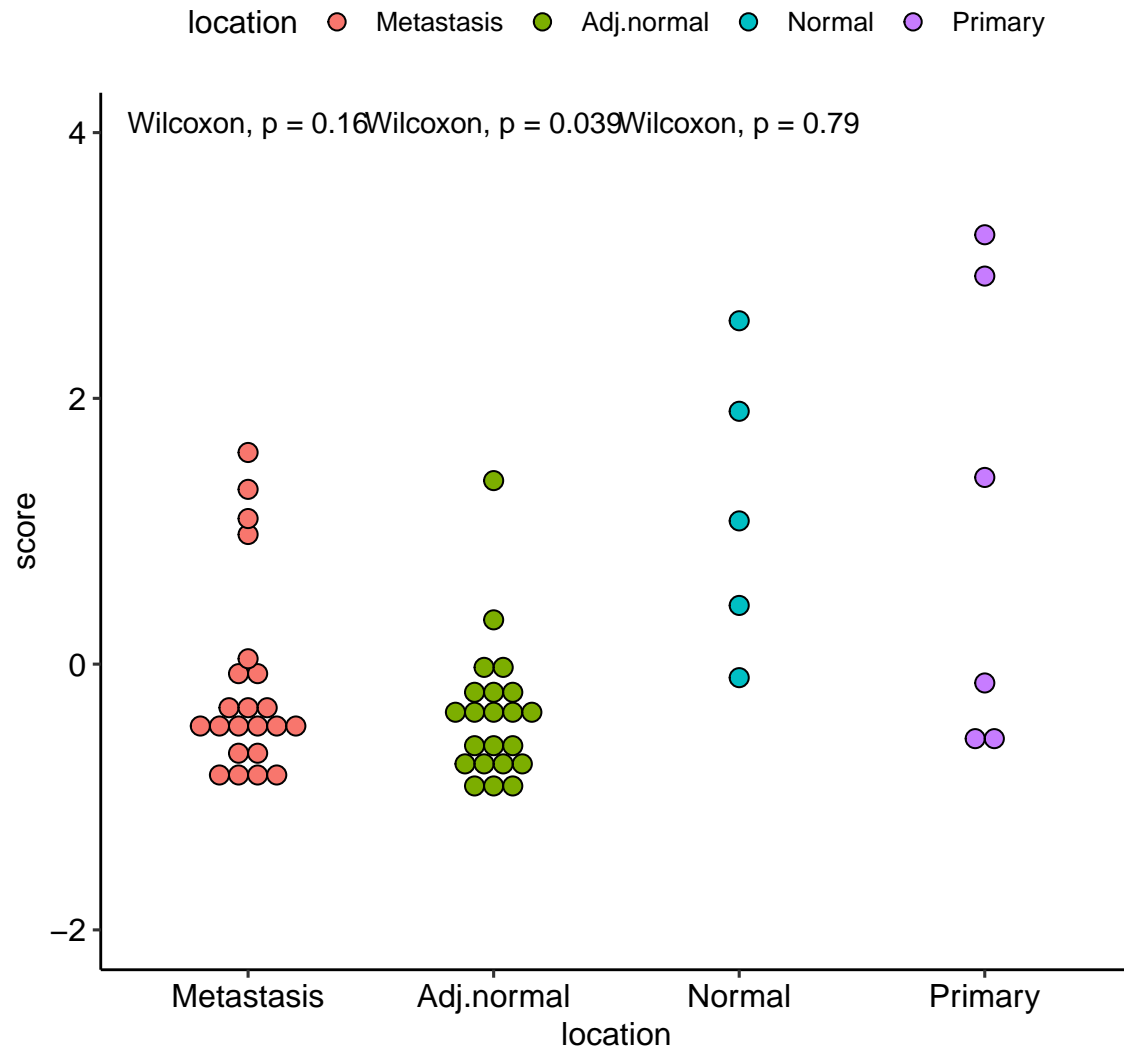
```
## Warning: Removed 2 rows containing non-finite values (stat_compare_means).
```

## Collisson.QM score





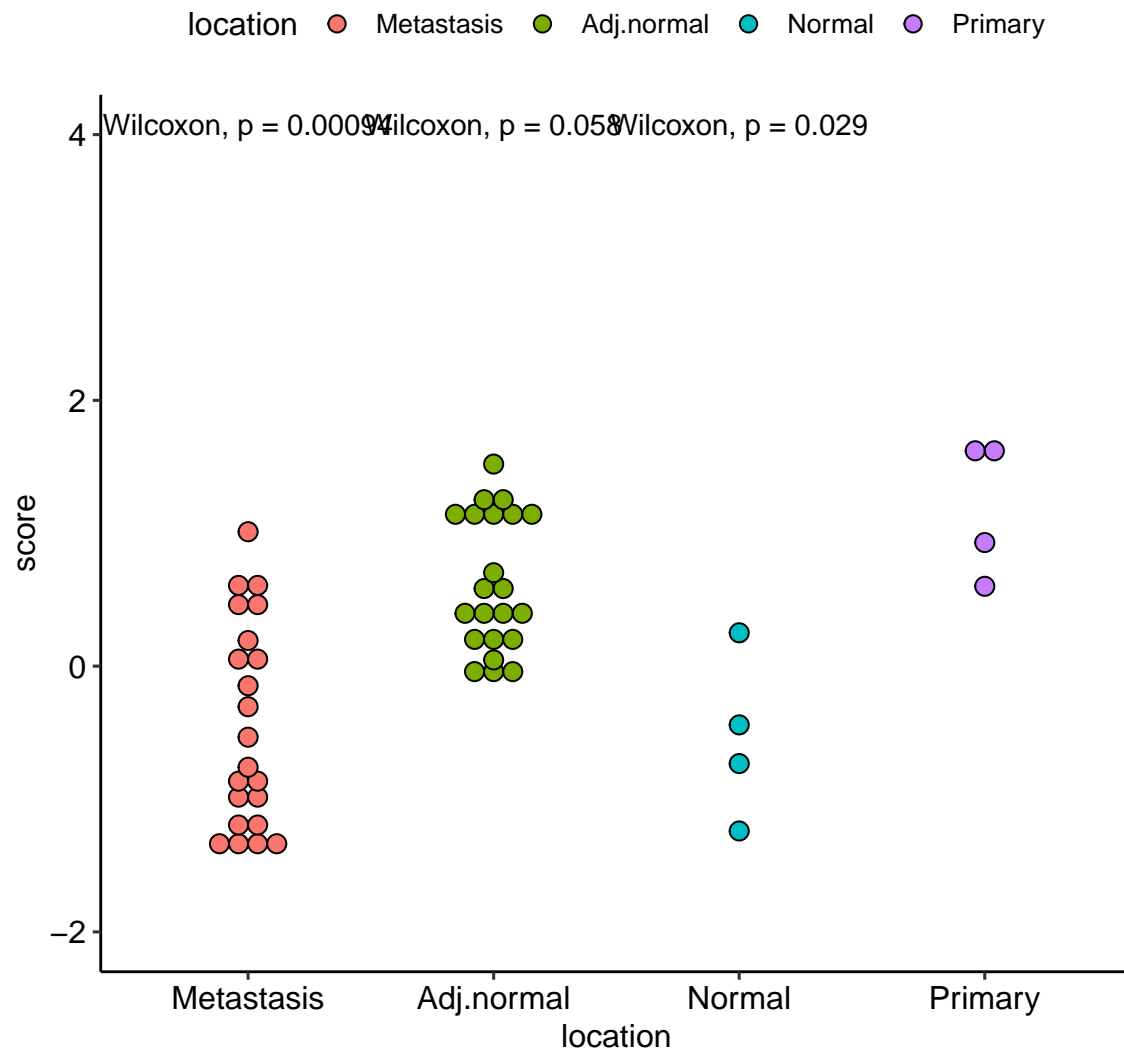
## ICGC.ADEX.Up score



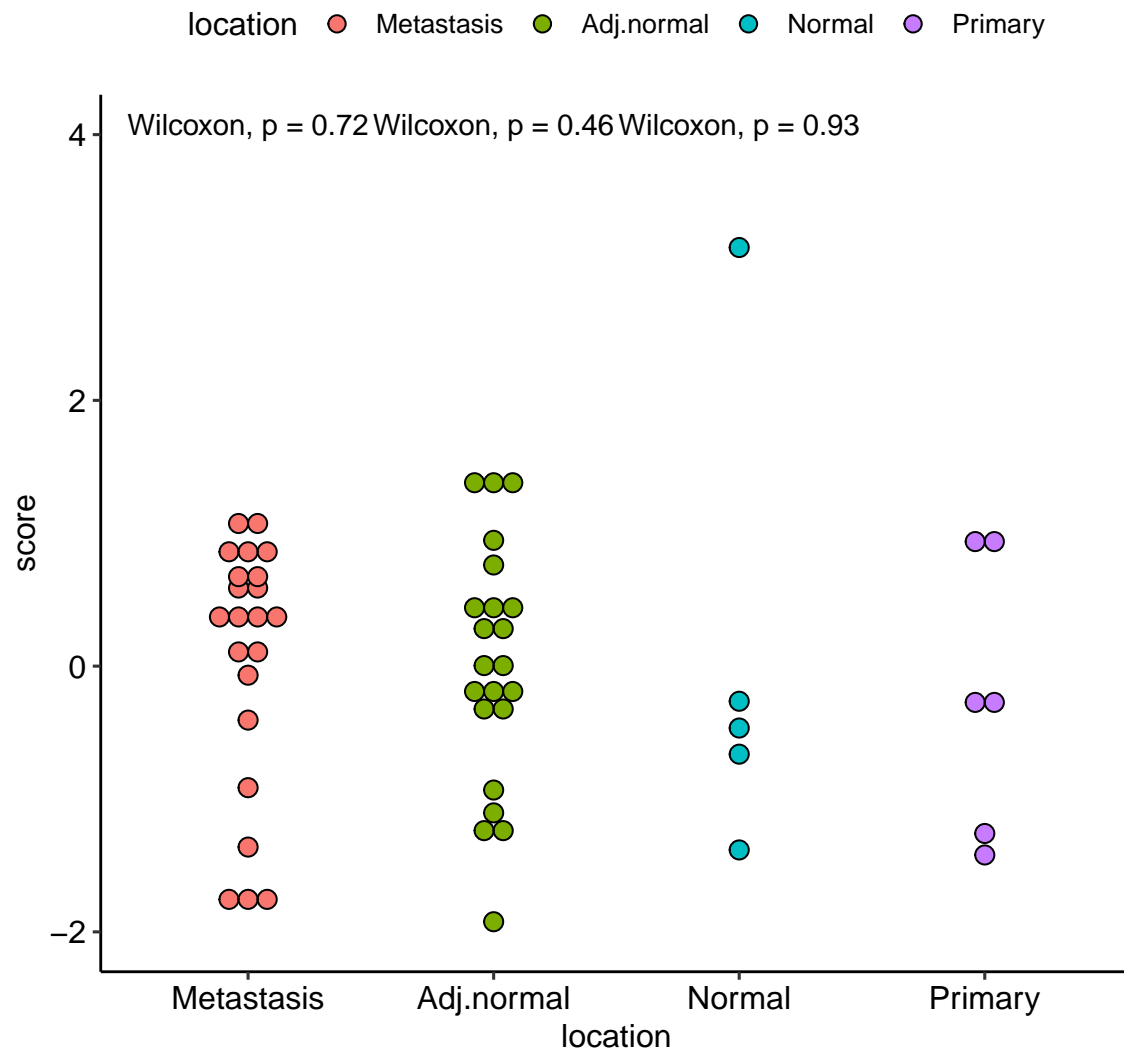
```
## Warning: Removed 3 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 3 rows containing non-finite values (stat_compare_means).
```

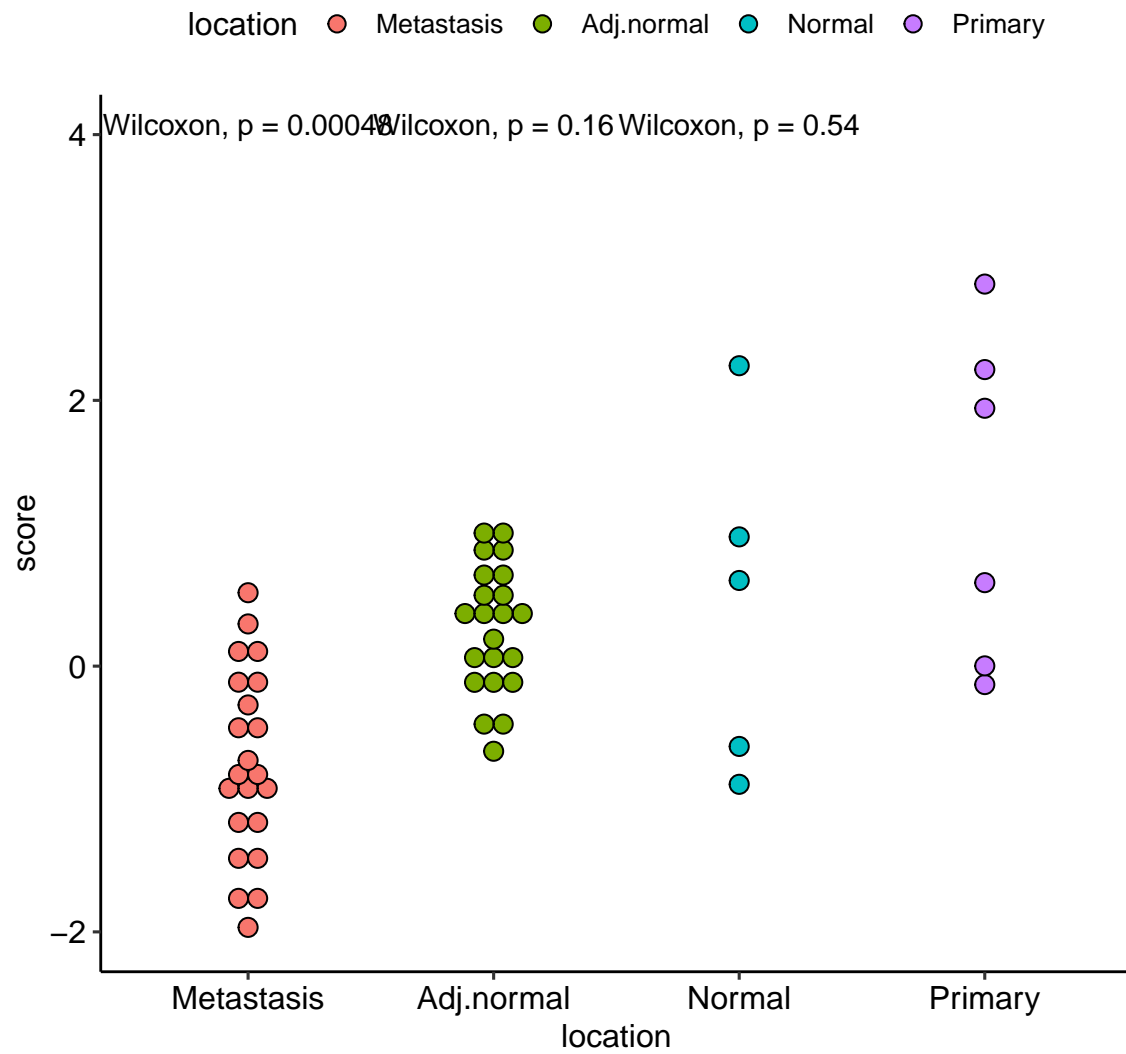
## ICGC.ADEX.Down score



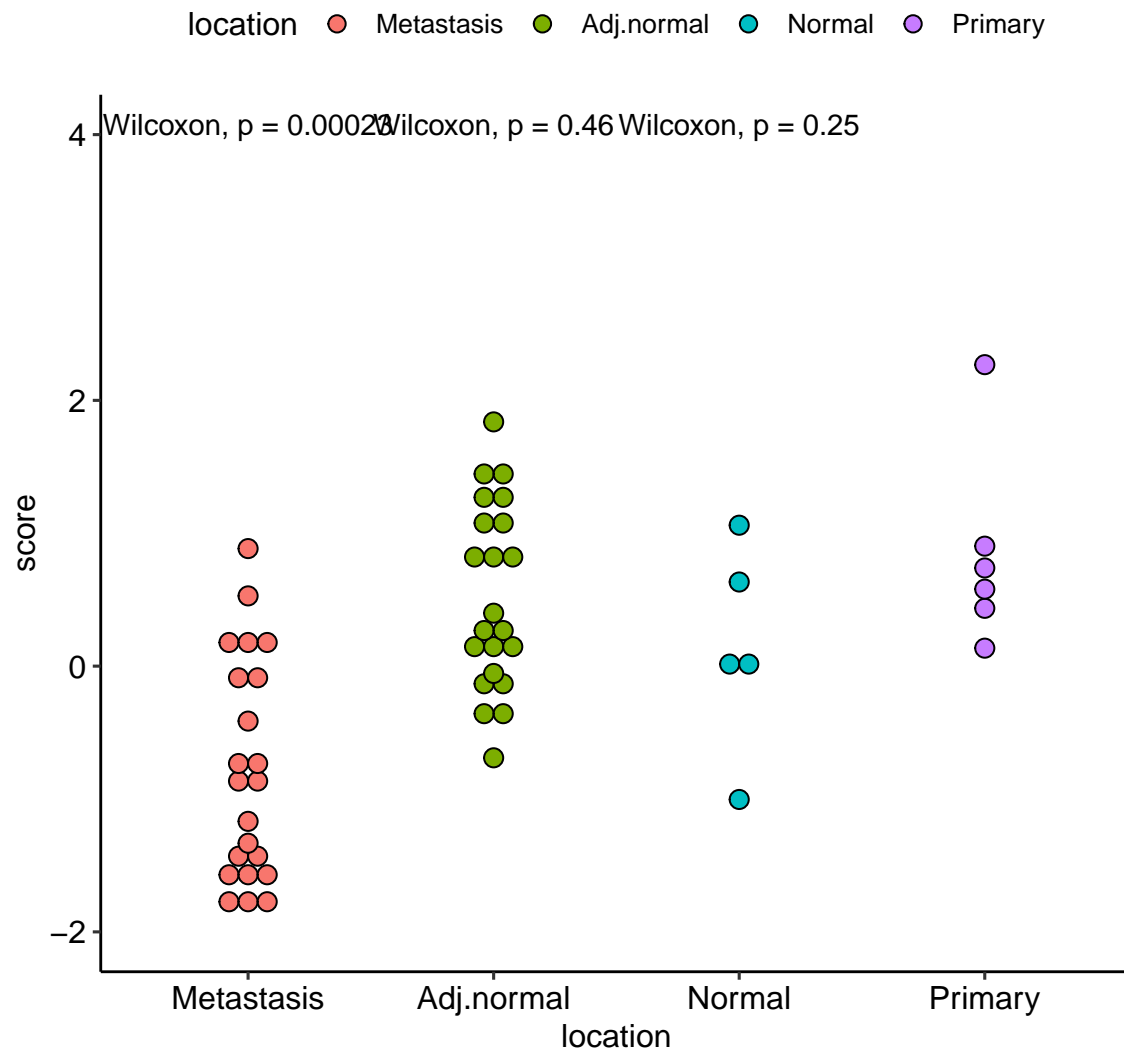
## ICGC.Immunogenic.Up score



# ICGC.Immunogenic.Down score



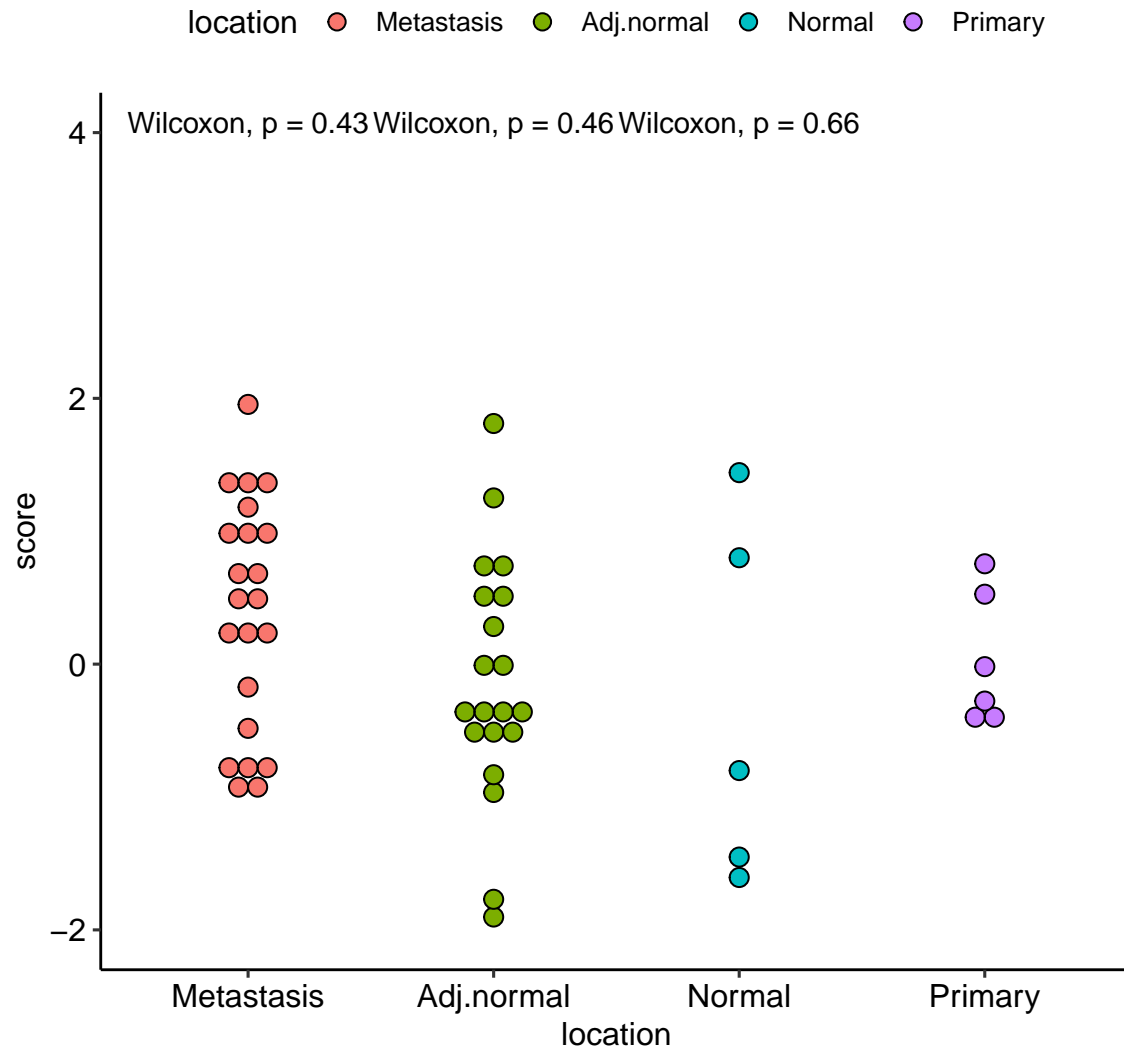
## ICGC.Progenitor.Up score



```
## Warning: Removed 2 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 2 rows containing non-finite values (stat_compare_means).
```

## ICGC.Progenitor.Down score

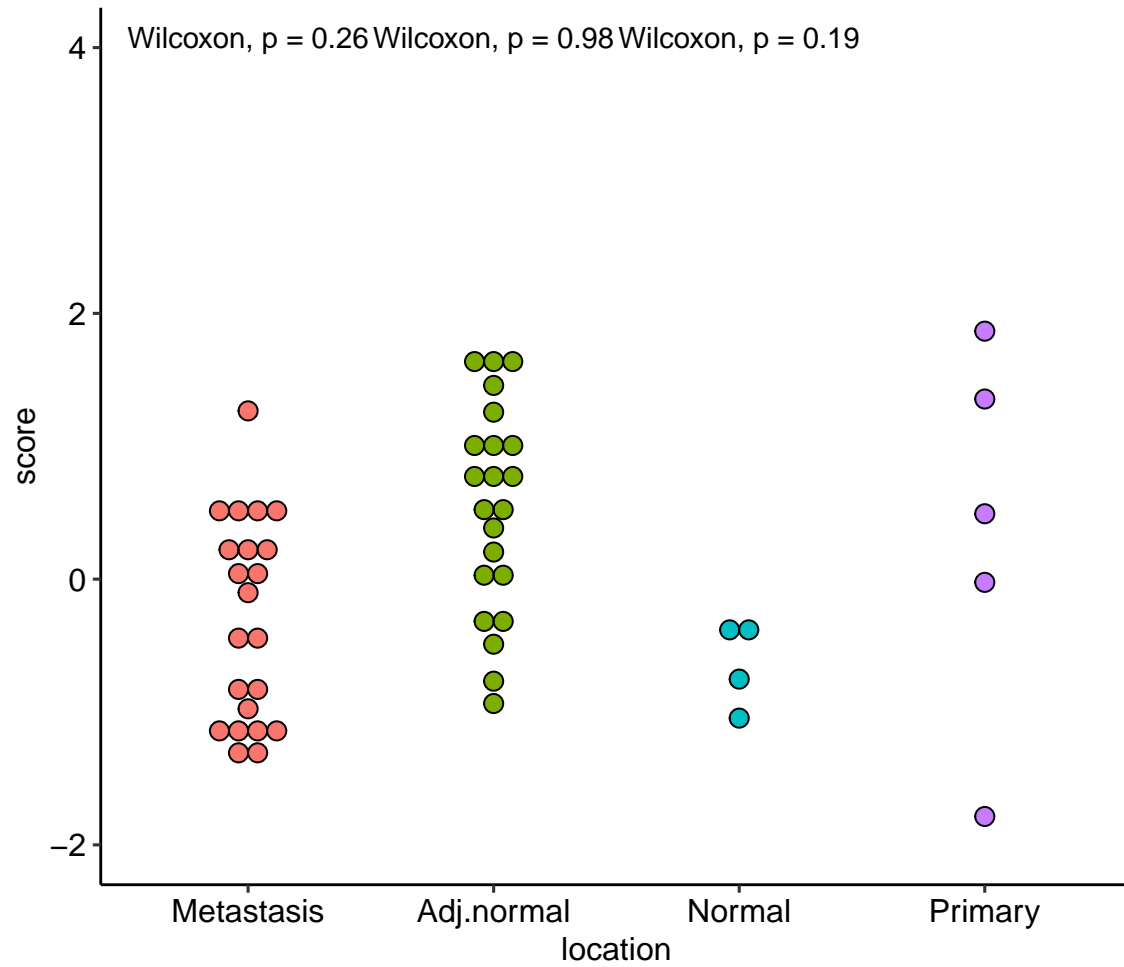


## Warning: Removed 2 rows containing non-finite values (stat\_bindot).

## Warning: Removed 2 rows containing non-finite values (stat\_compare\_means).

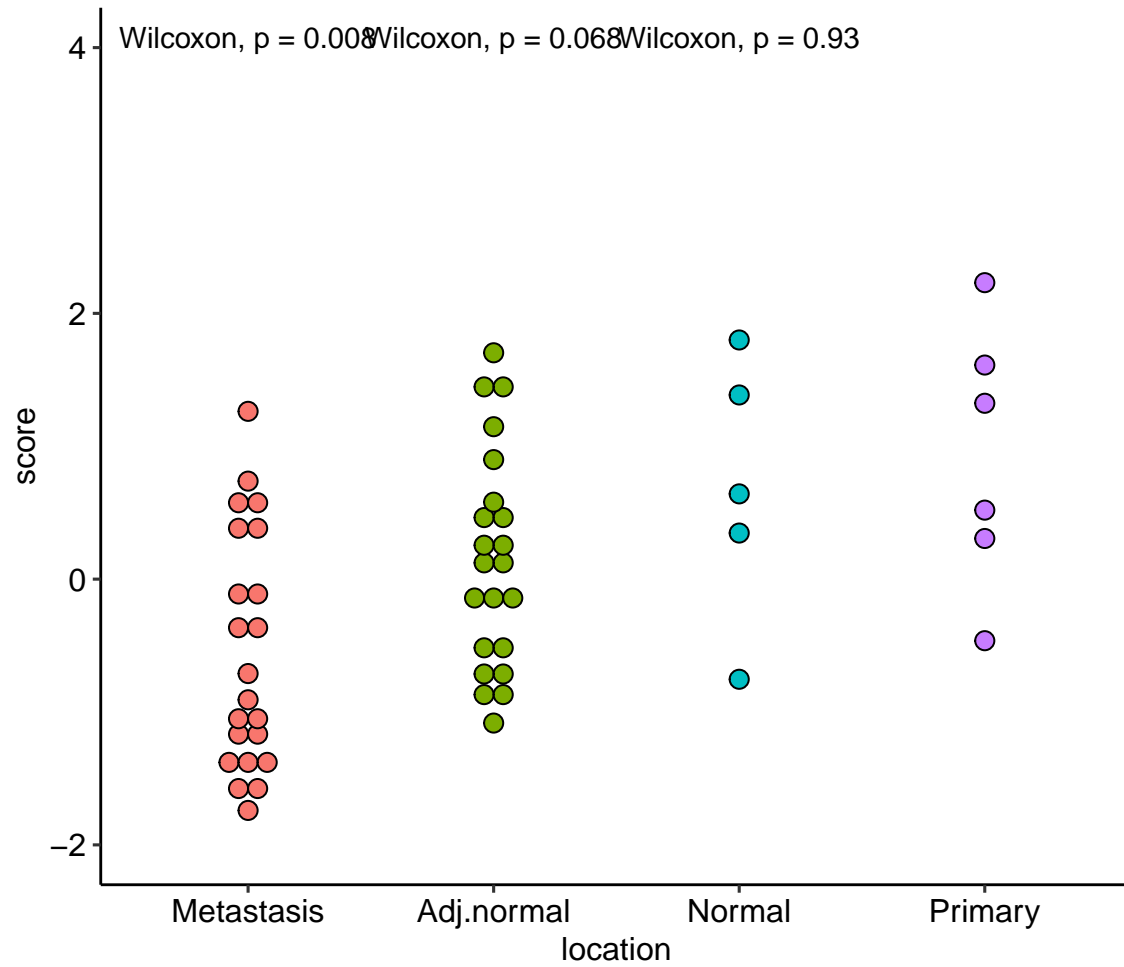
## ICGC.Squamous.Up score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



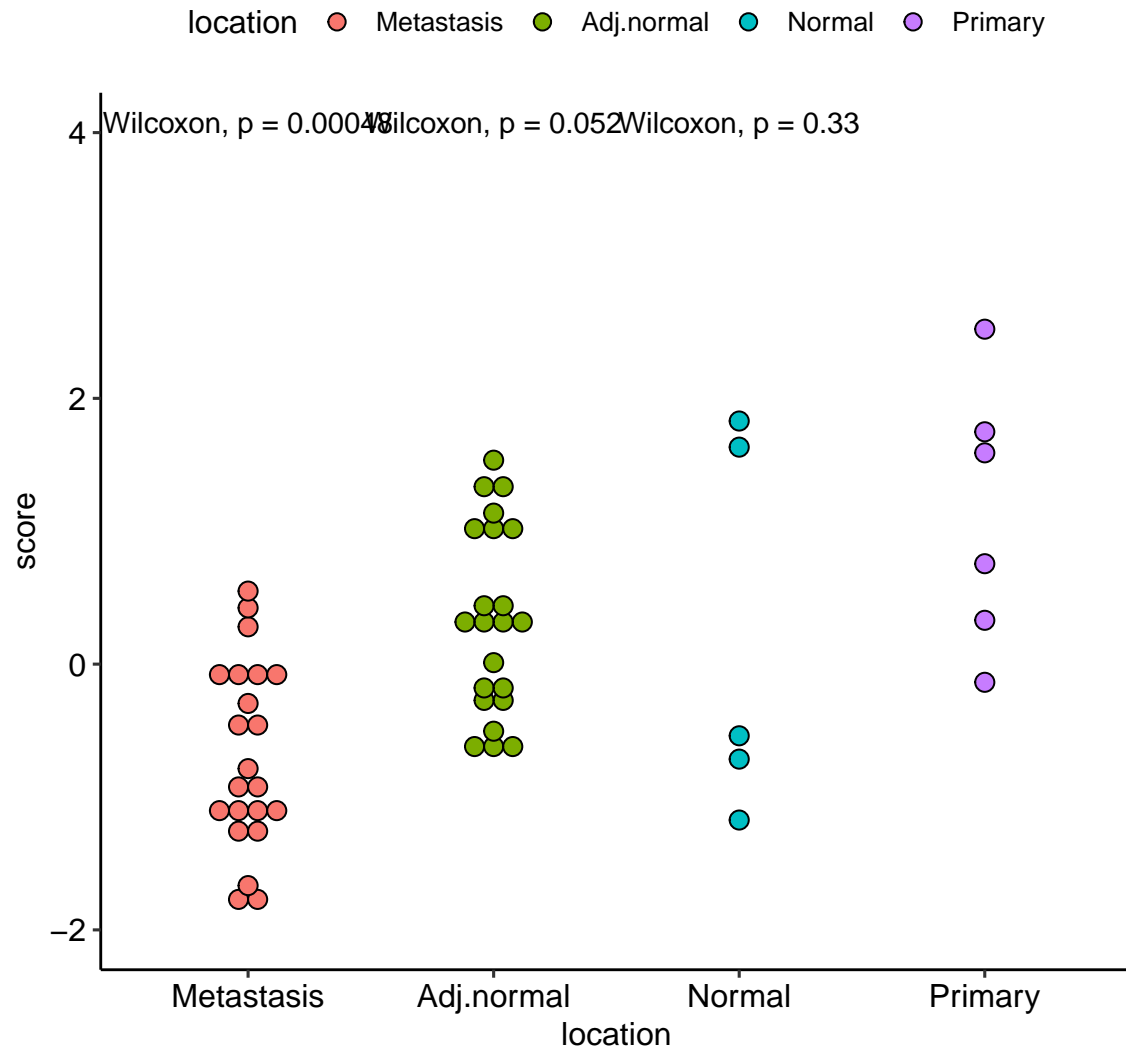
## ICGC.Squamous.Down score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

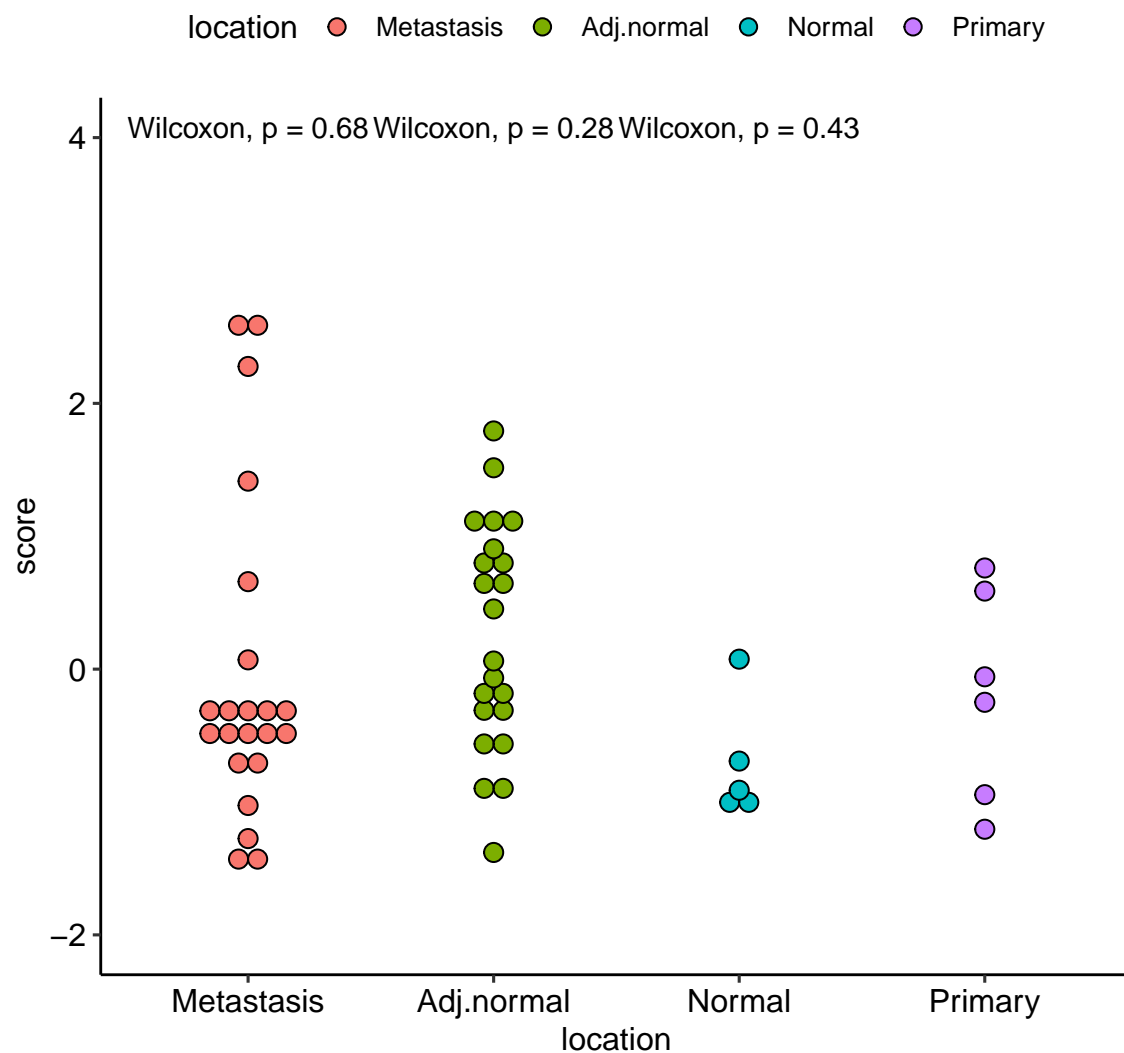




# ICGC.SAM score



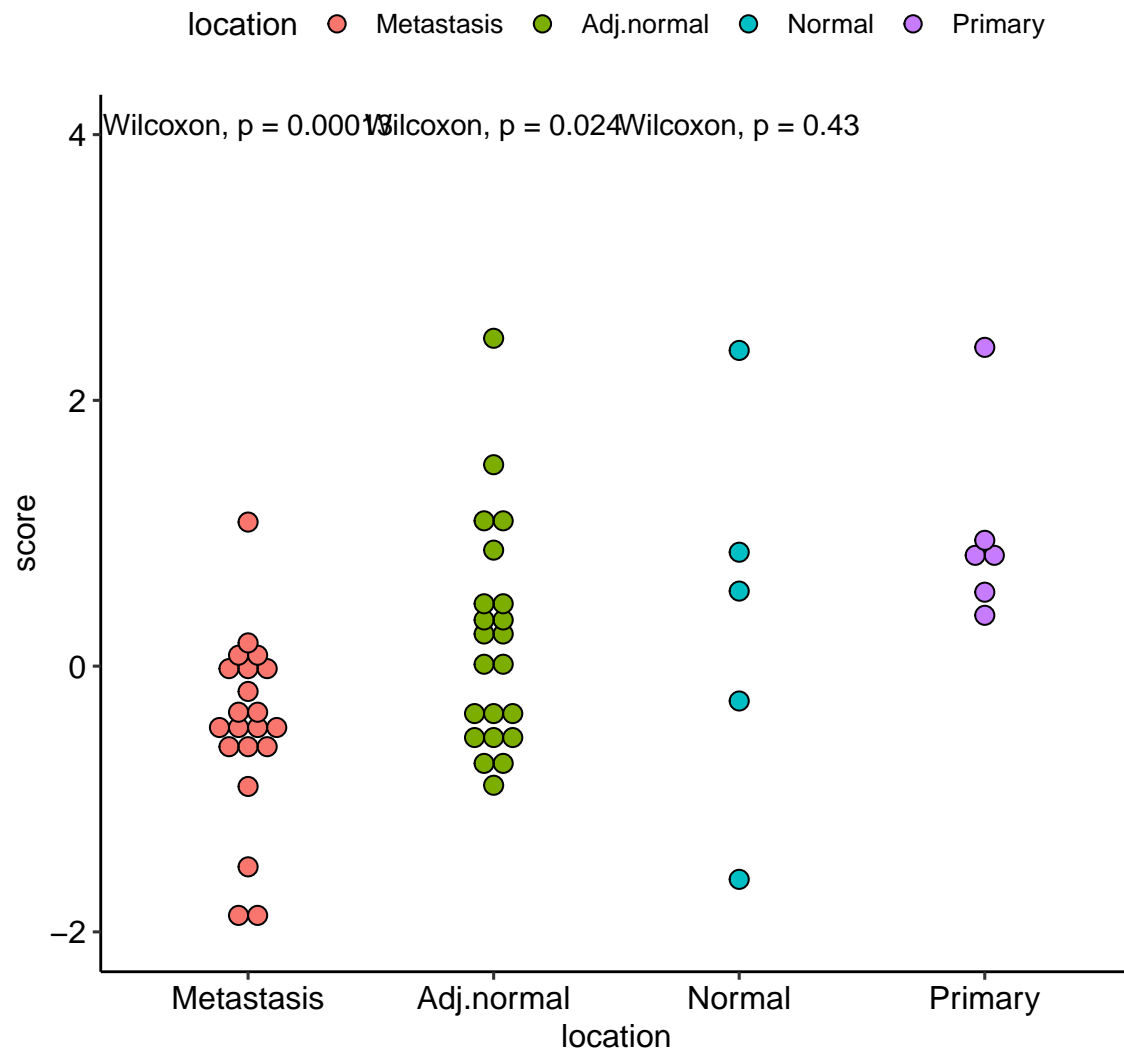
## Scarpa\_PanNET score



```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

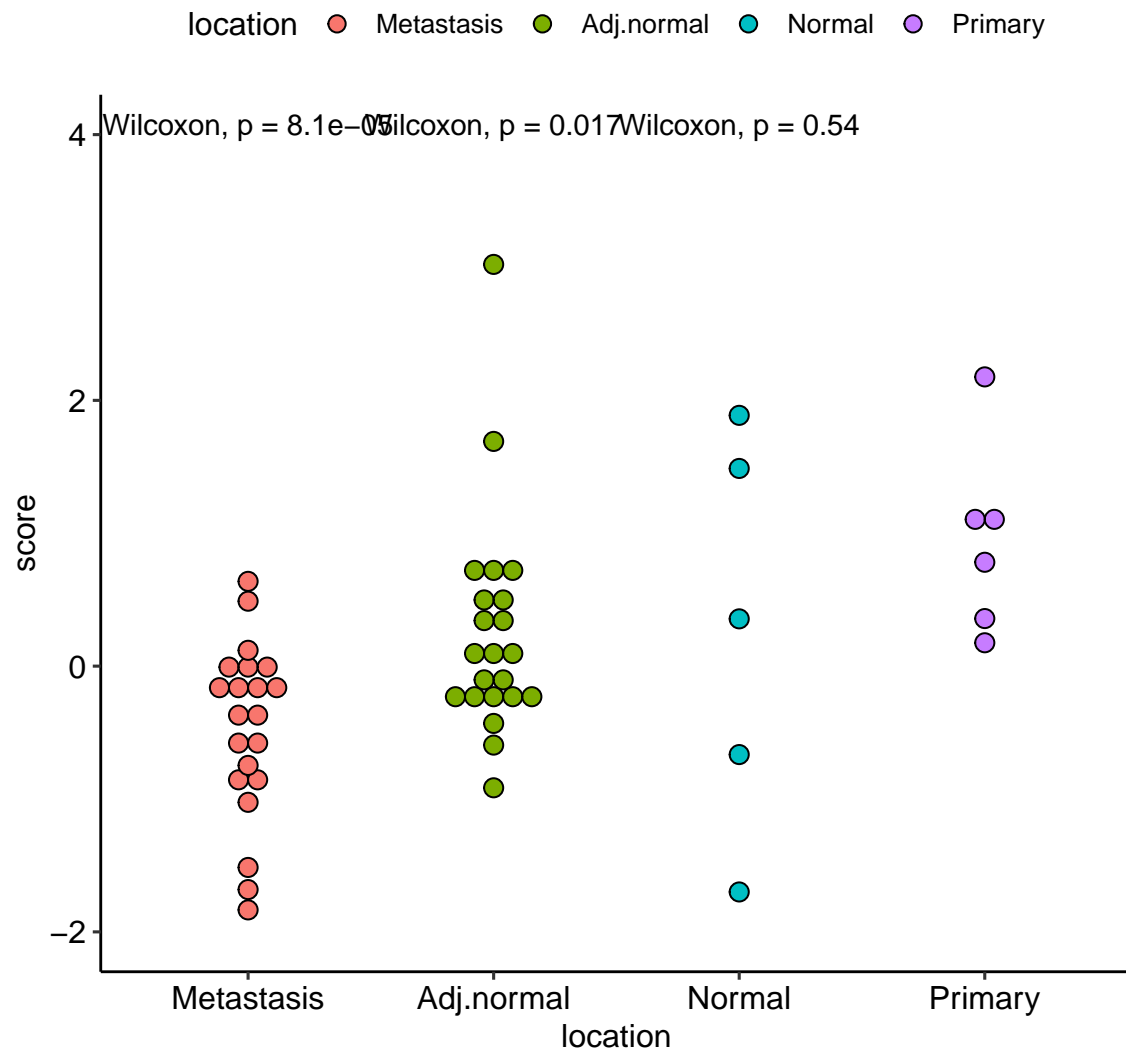
## Moffitt.Top5s score



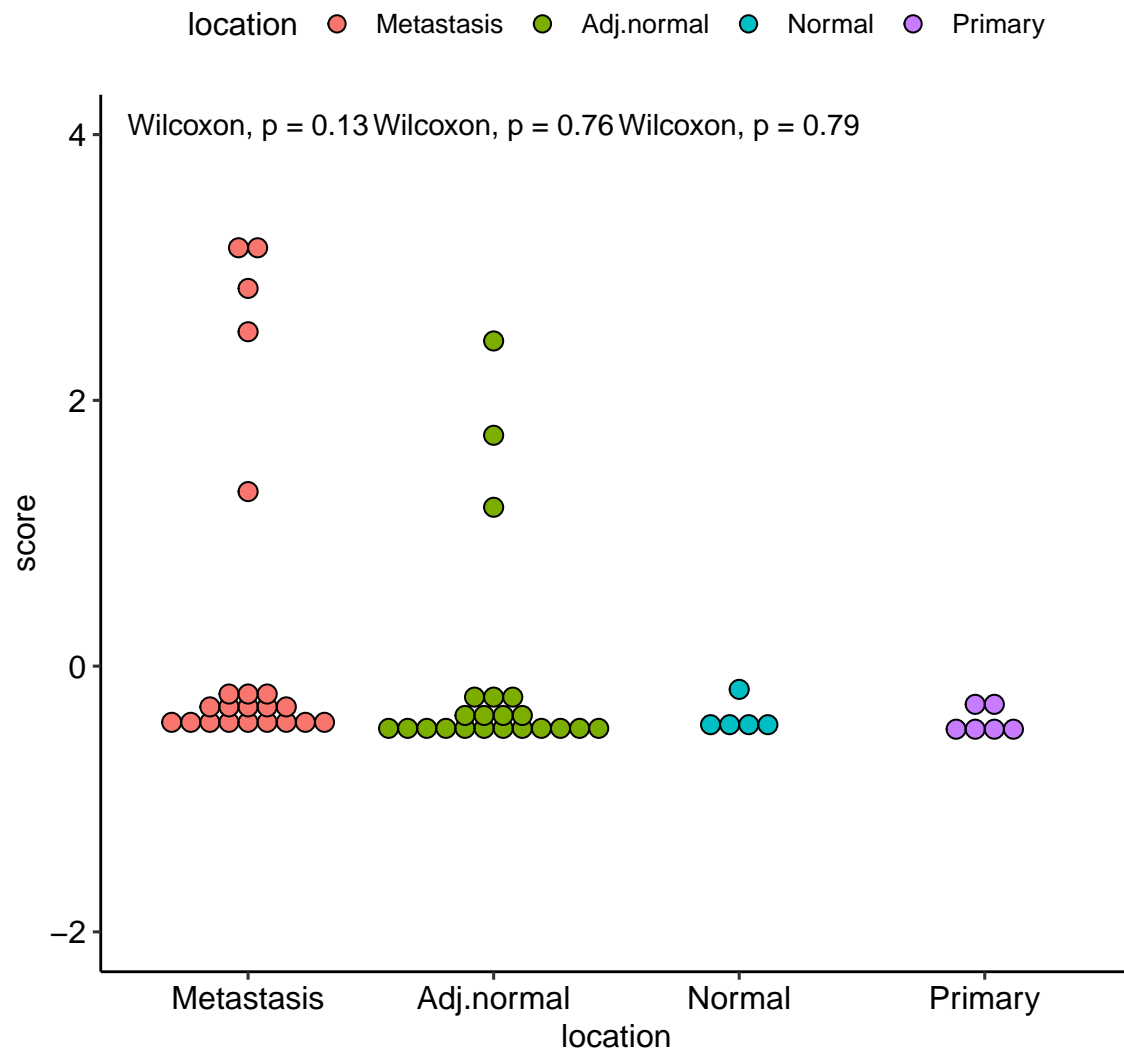
```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

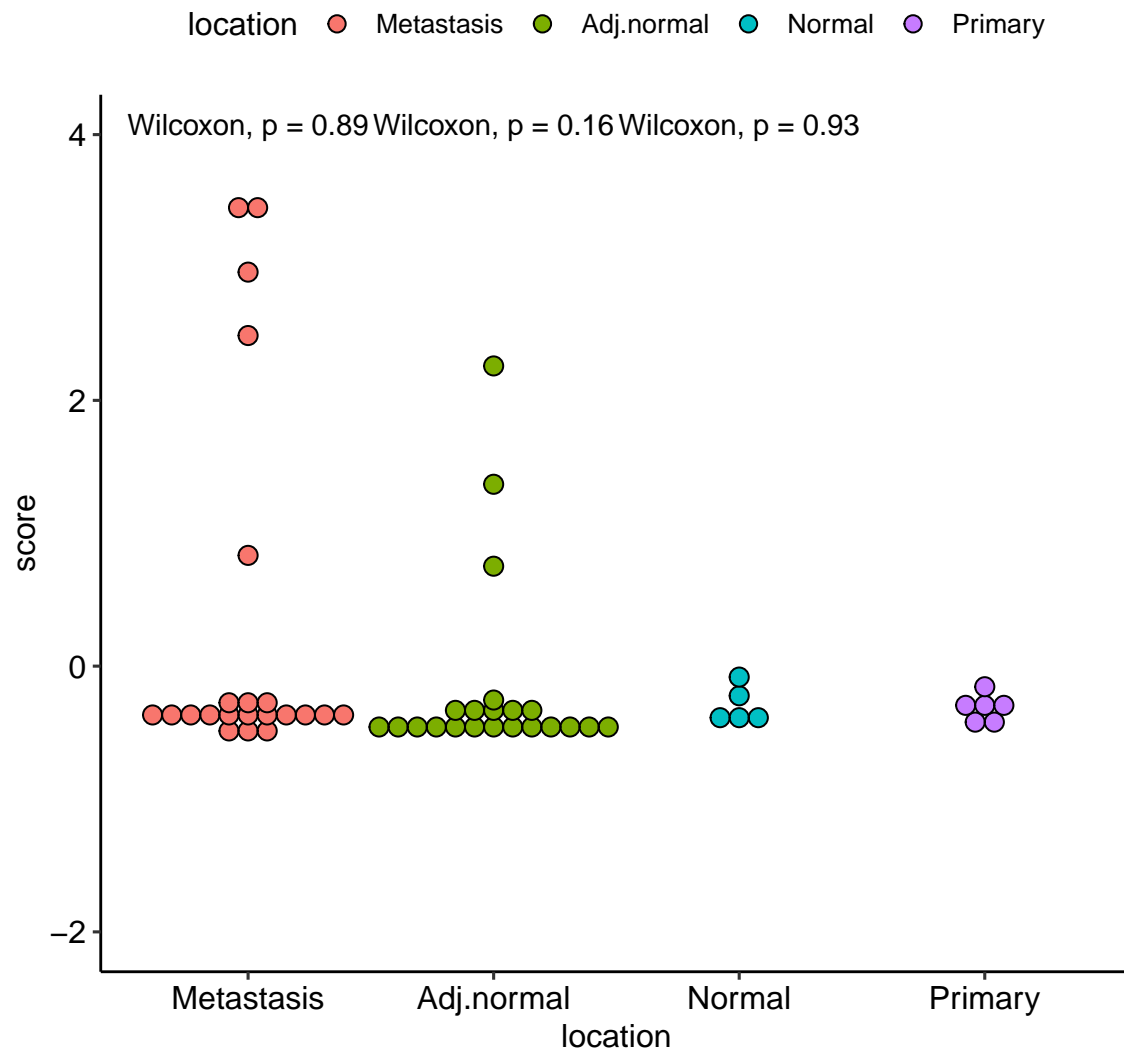
# Moffitt.Top25s score



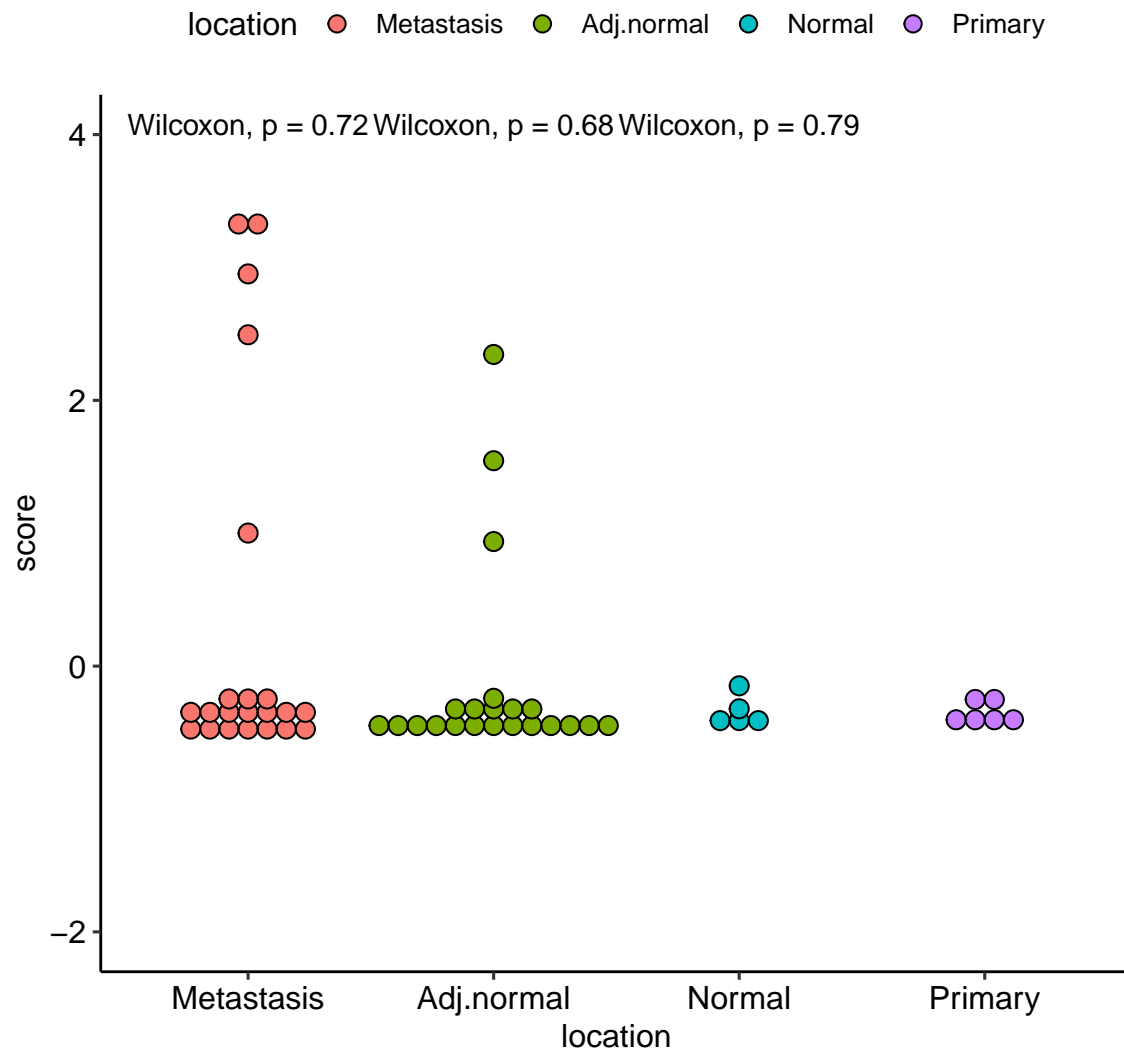
## Moffitt.F1\_Liver.top25 score



## Moffitt.F1\_Liver.top250 score



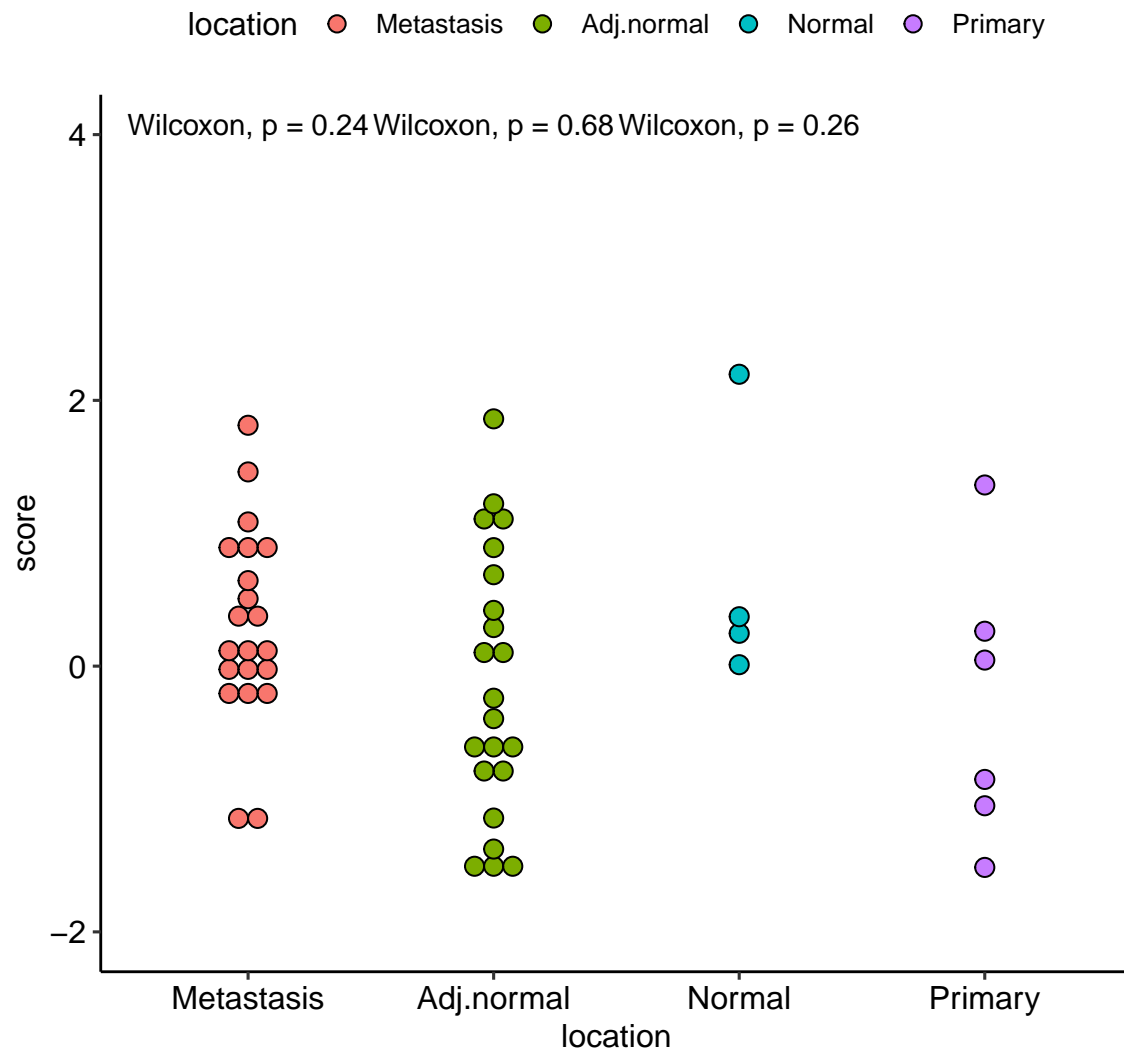
## Moffitt.F1\_Liver.top100 score



```
## Warning: Removed 2 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 2 rows containing non-finite values (stat_compare_means).
```

## Moffitt.F2.top25 score

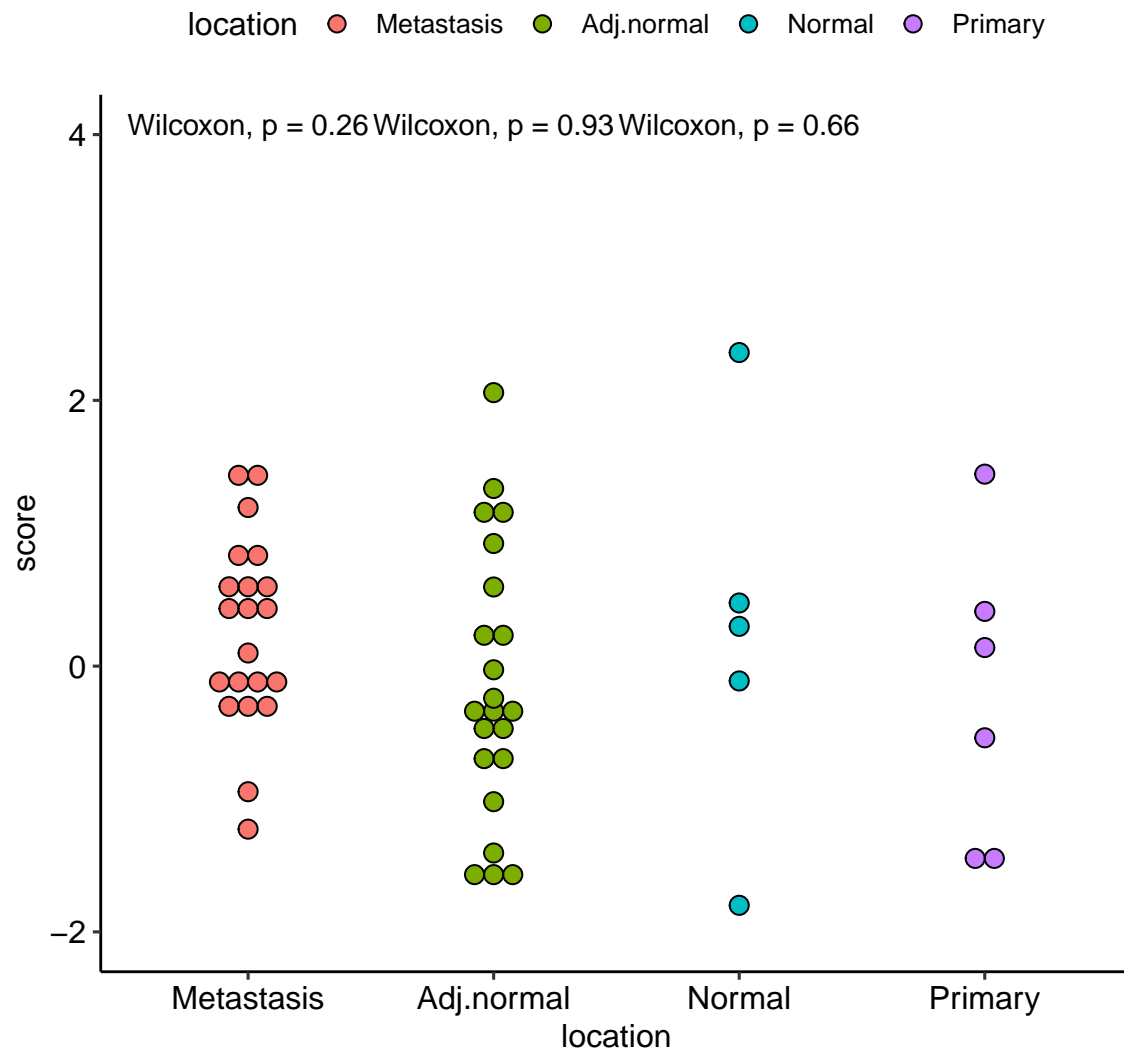


```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```



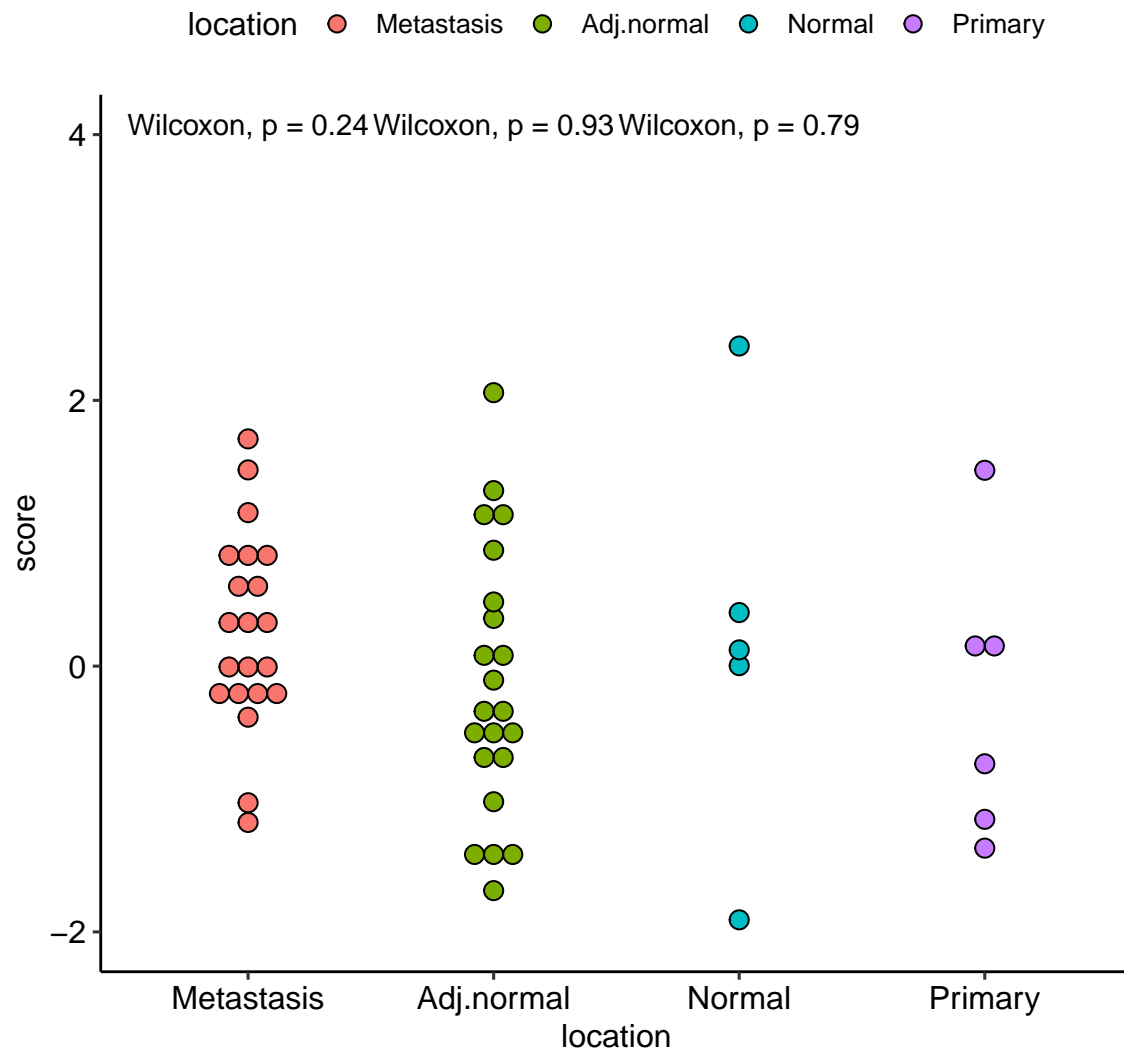
## Moffitt.F2.top250 score



```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

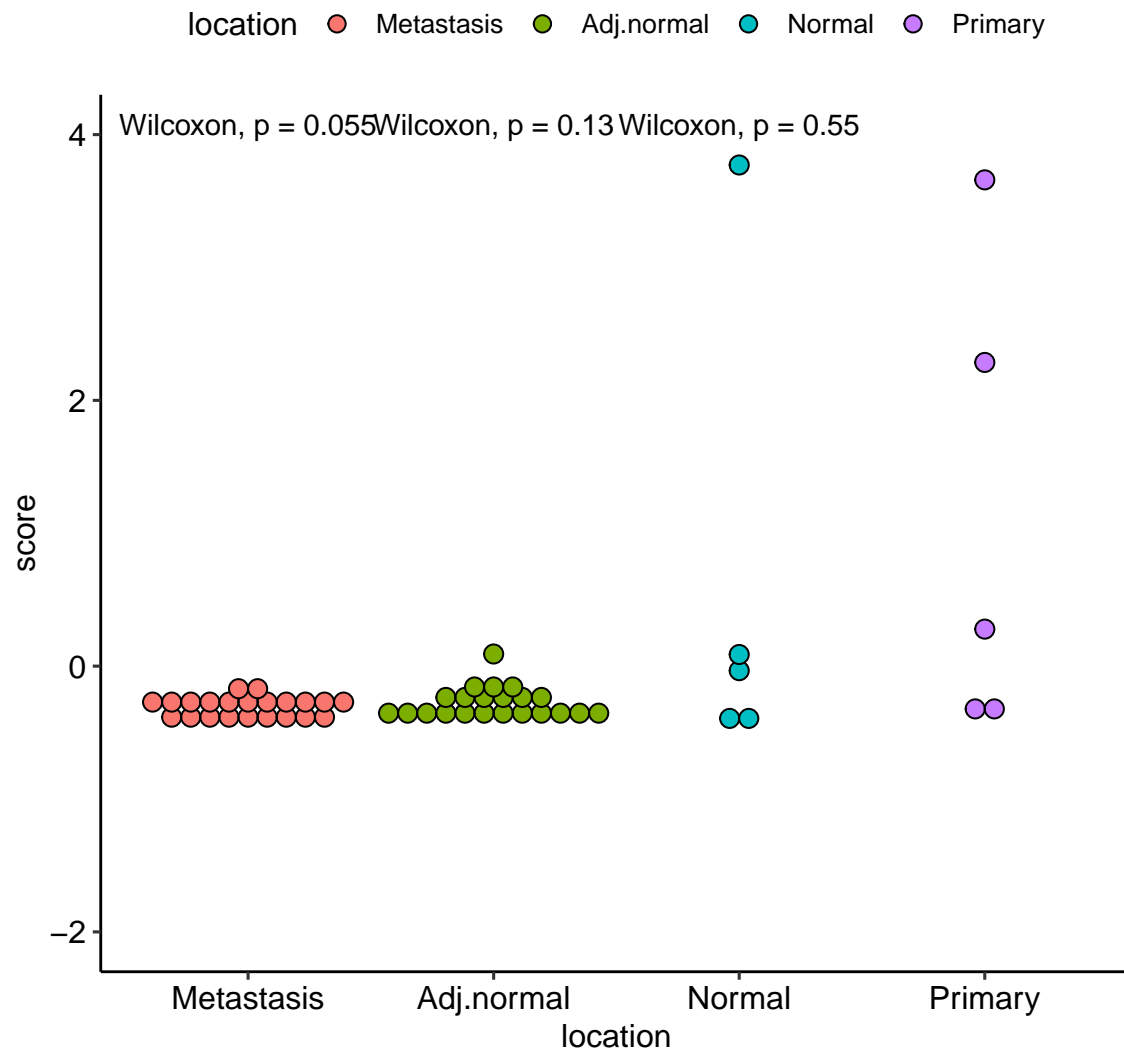
## Moffitt.F2.top100 score



```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

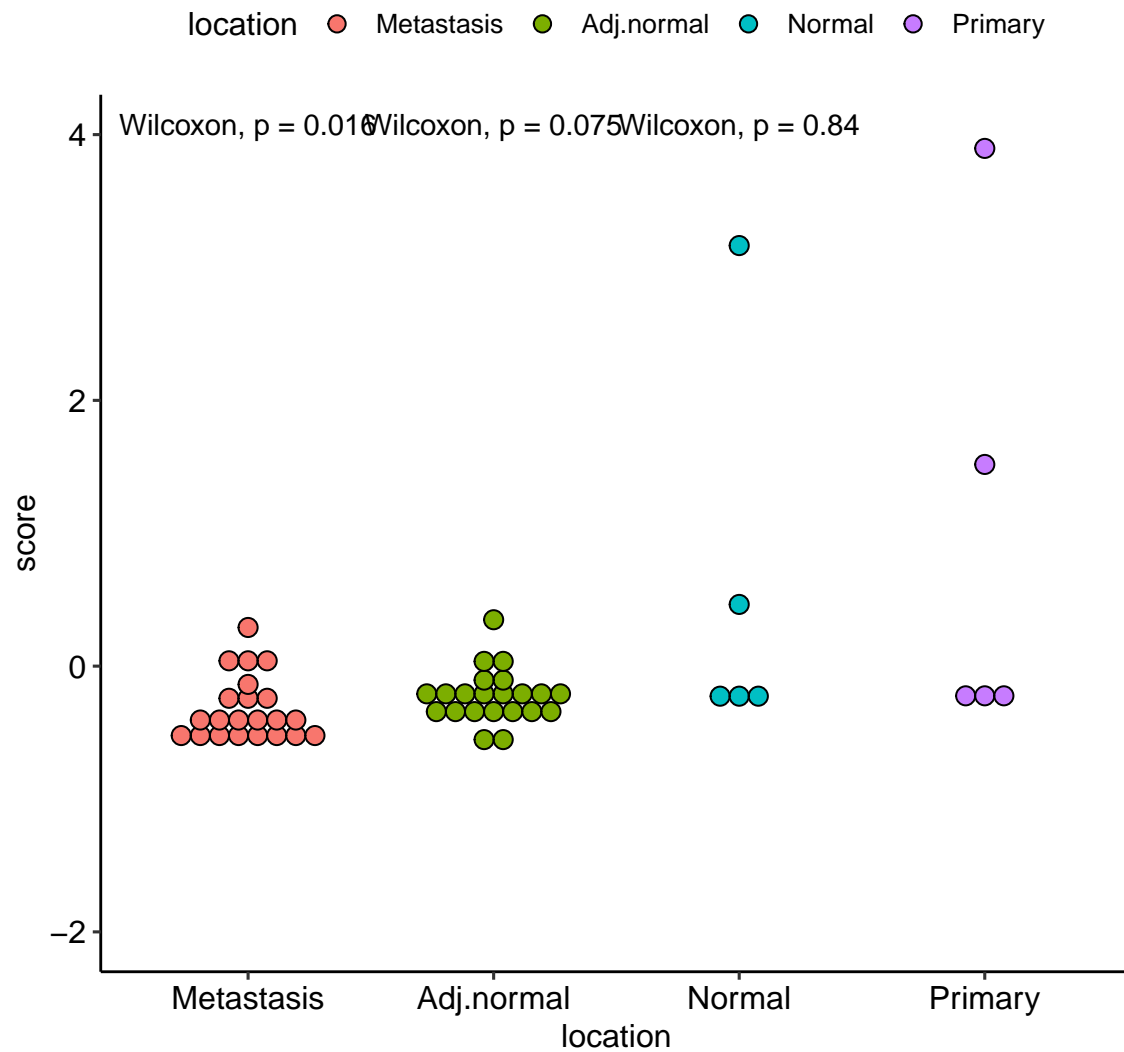
## Moffitt.F3\_Exocrine.top25 score



```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

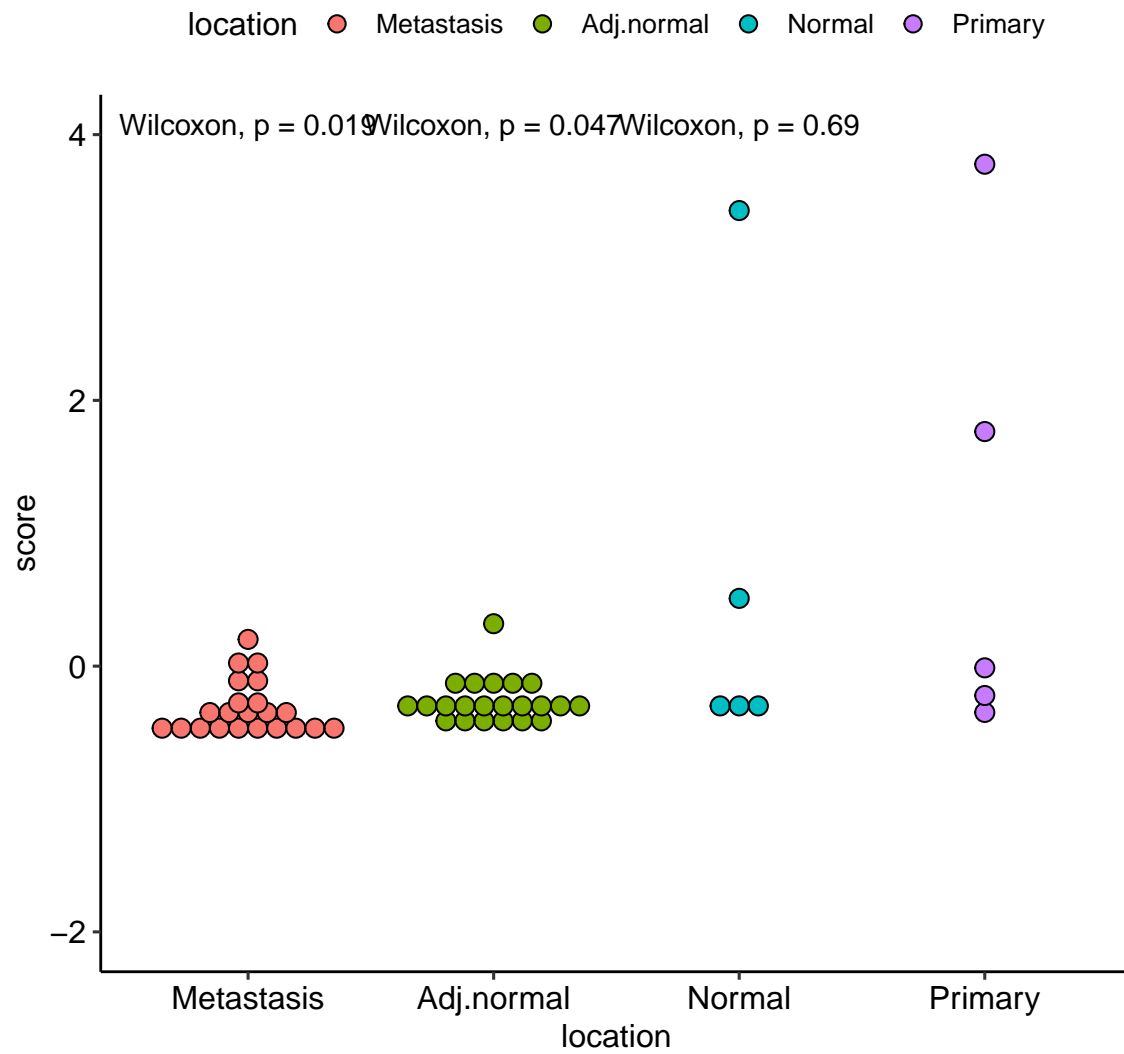
## Moffitt.F3\_Exocrine.top250 score



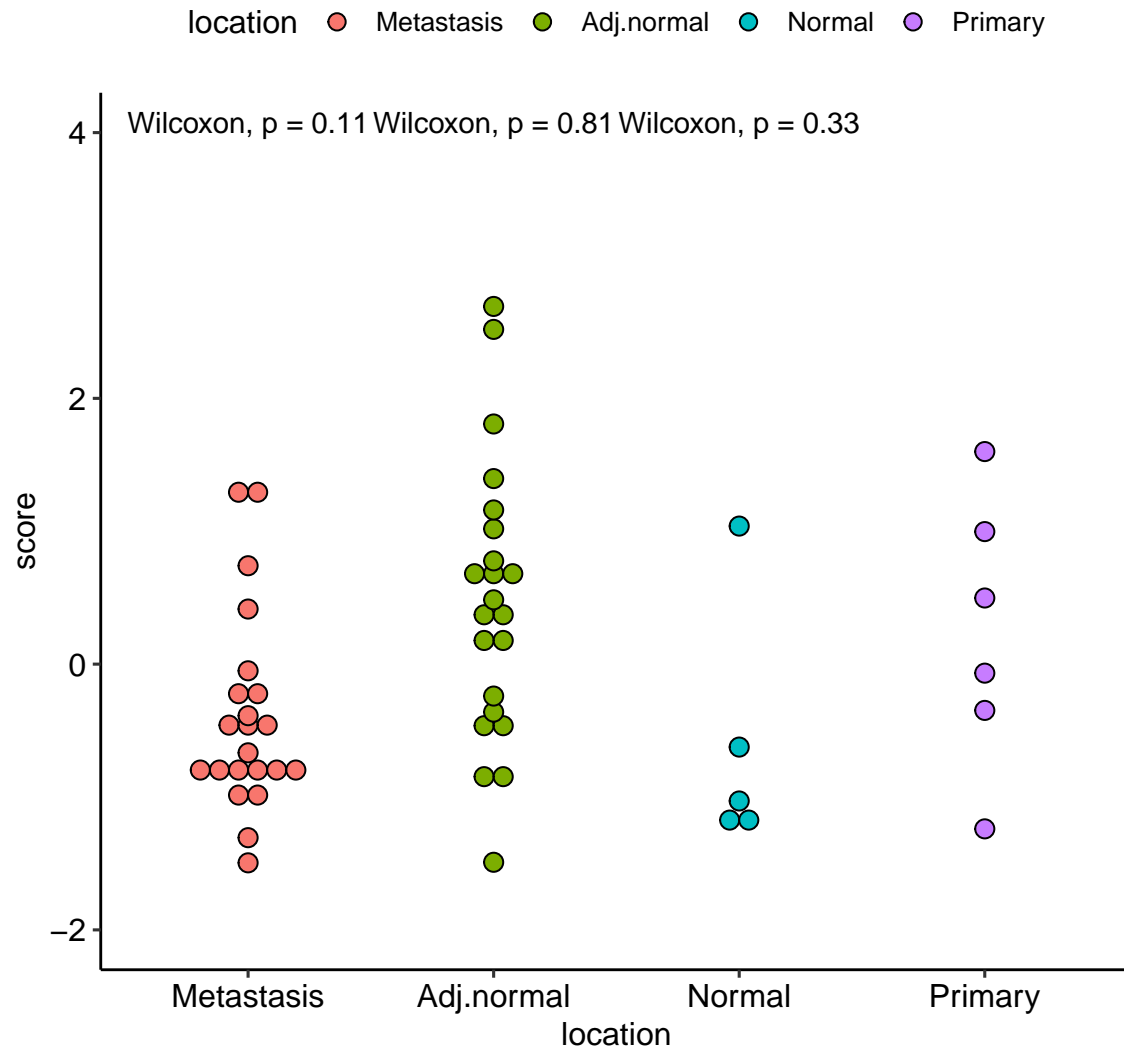
```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

## Moffitt.F3\_Exocrine.top100 score



## Moffitt.F4\_CellCycle.top25 score

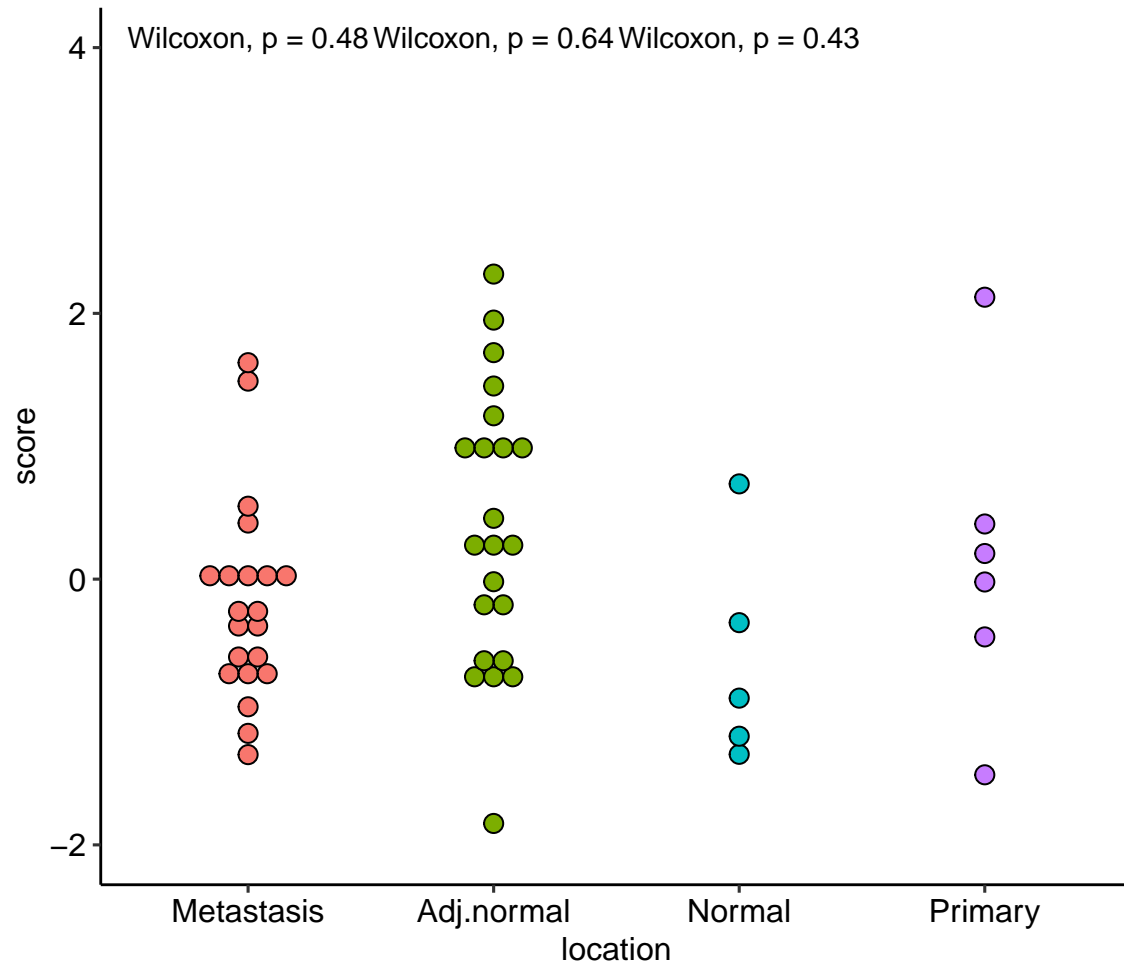


## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

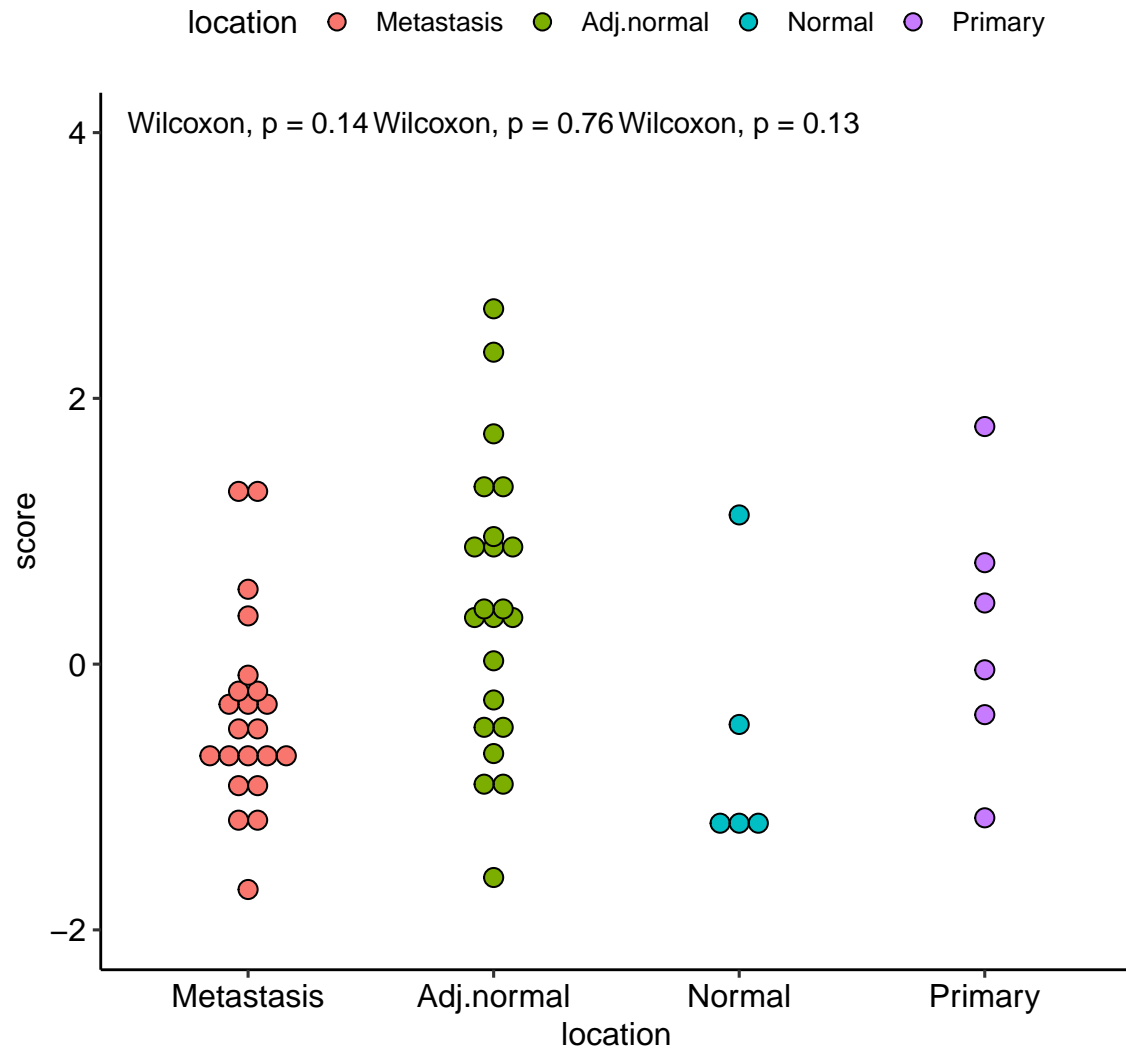
## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

## Moffitt.F4\_CellCycle.top250 score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

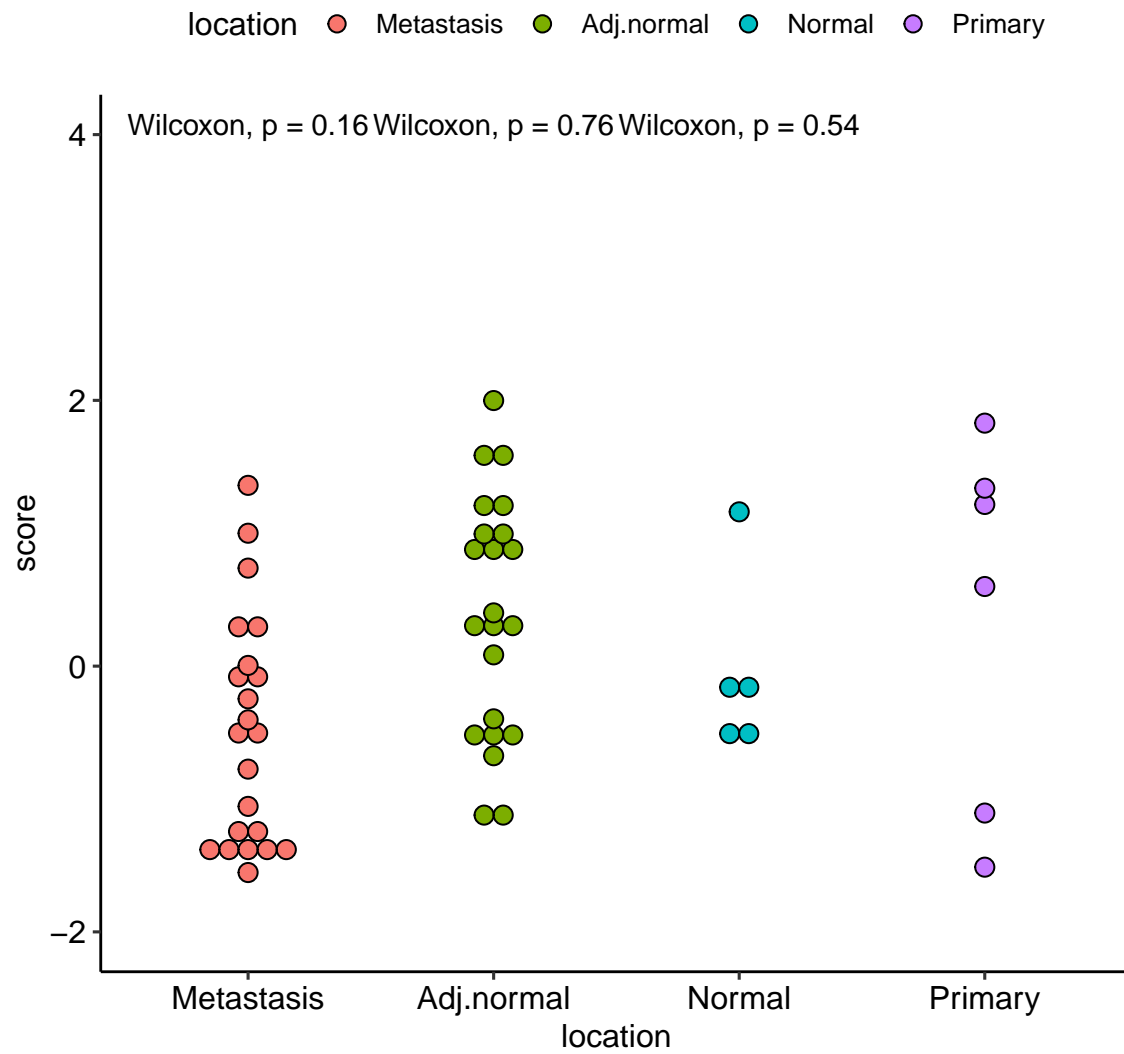


# Moffitt.F4\_CellCycle.top100 score



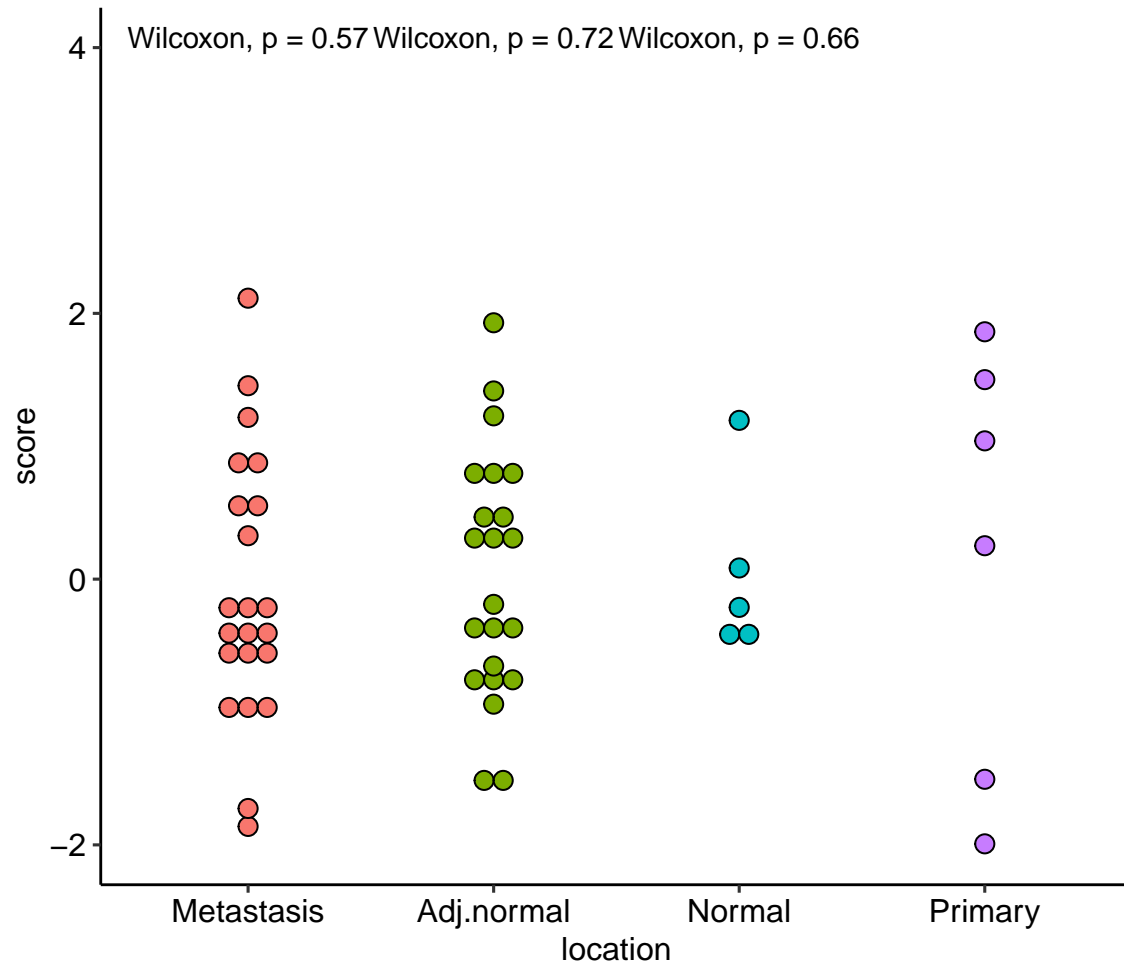


## Moffitt.F5\_ActivatedStroma.top25 score

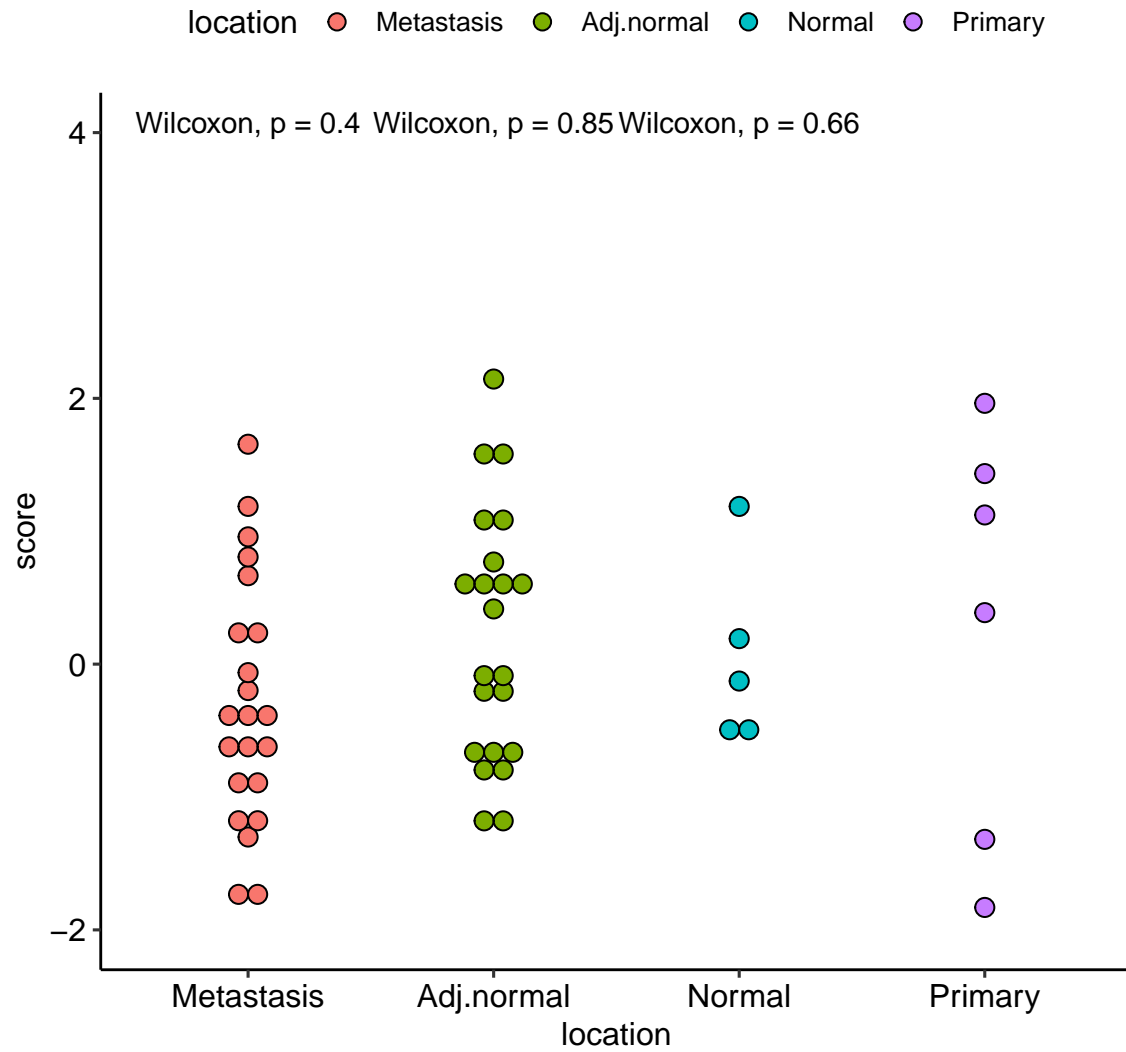


## Moffitt.F5\_ActivatedStroma.top250 score

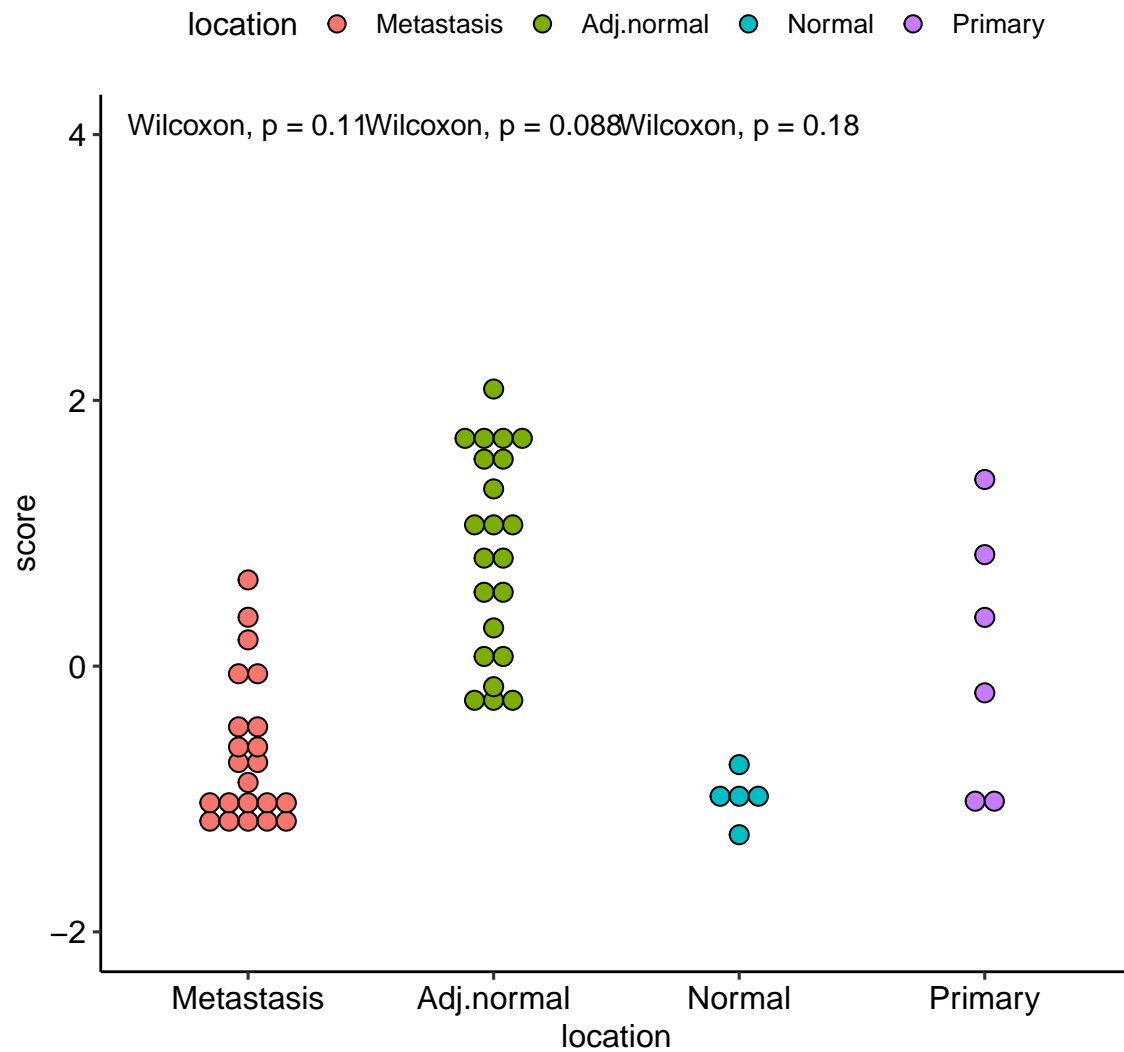
location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



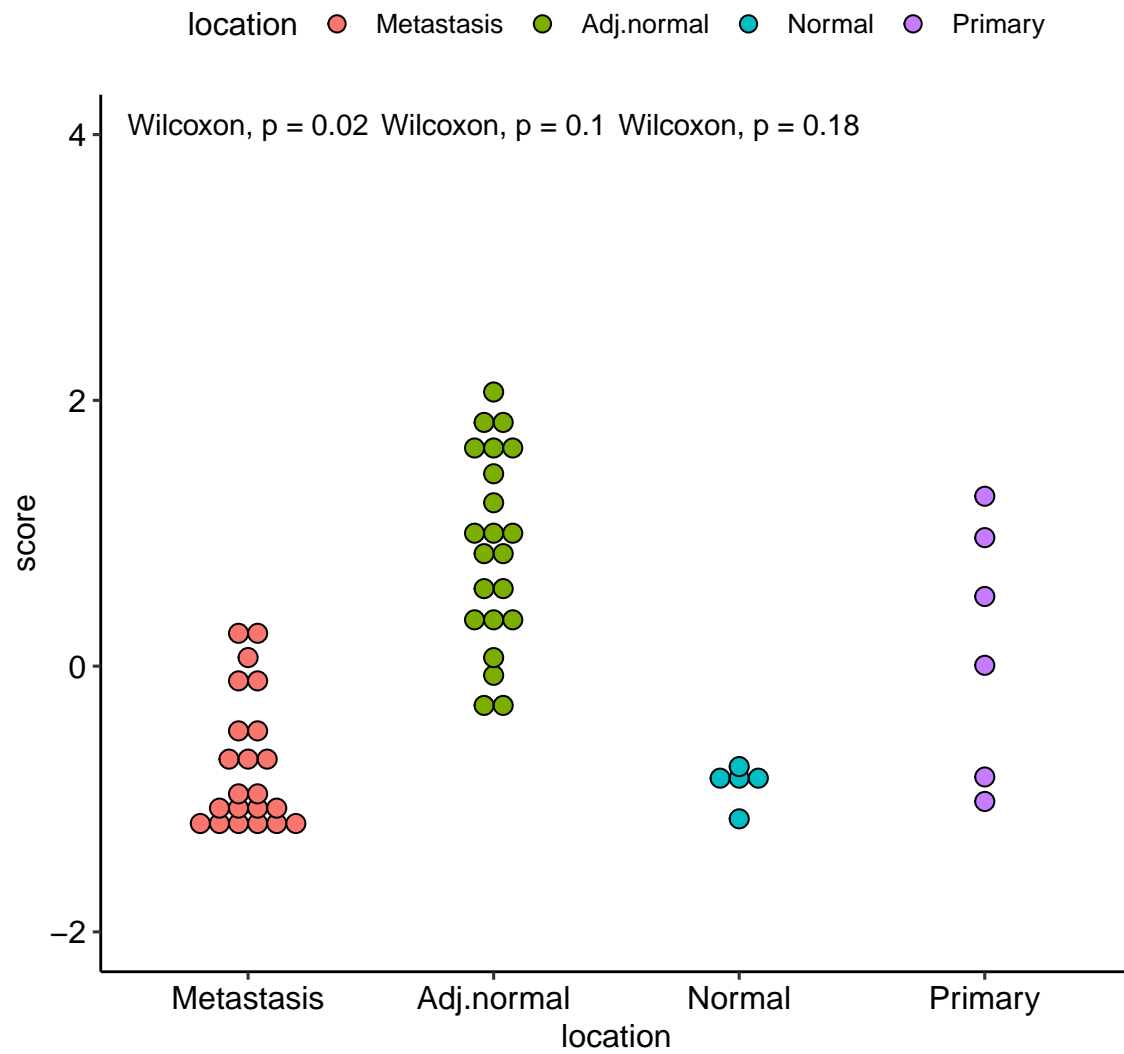
# Moffitt.F5\_ActivatedStroma.top100 score



## Moffitt.F6\_BasalLike.top25 score

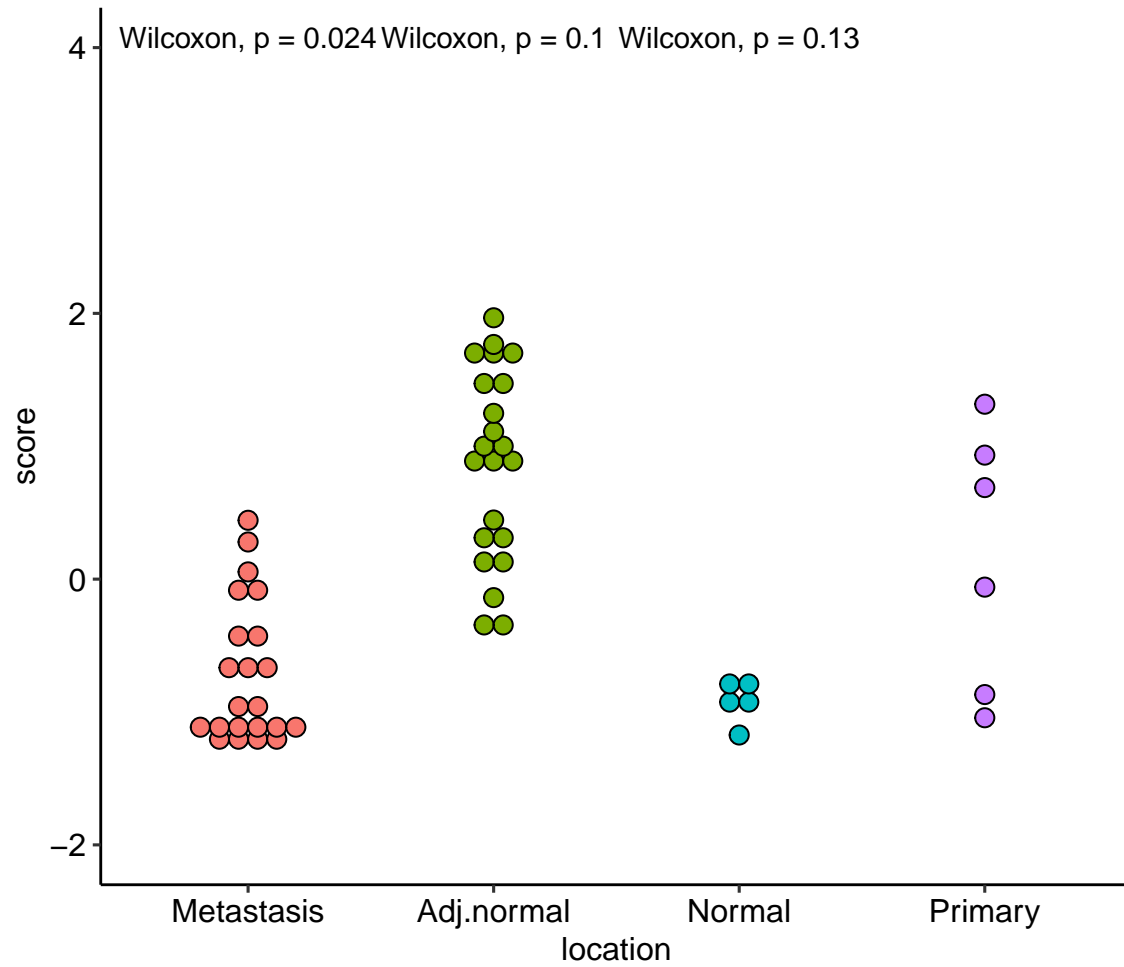


## Moffitt.F6\_BasalLike.top250 score



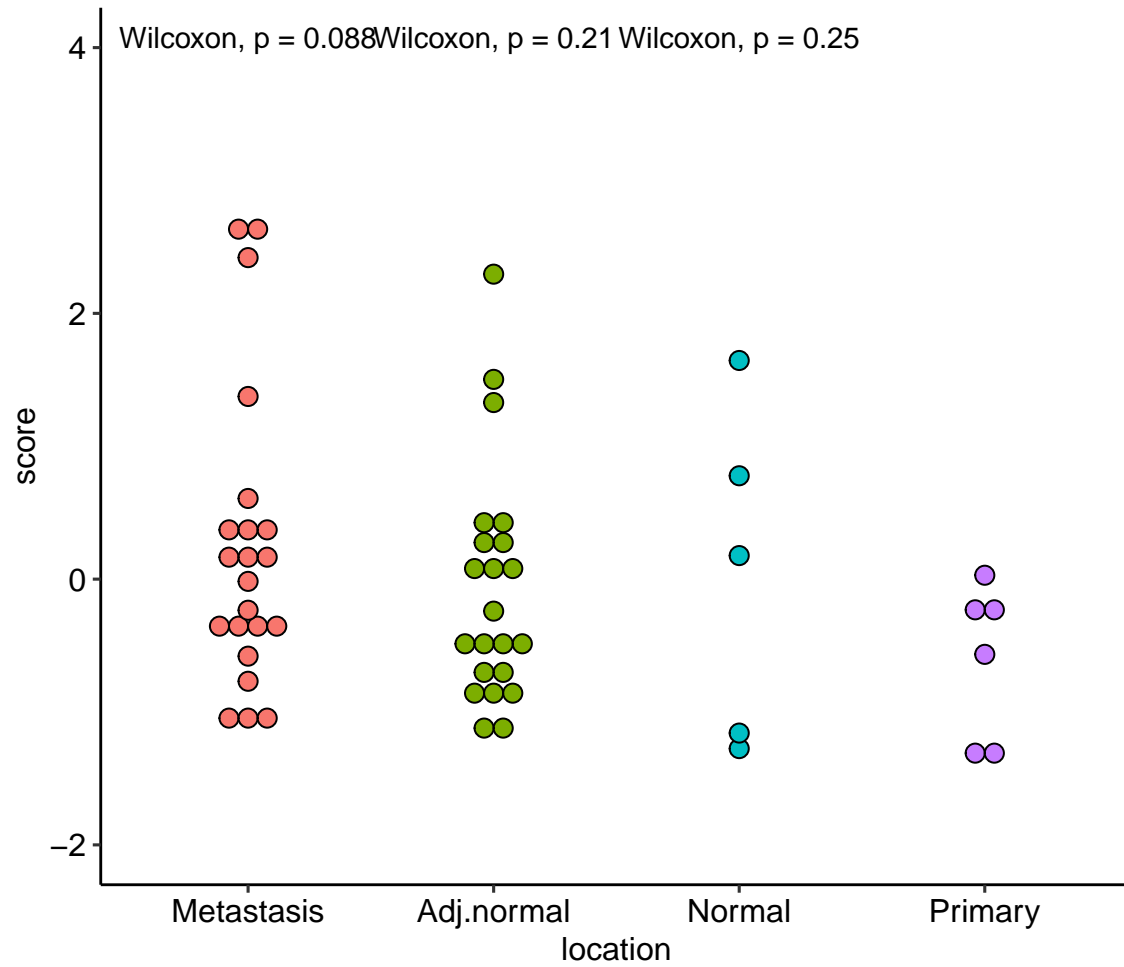
## Moffitt.F6\_BasalLike.top100 score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

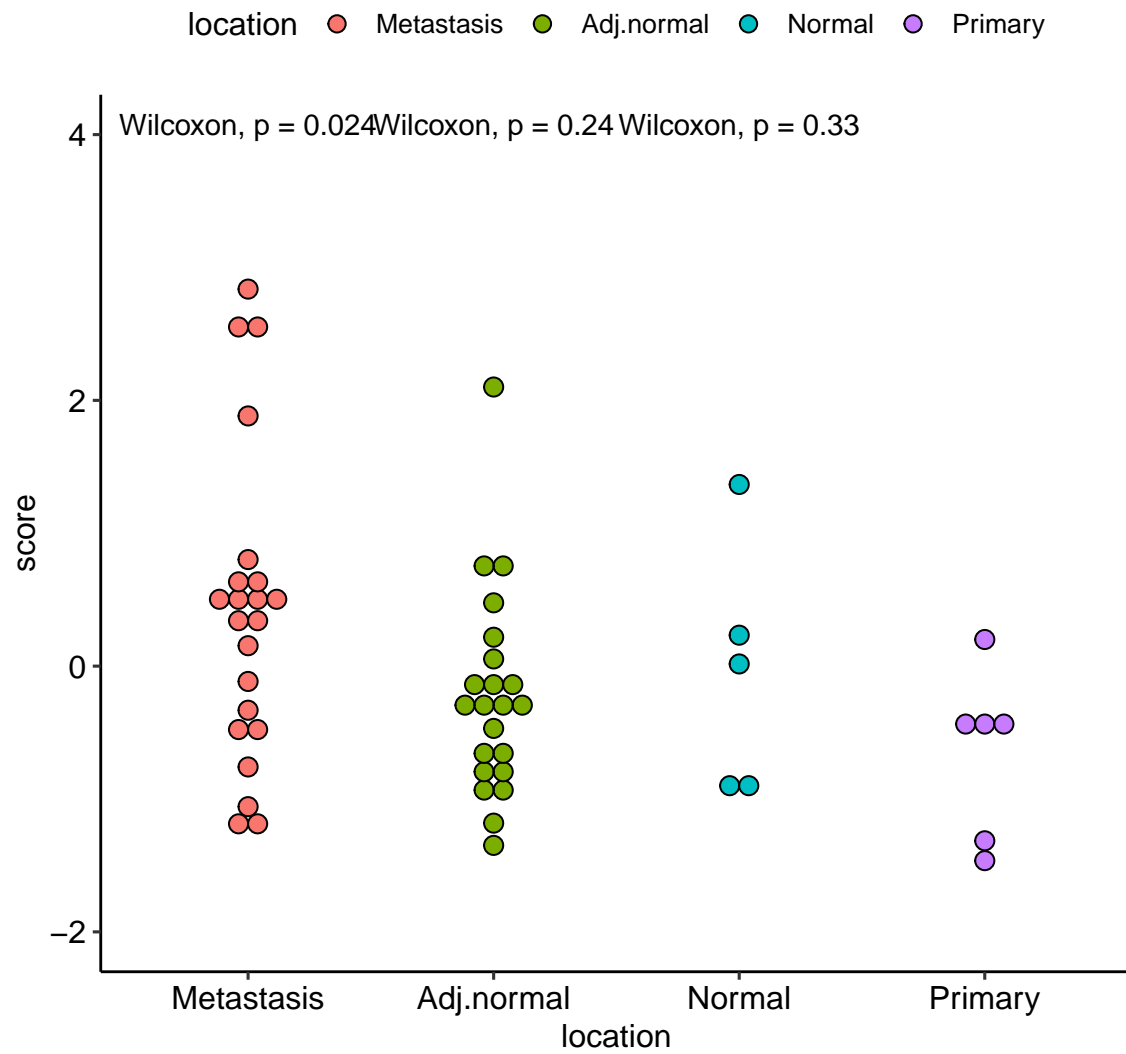


## Moffitt.F7\_Immune.top25 score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



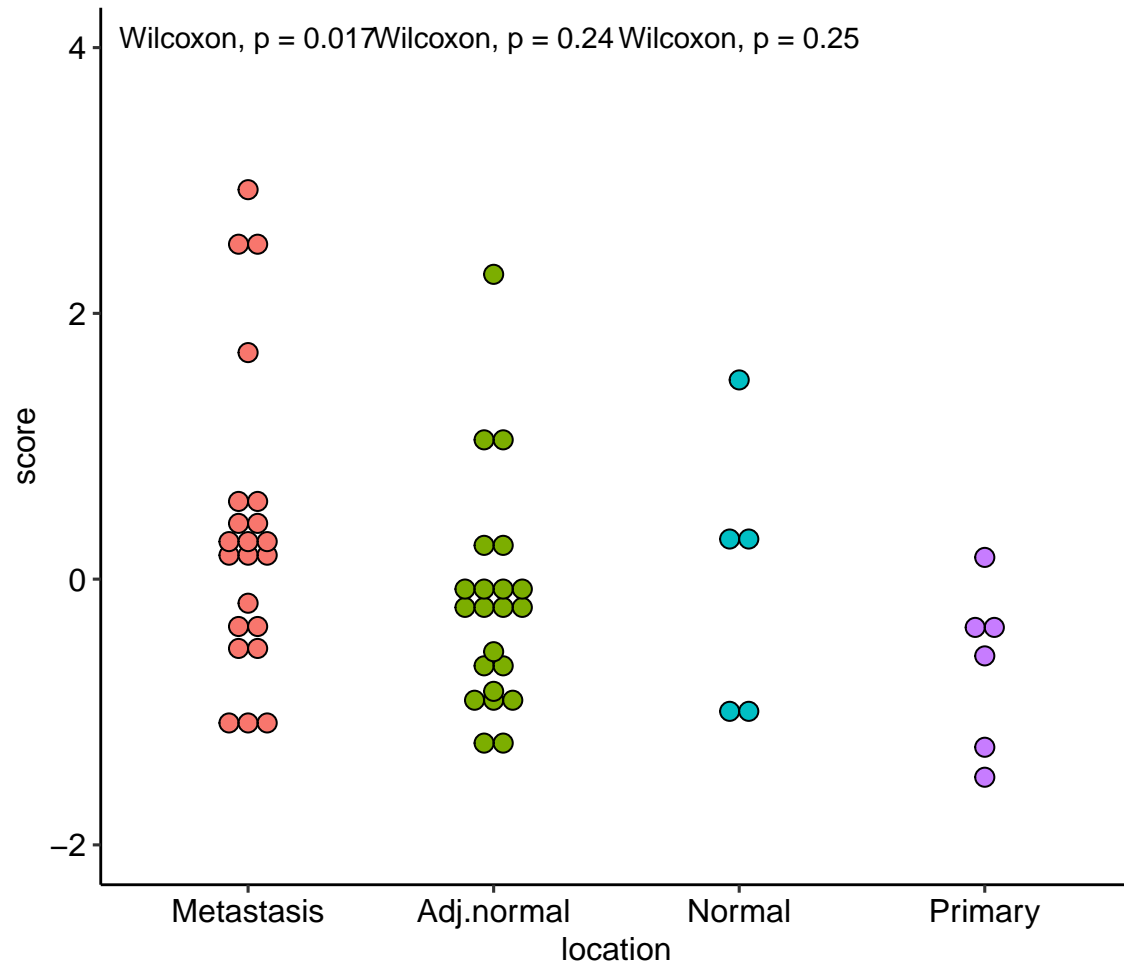
## Moffitt.F7\_Immune.top250 score



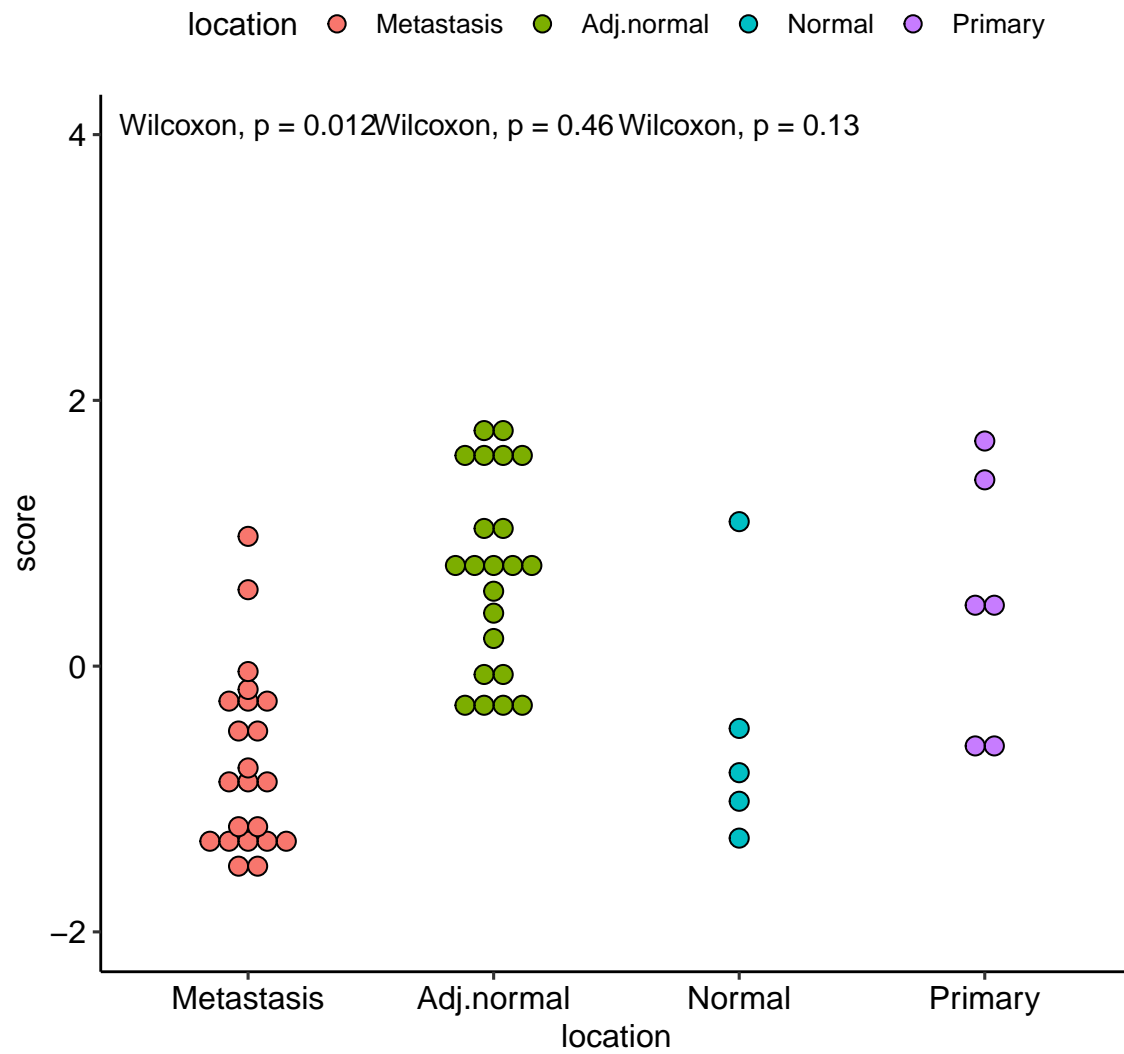


## Moffitt.F7\_Immune.top100 score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

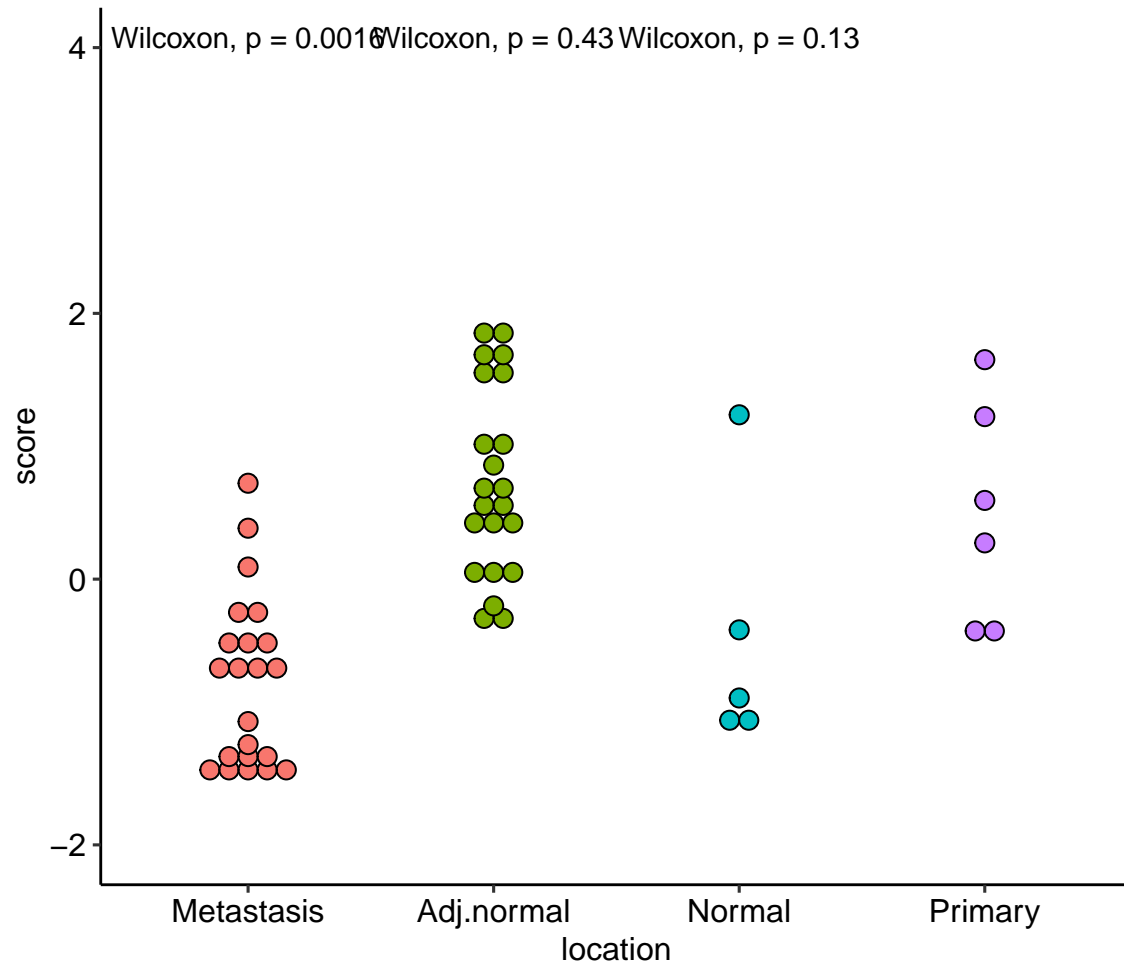


# Moffitt.F8\_Classical.top25 score



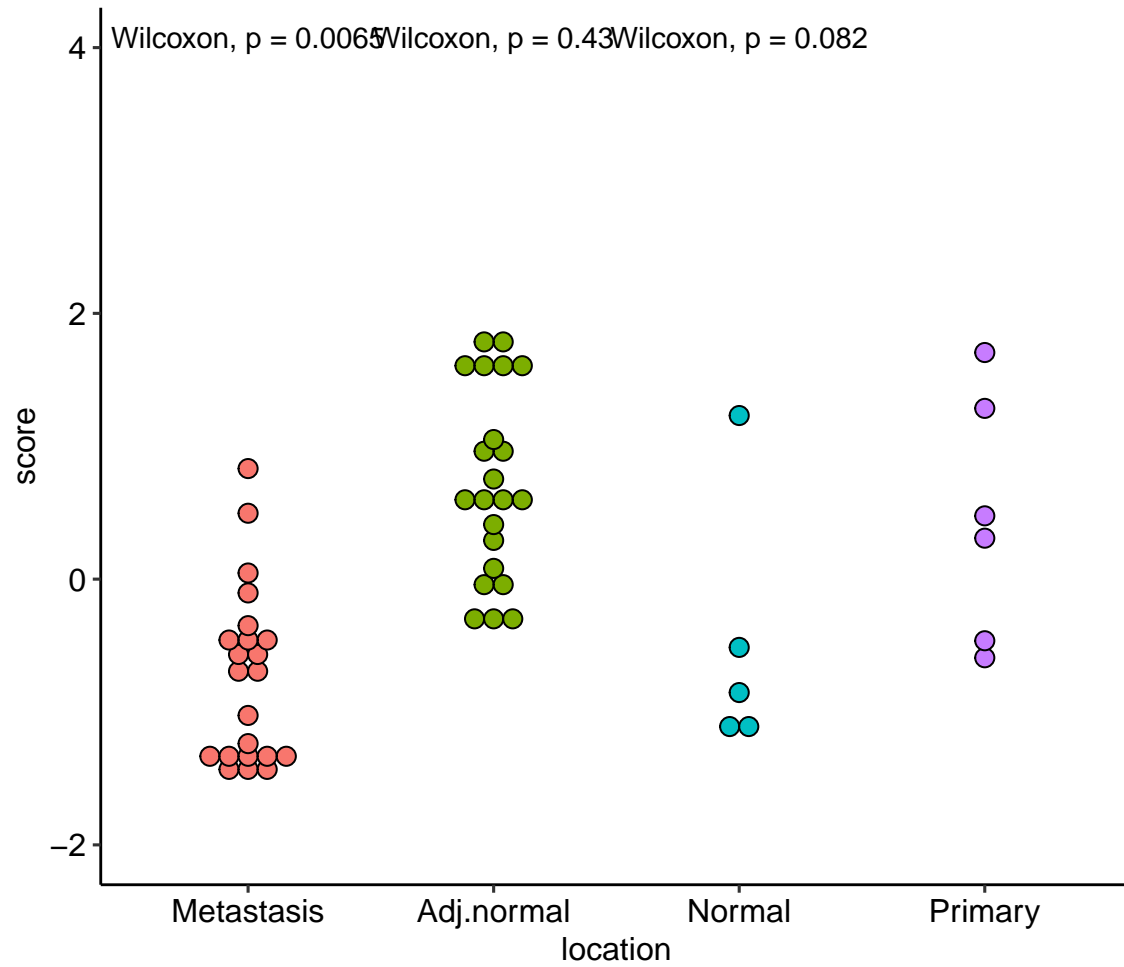
## Moffitt.F8\_Classical.top250 score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

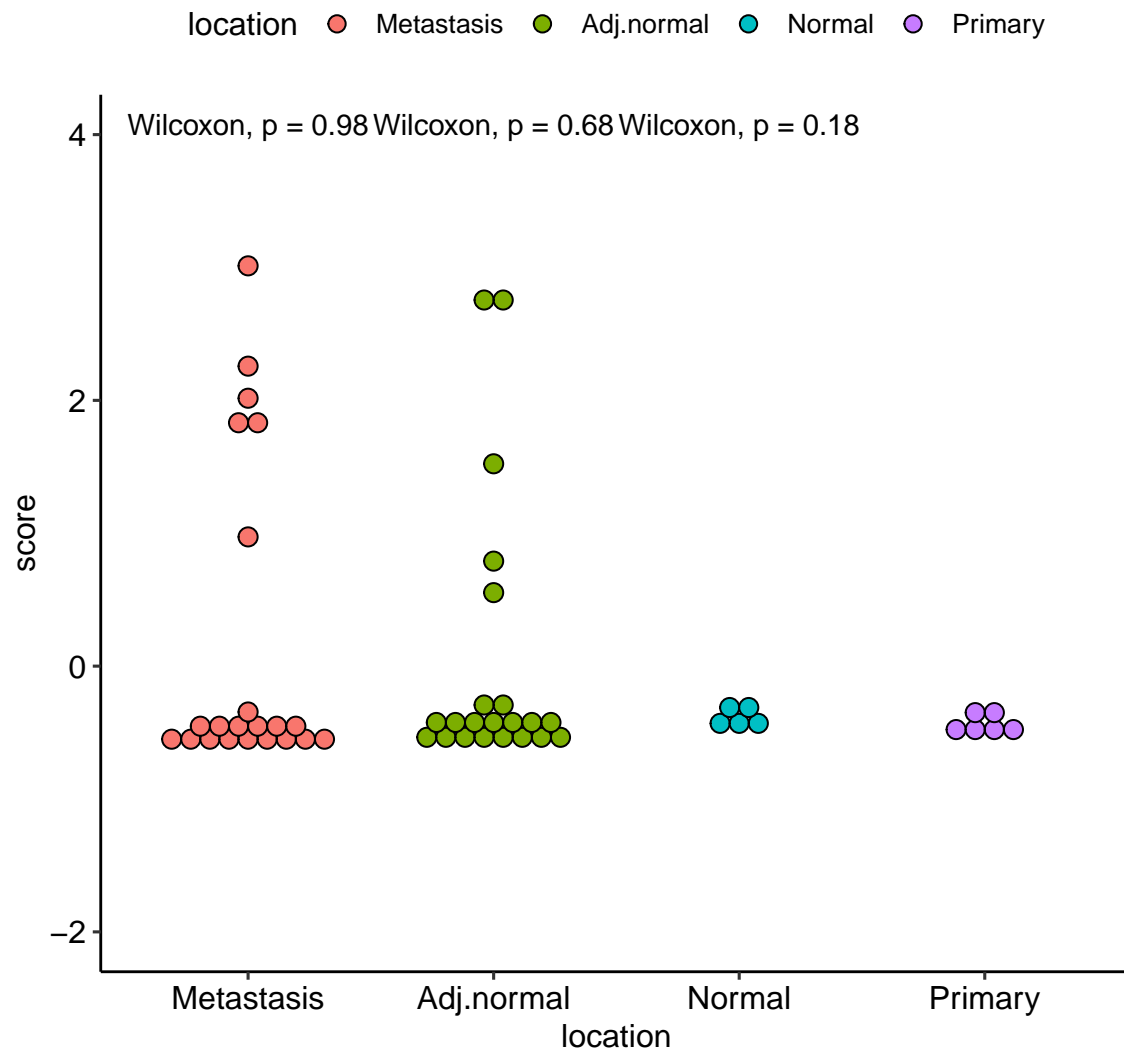


## Moffitt.F8\_Classical.top100 score

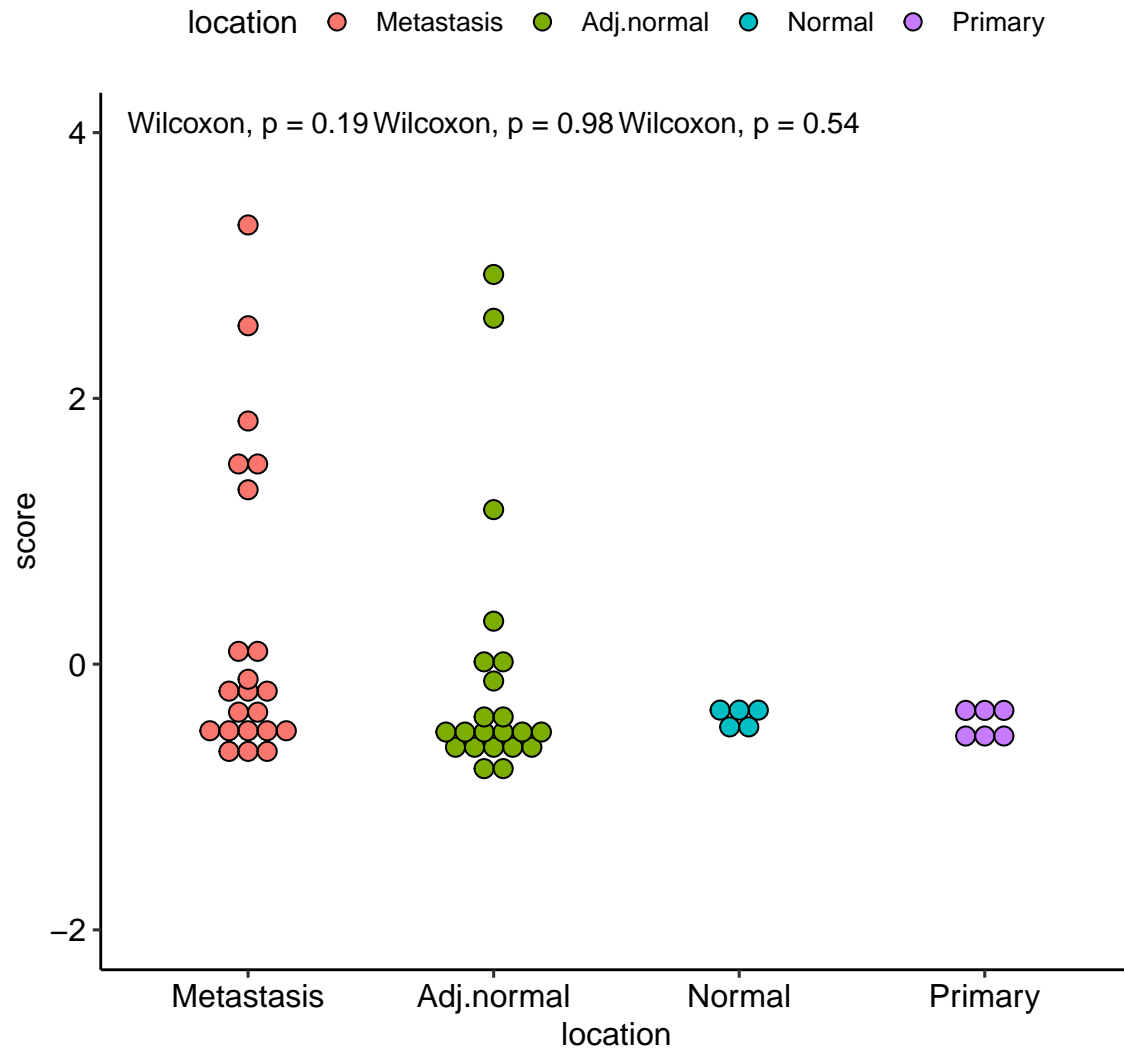
location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



## Moffitt.F9\_Lung.top25 score

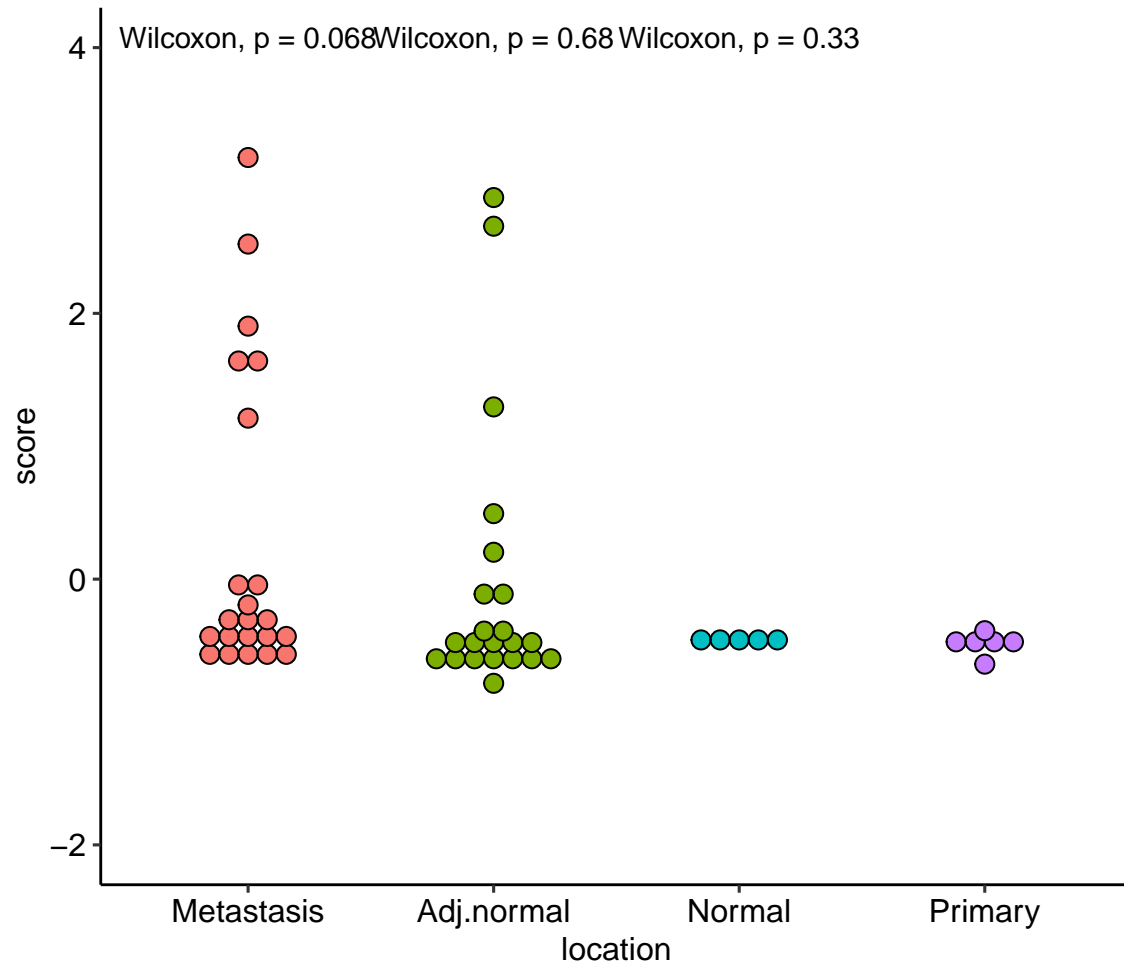


## Moffitt.F9\_Lung.top250 score

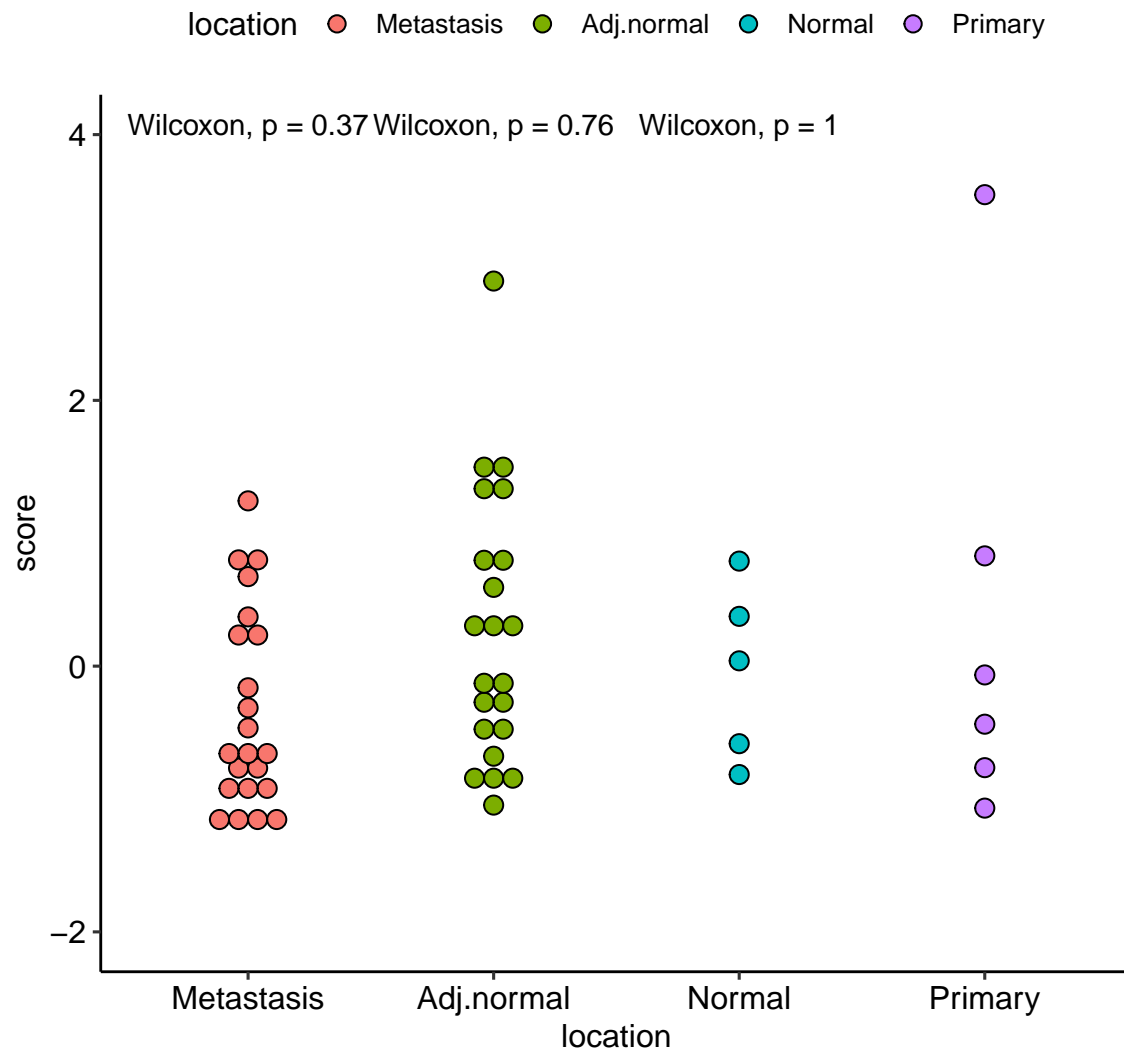


## Moffitt.F9\_Lung.top100 score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



## Moffitt.F10.top25 score

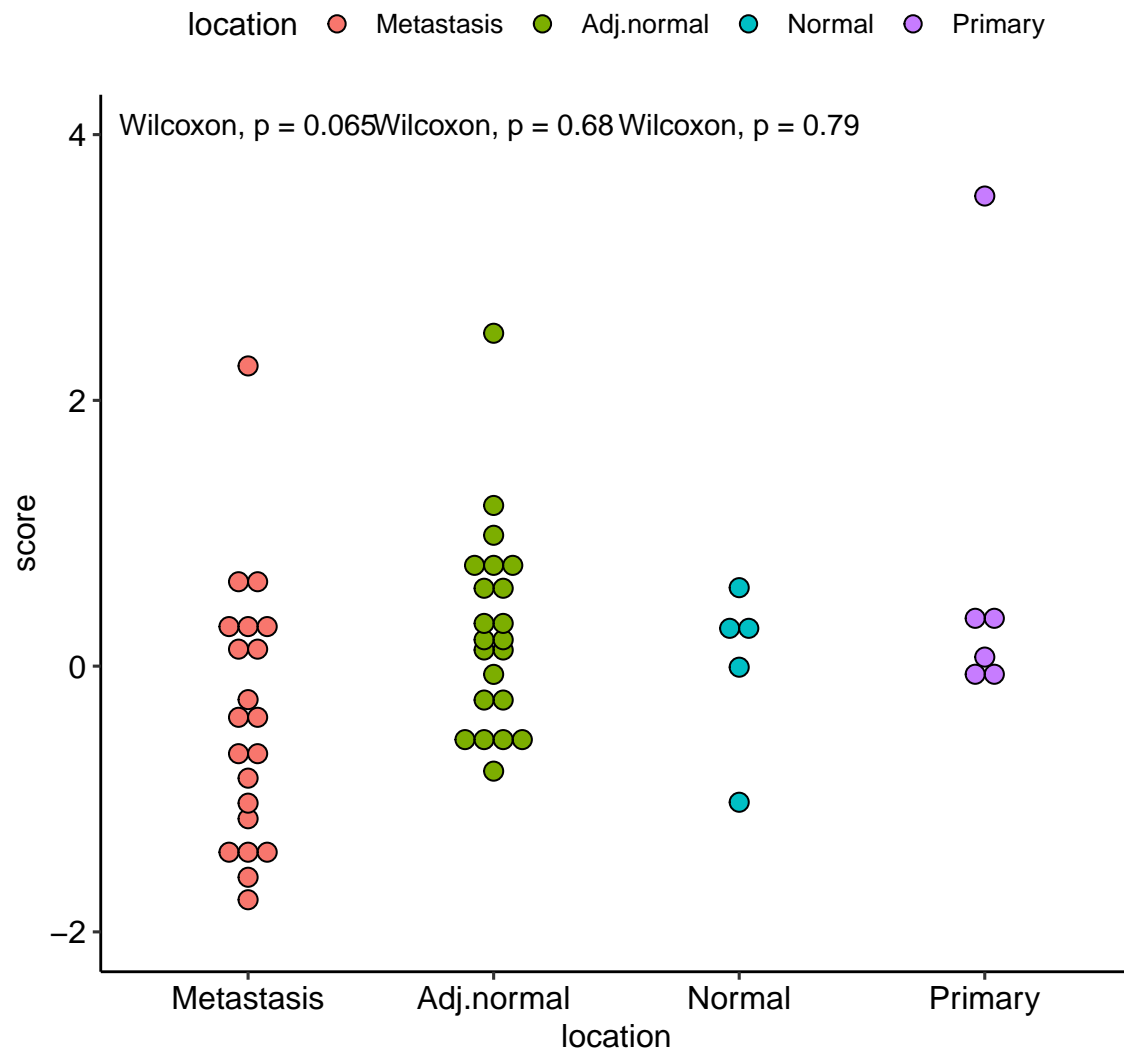


```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

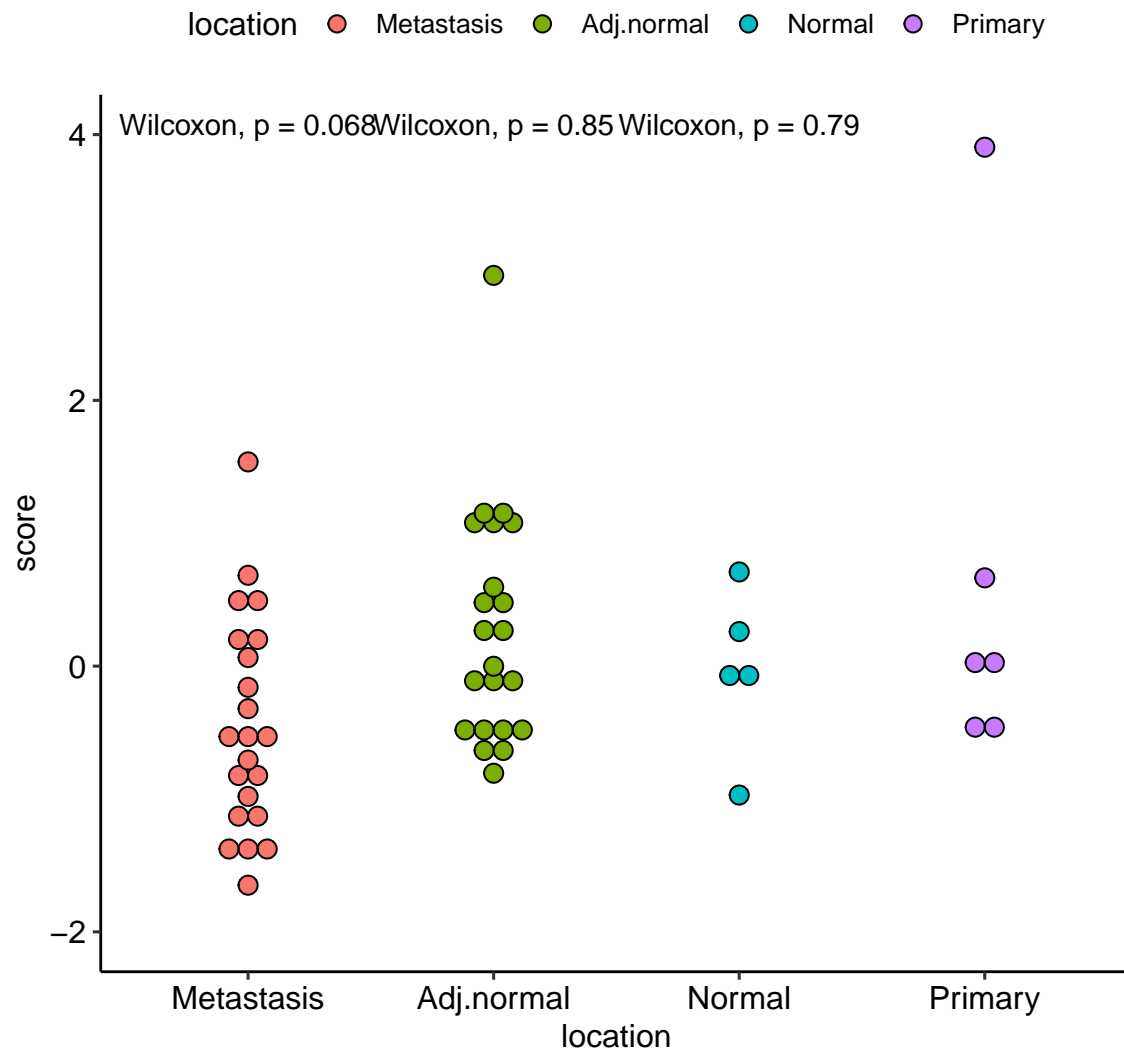
```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```



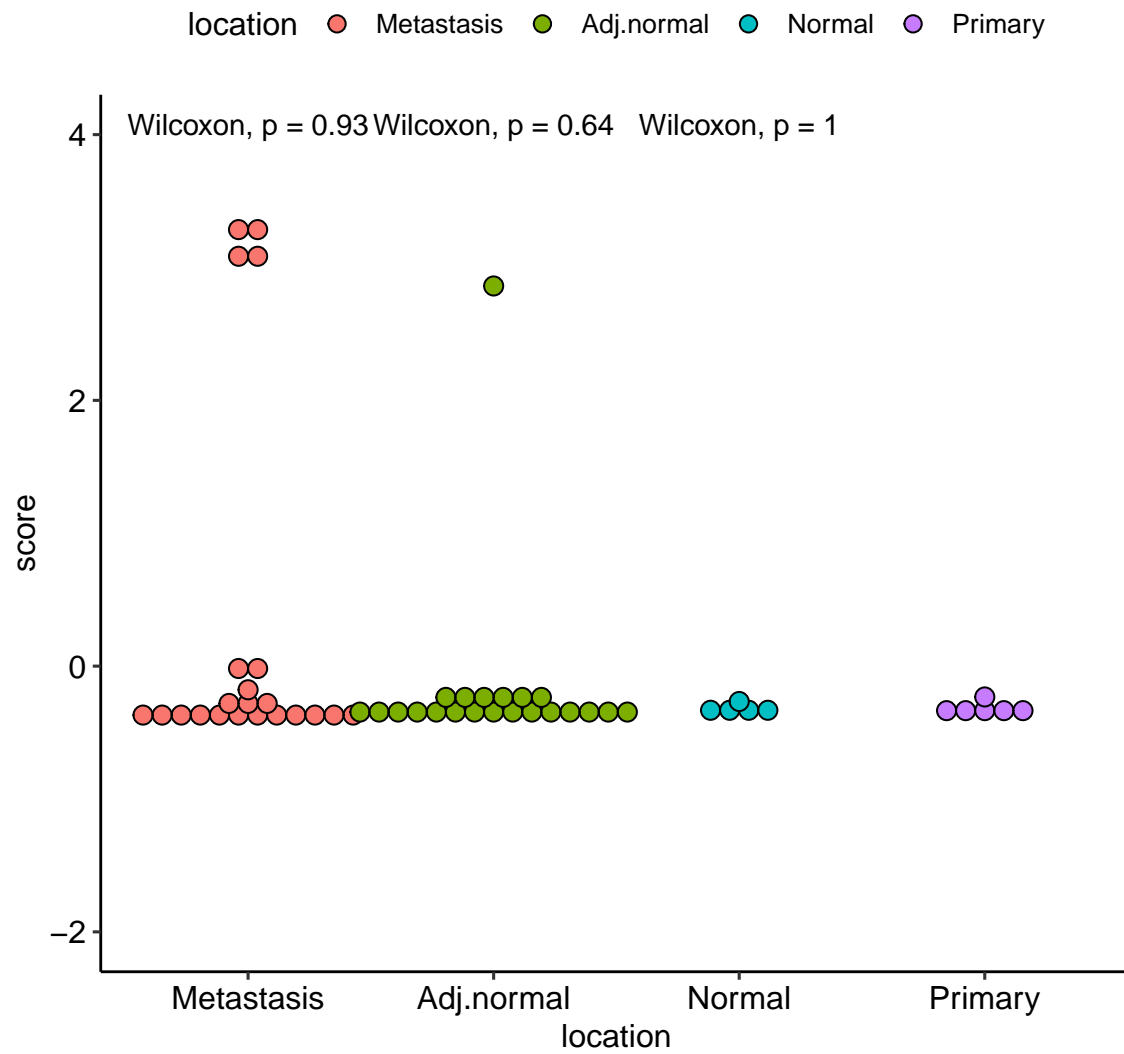
## Moffitt.F10.top250 score



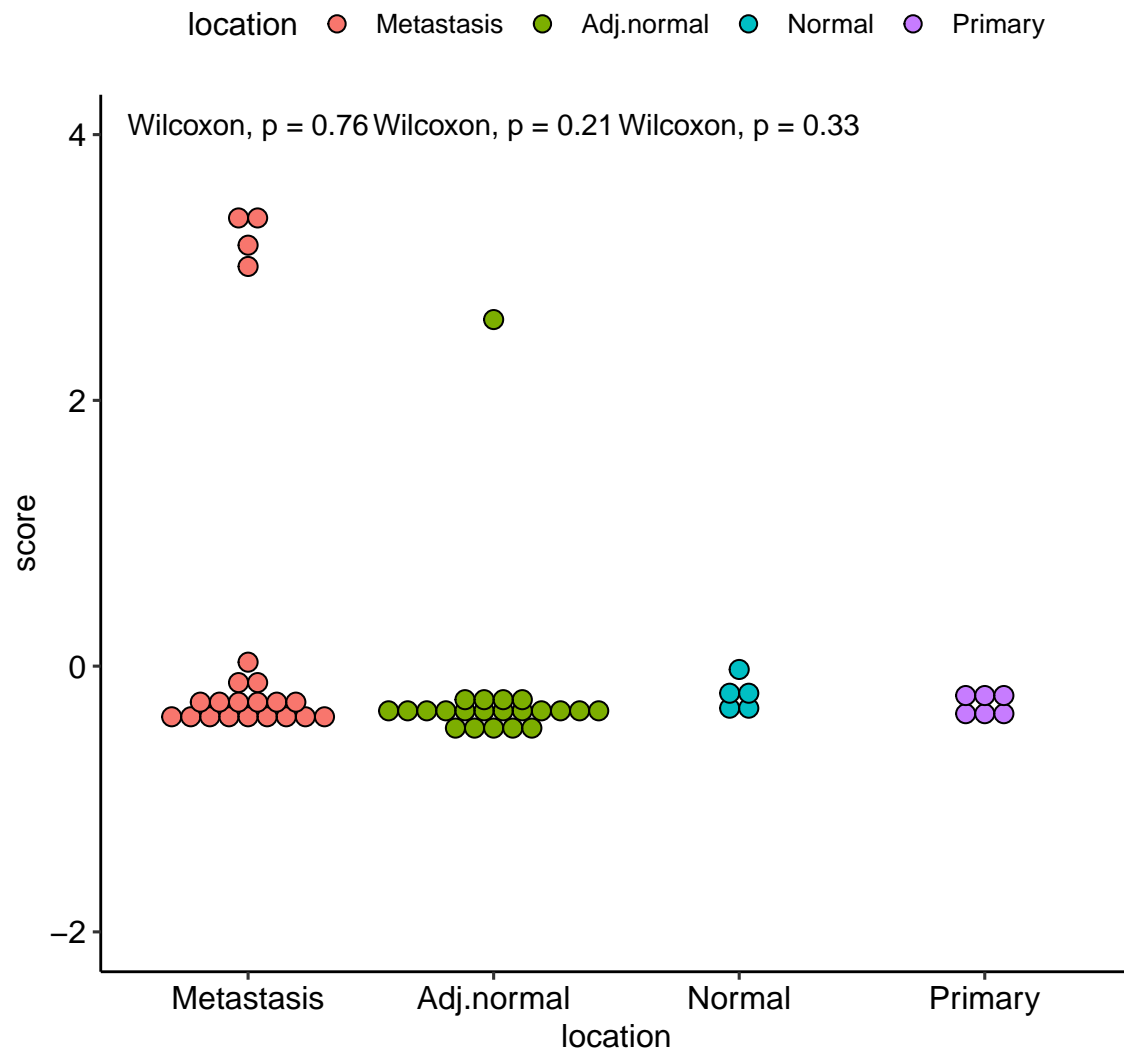
## Moffitt.F10.top100 score



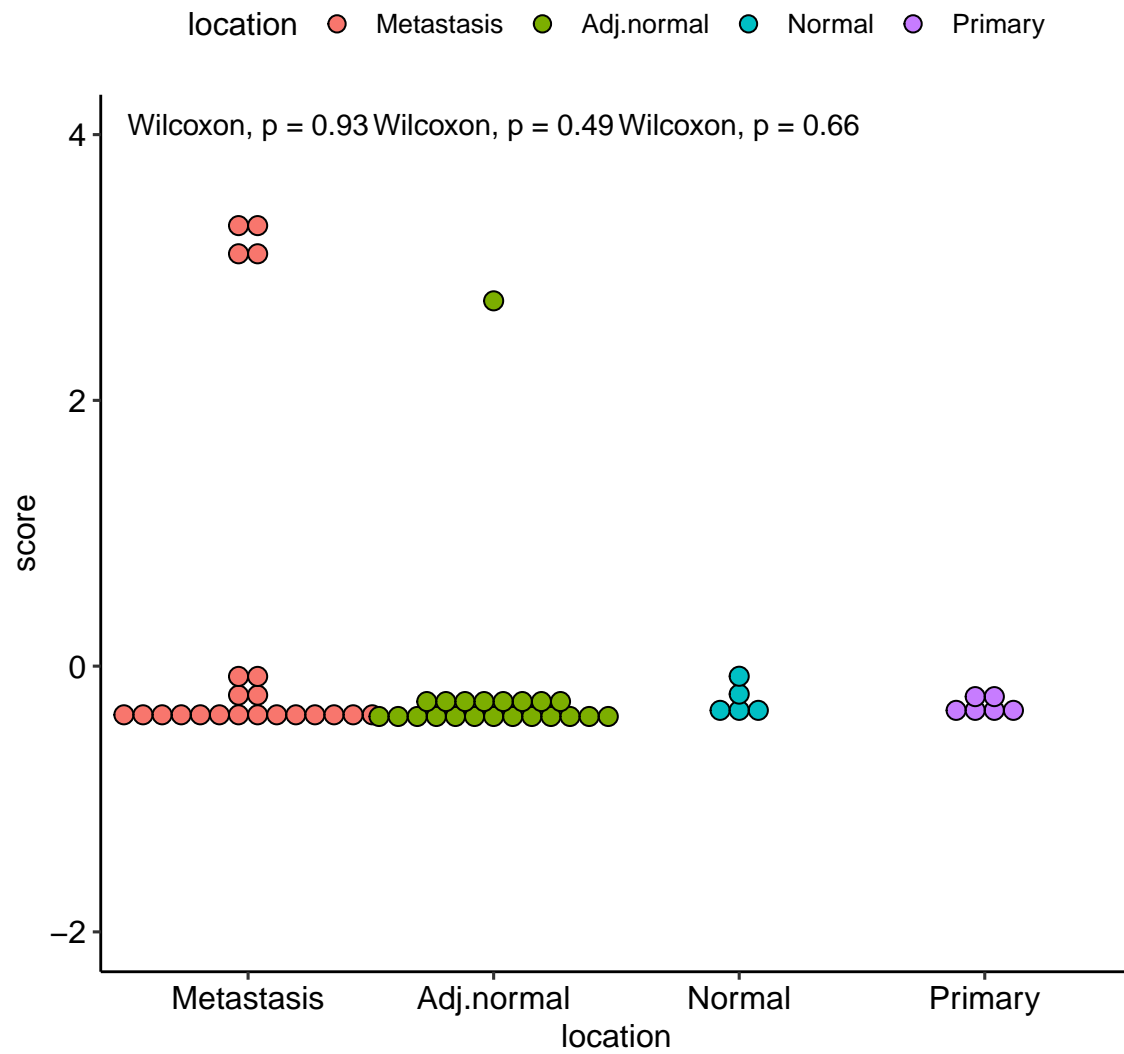
## Moffitt.F11\_Muscle.top25 score



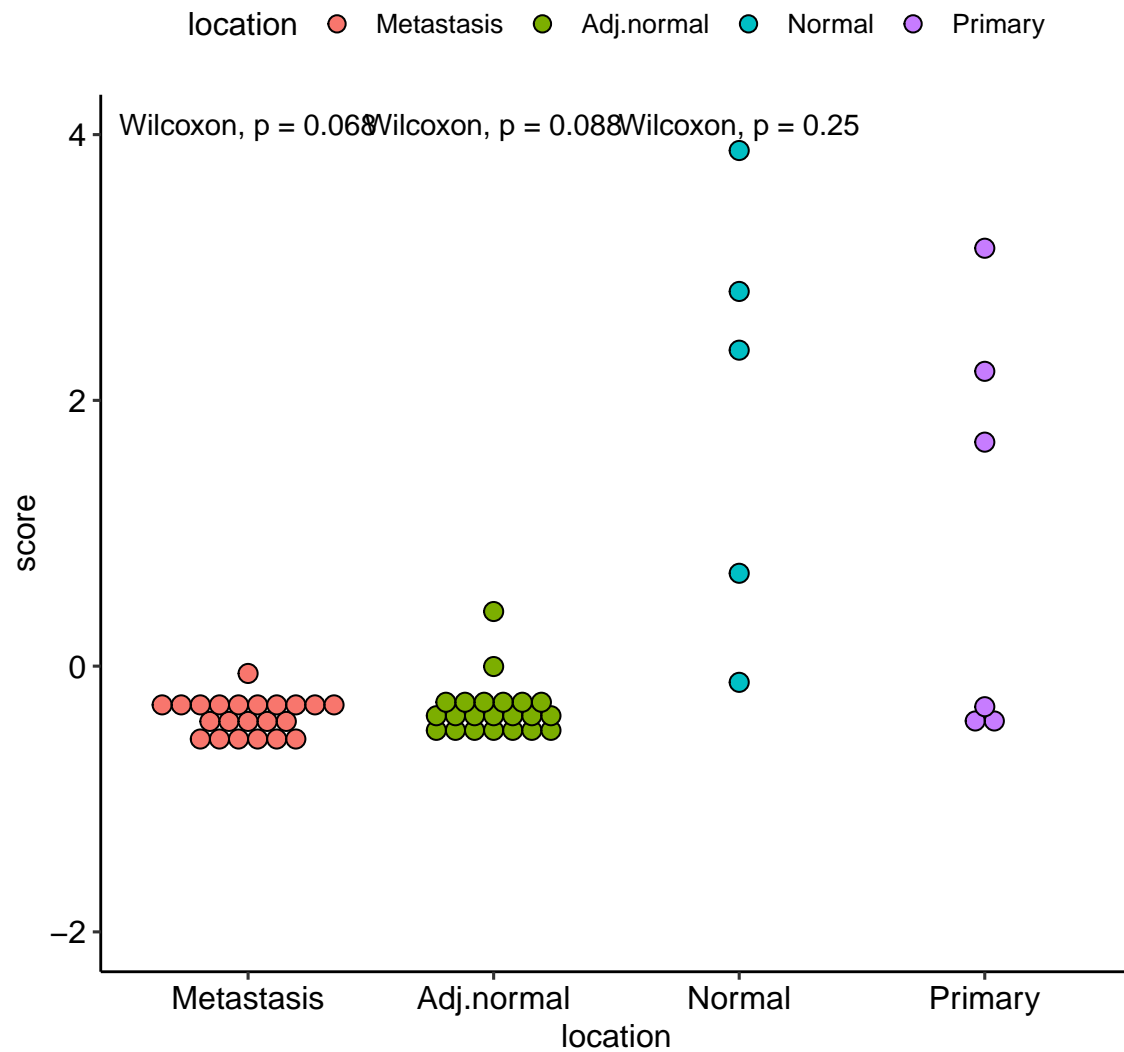
## Moffitt.F11\_Muscle.top250 score



## Moffitt.F11\_Muscle.top100 score



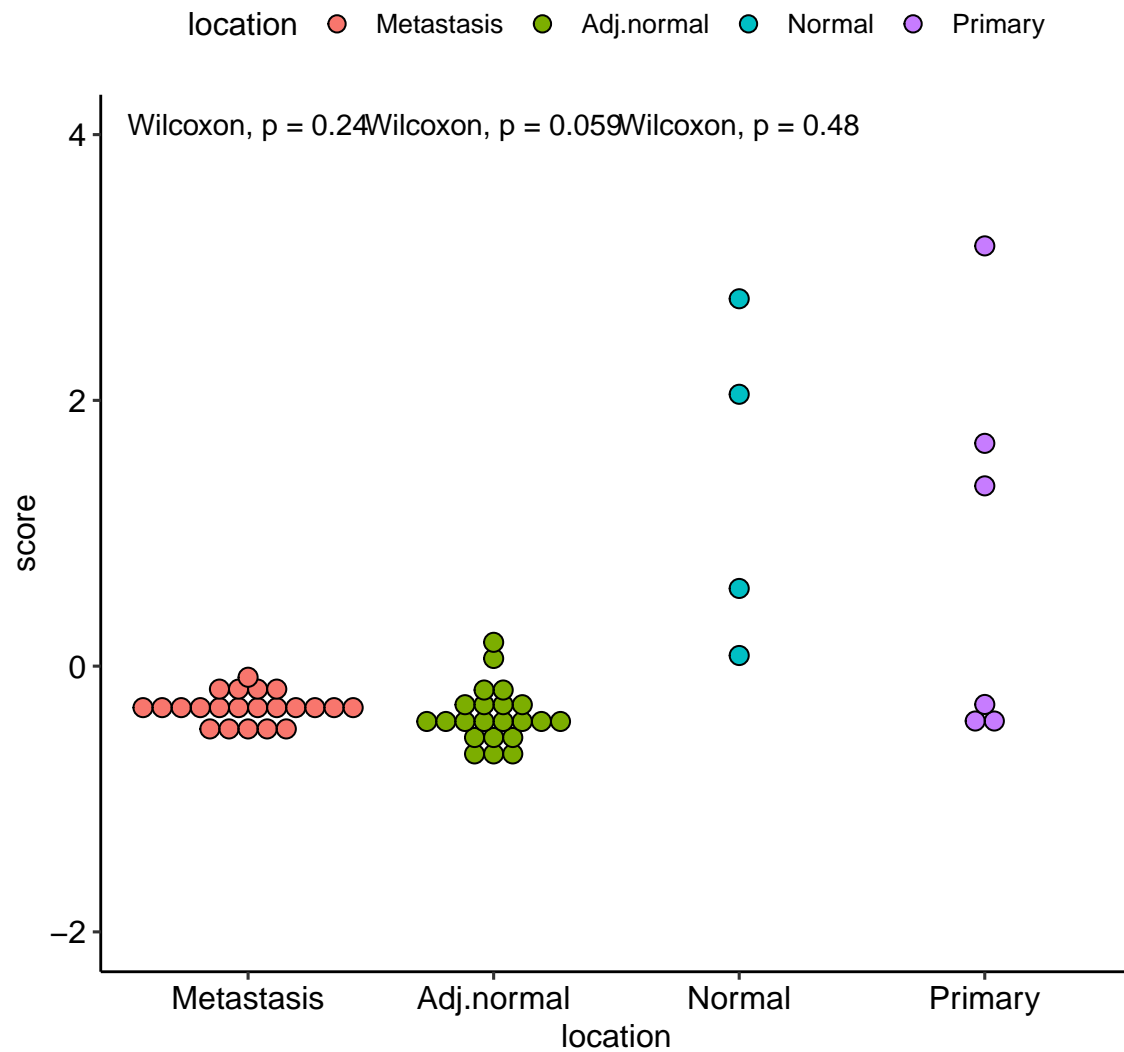
## Moffitt.F12\_Endocrine.top25 score



```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

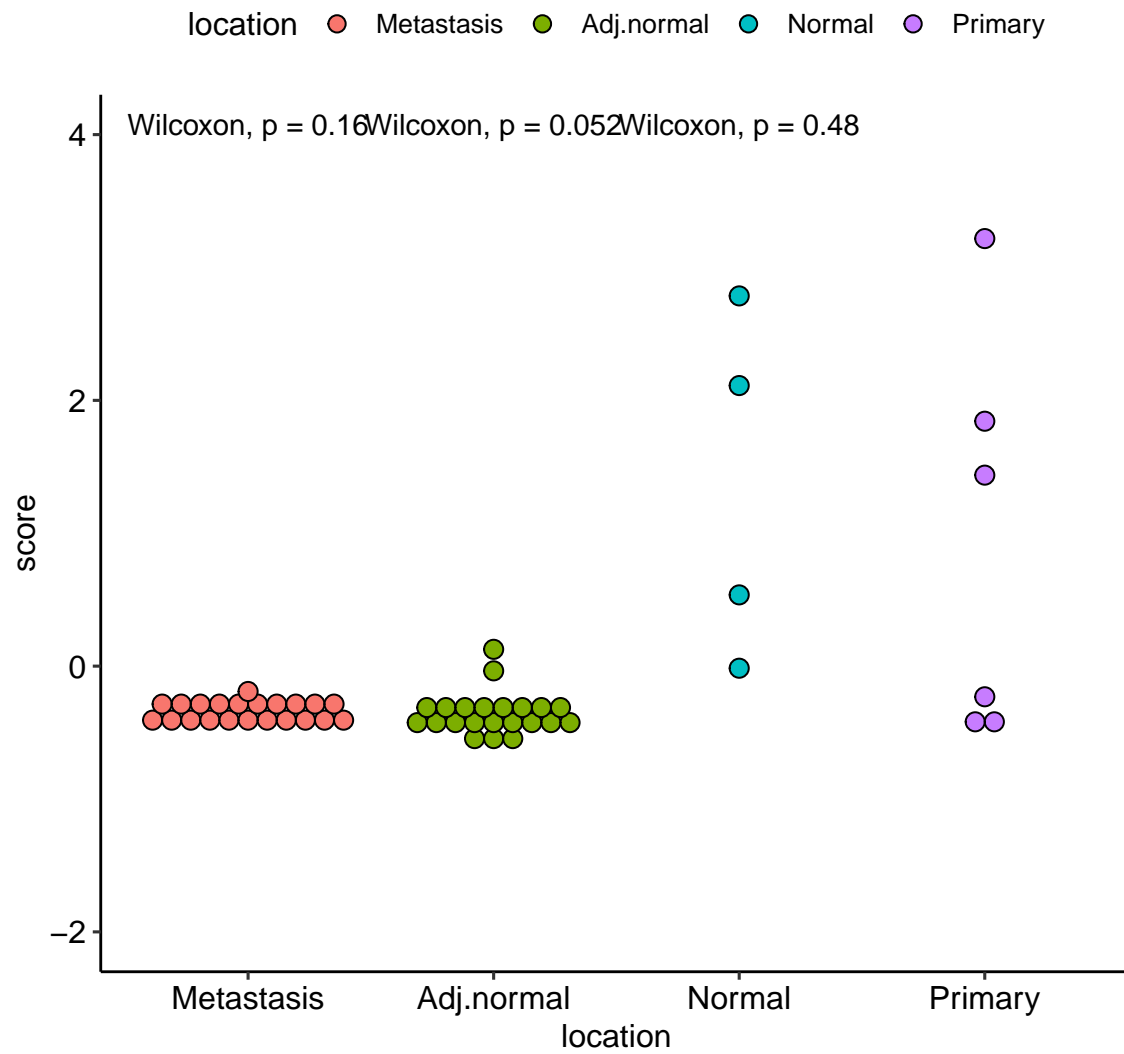
## Moffitt.F12\_Endocrine.top250 score



```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

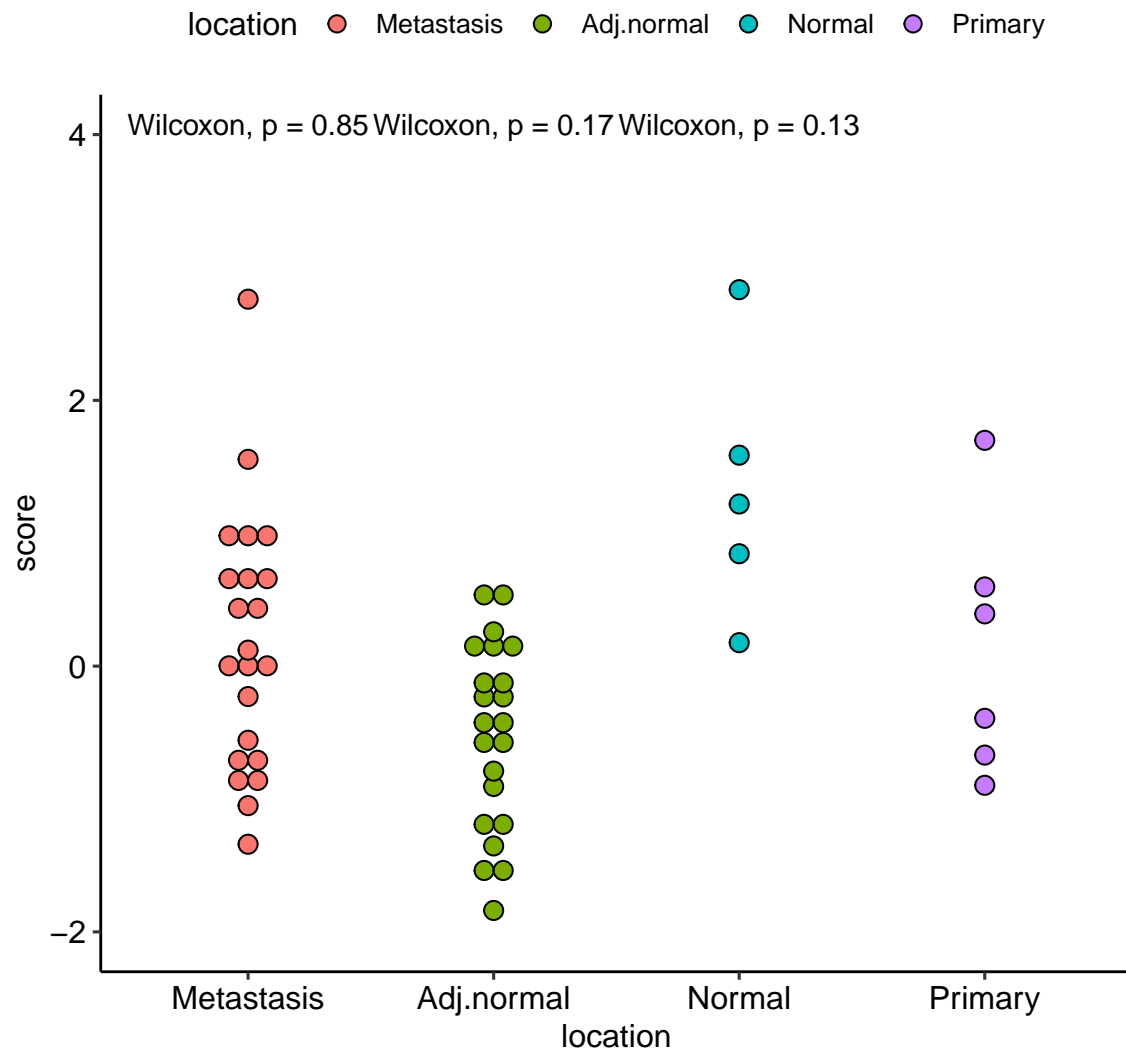
```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

## Moffitt.F12\_Endocrine.top100 score

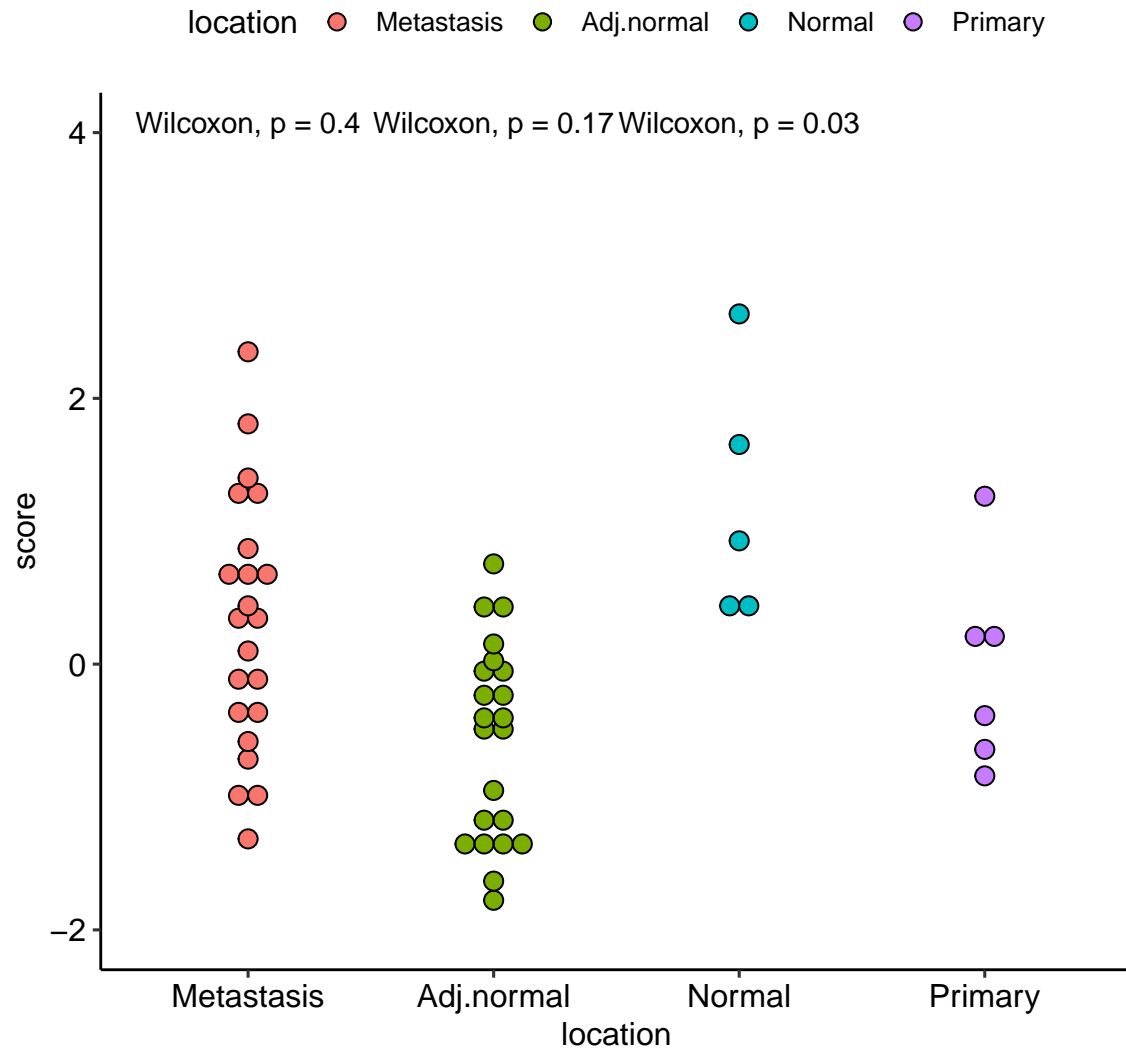




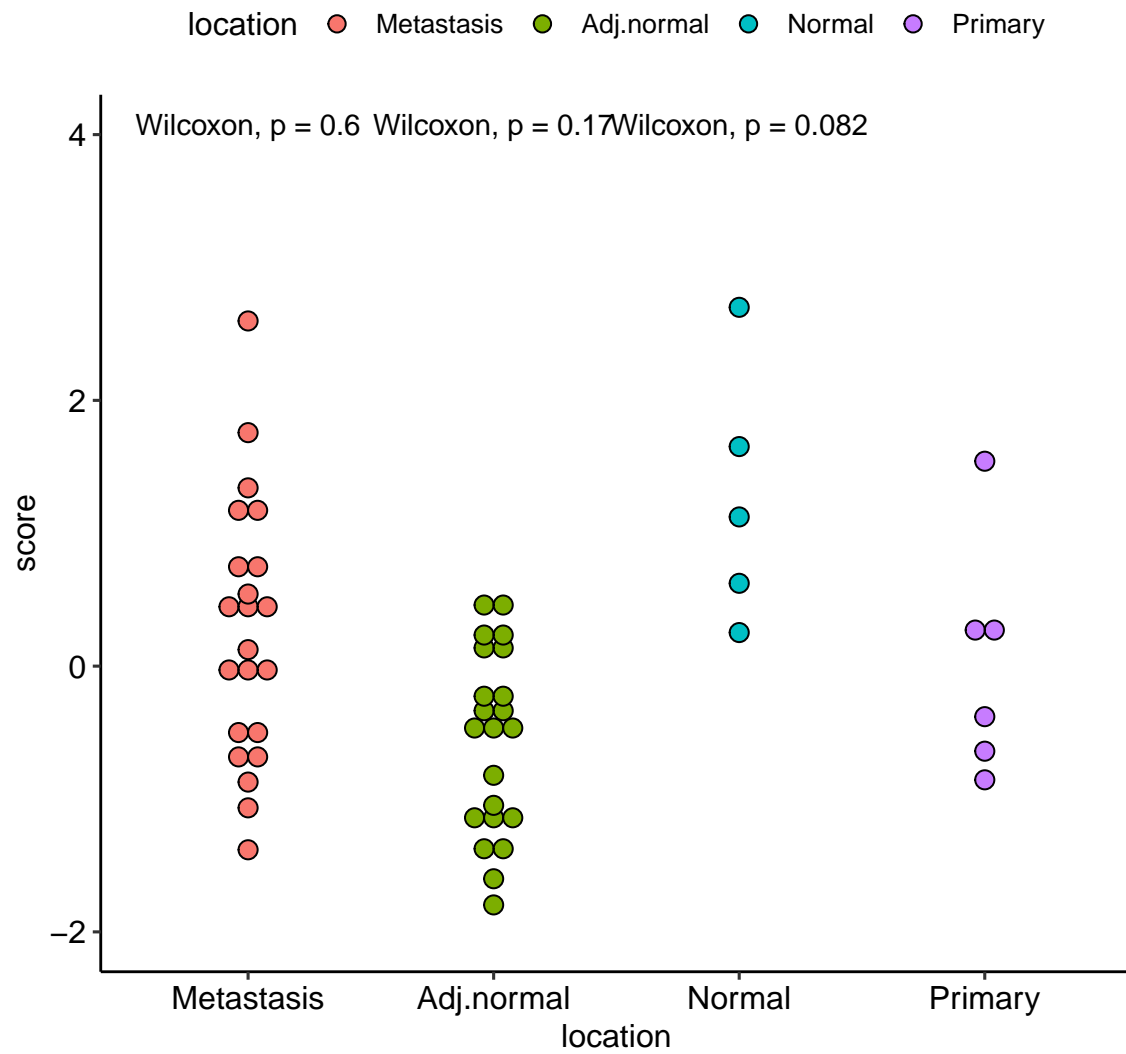
# Moffitt.F13\_NormalStroma.top25 score



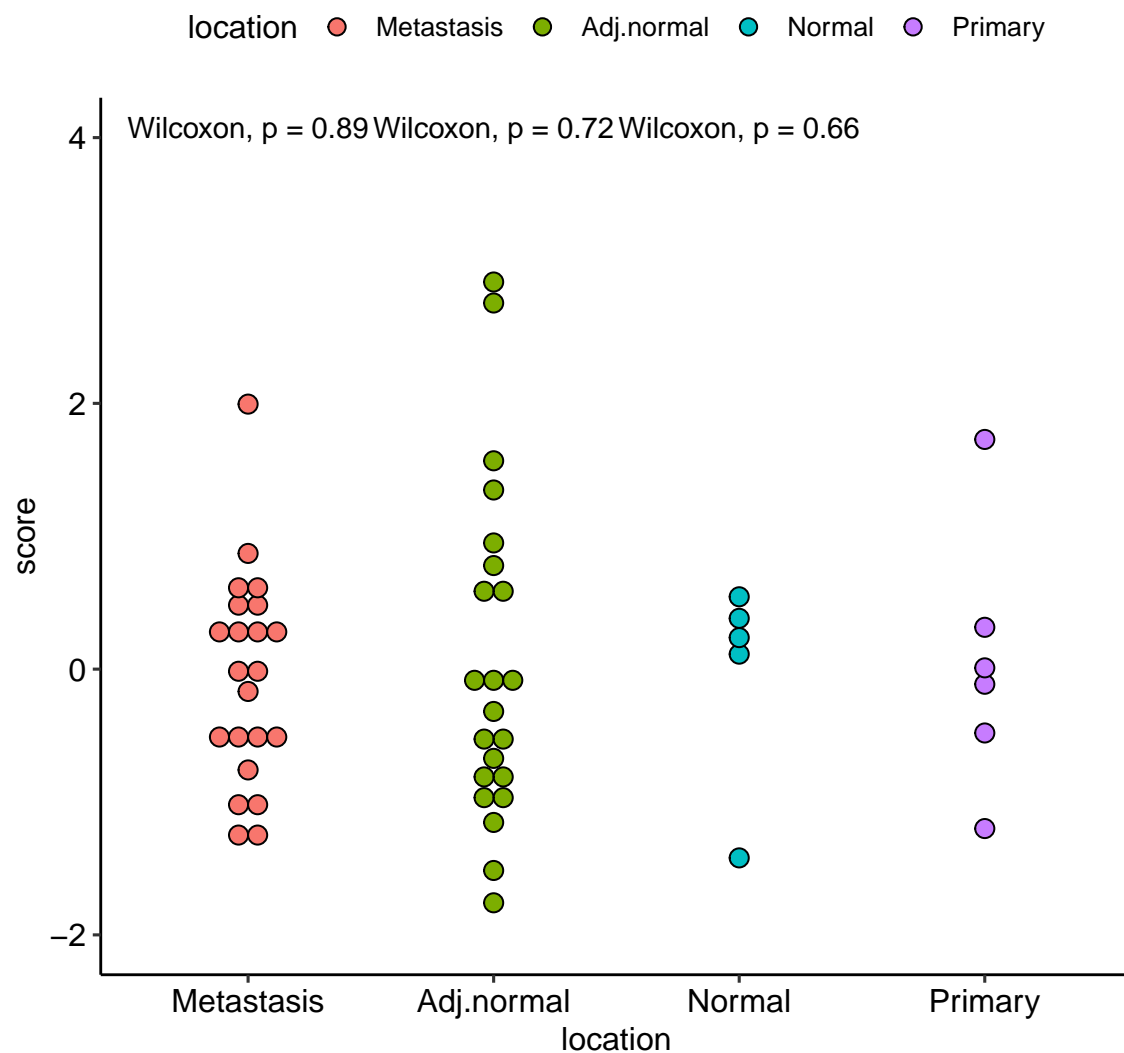
# Moffitt.F13\_NormalStroma.top250 score



# Moffitt.F13\_NormalStroma.top100 score



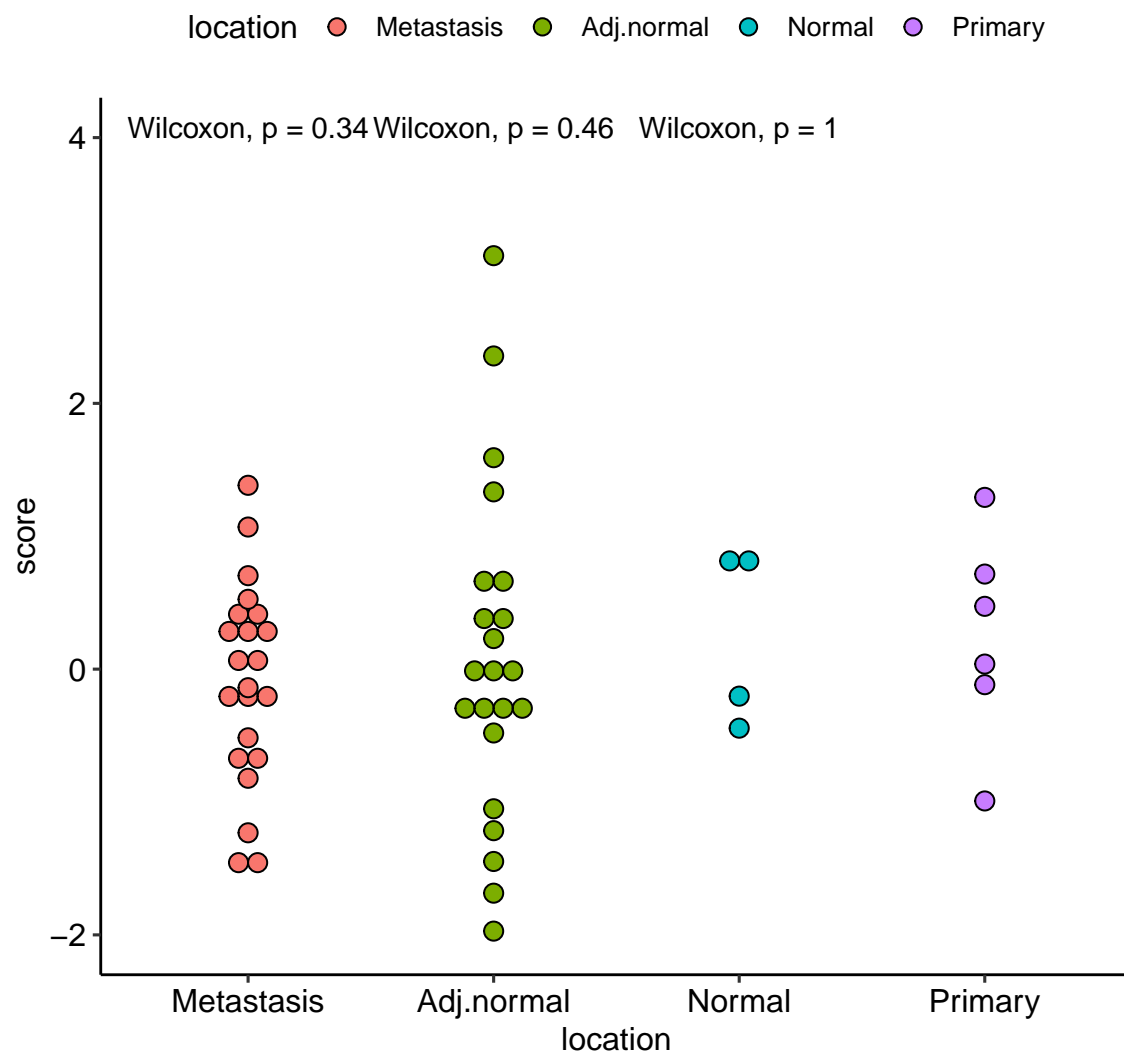
## Moffitt.F14.top25 score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

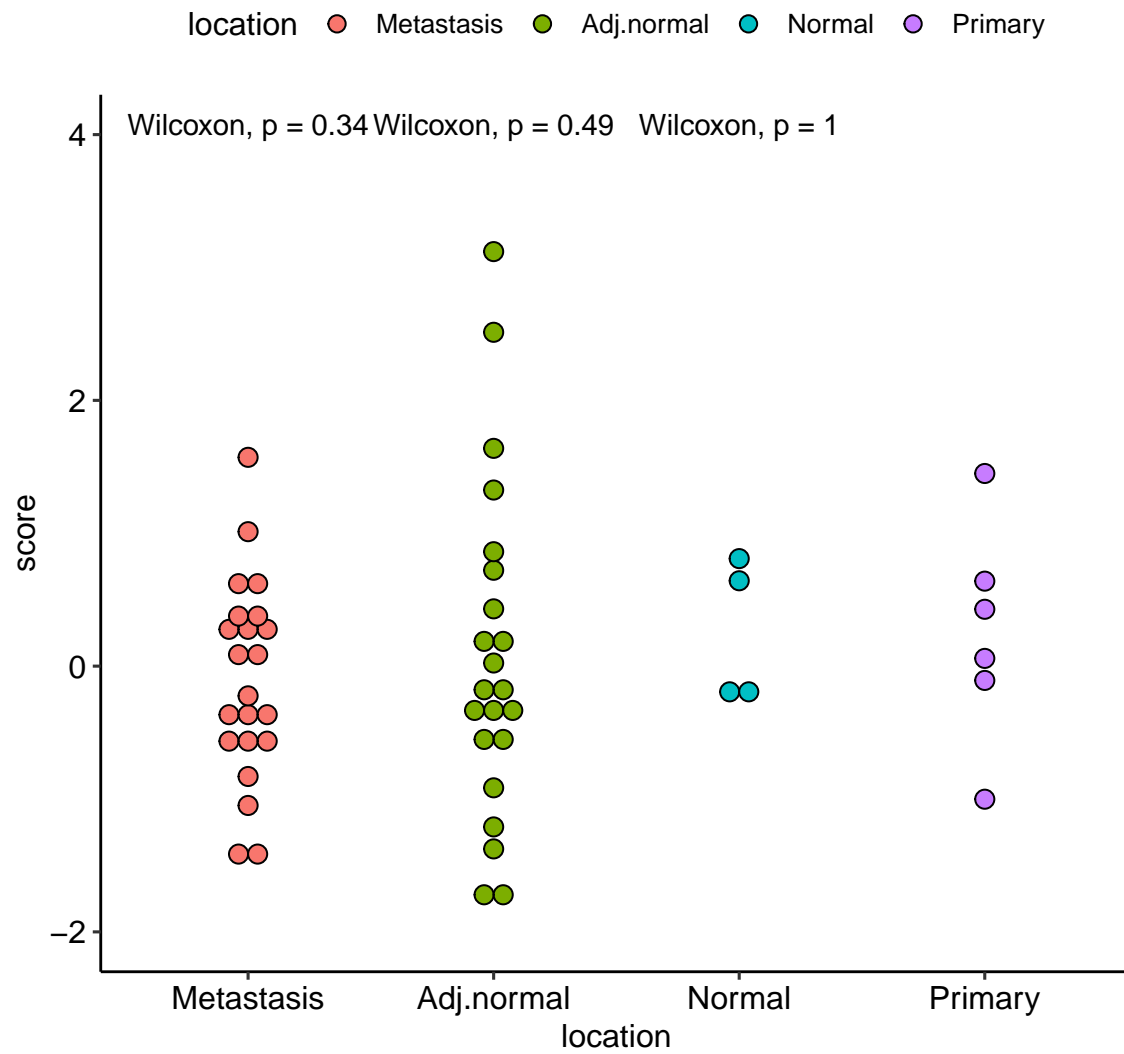
## Moffitt.F14.top250 score



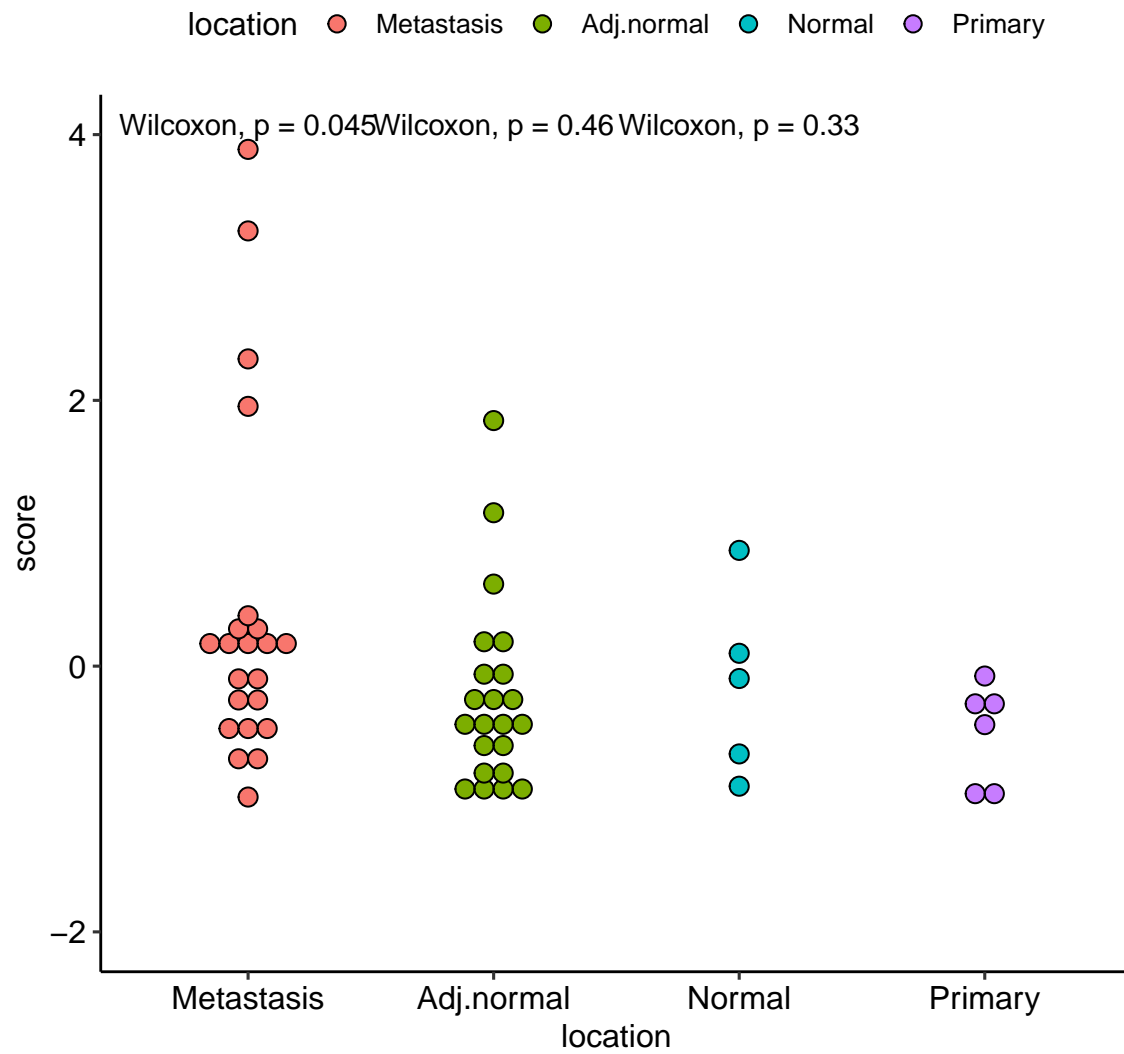
## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

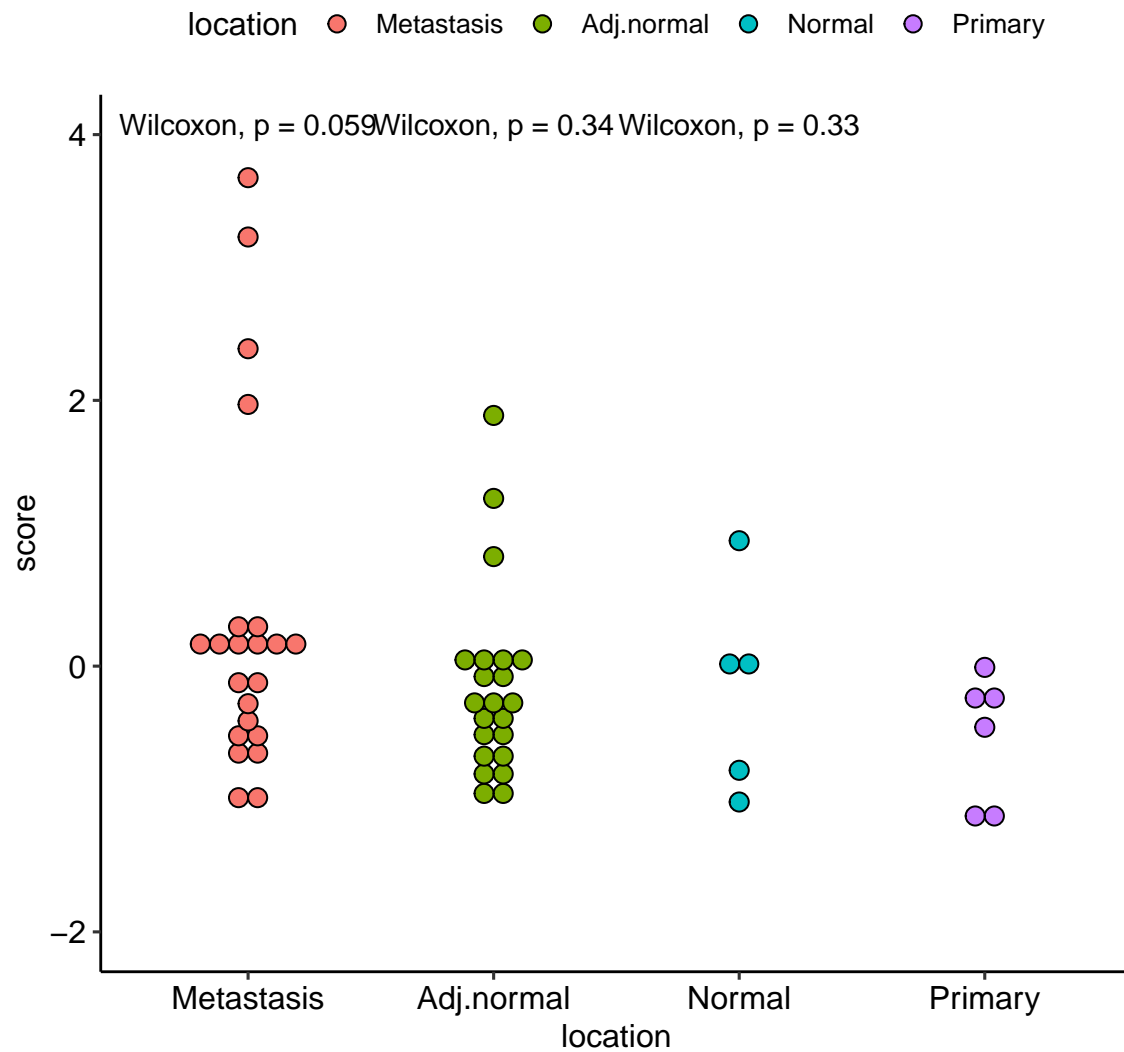
## Moffitt.F14.top100 score



## CIBERSORT.B.cells.naive score



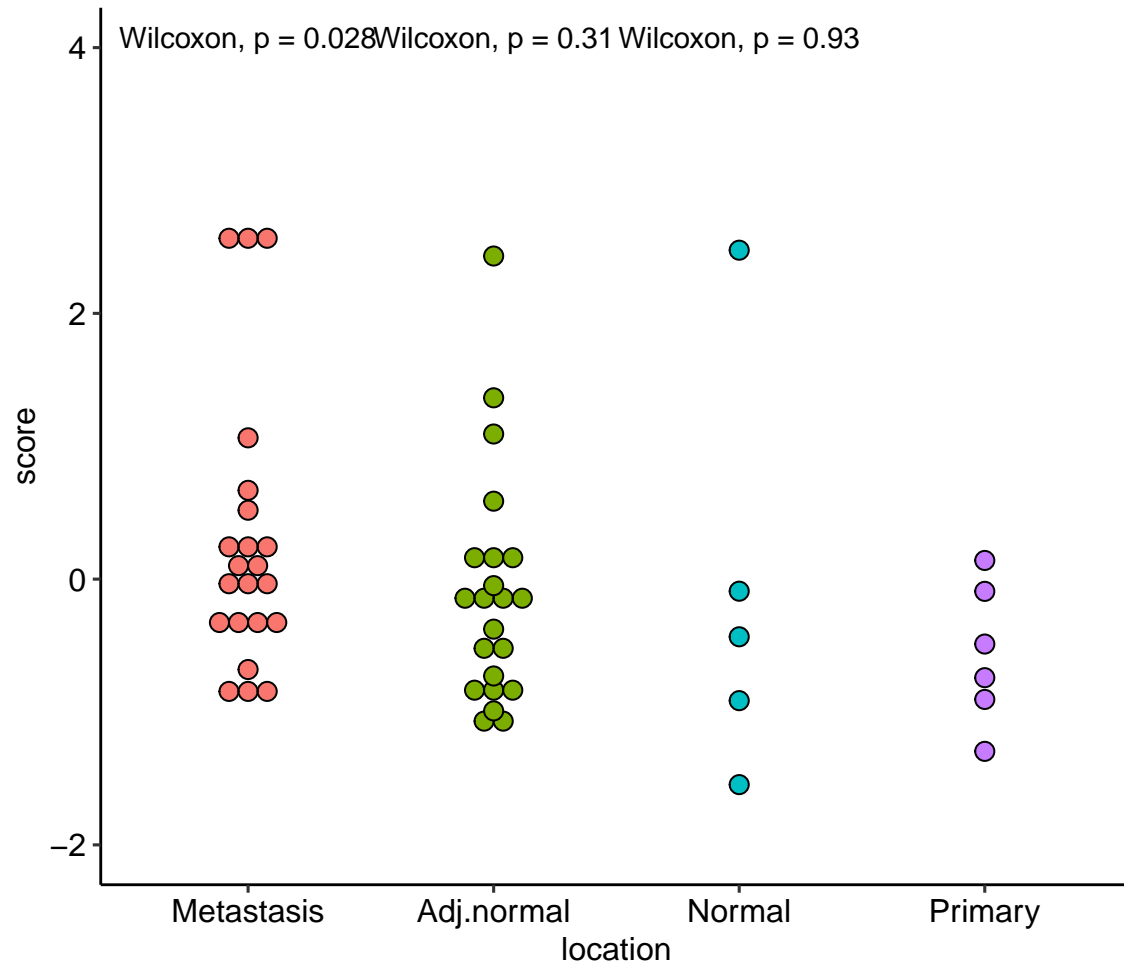
CIBERSORT.B.cells.memory score



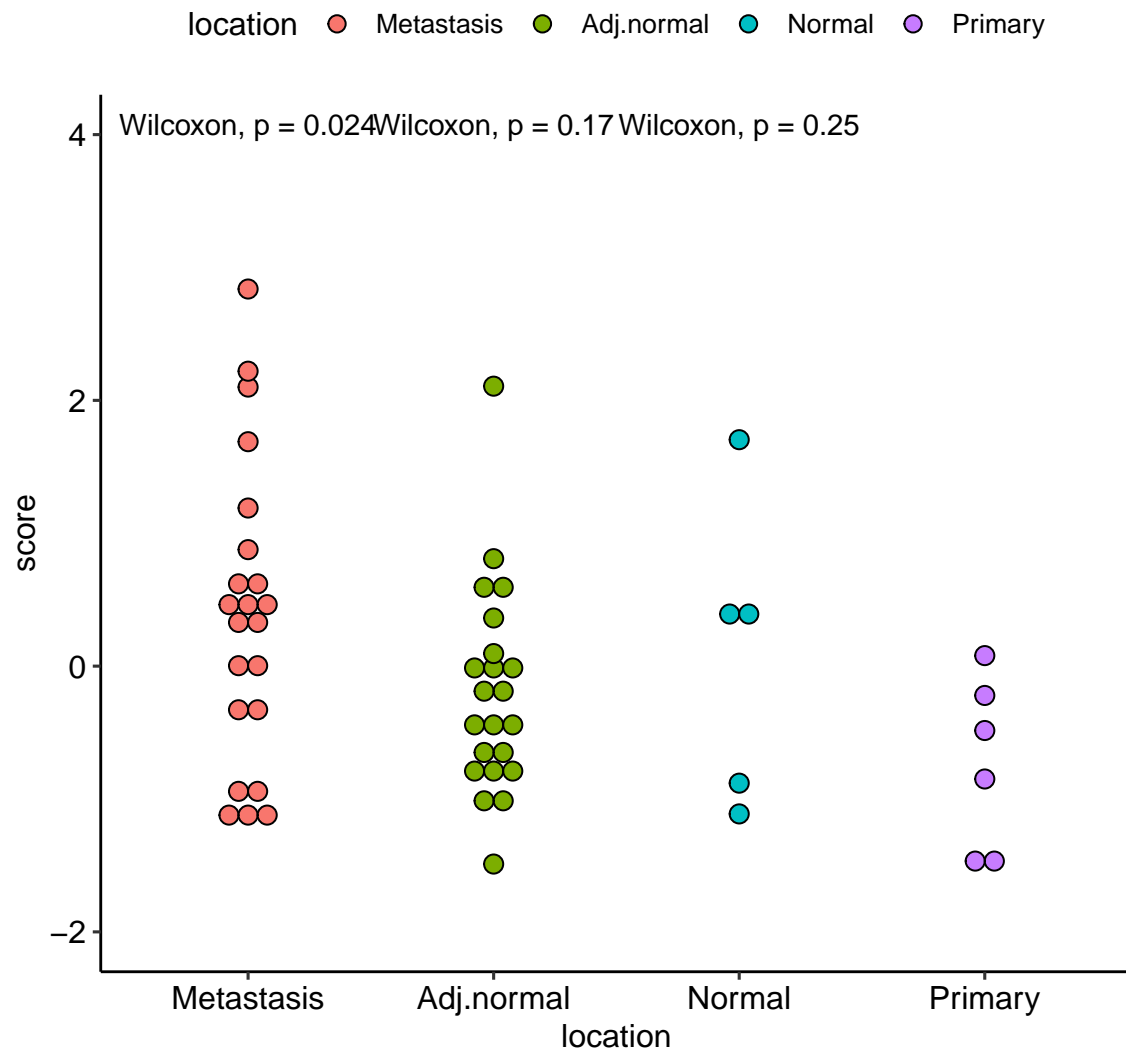


### CIBERSORT.Plasma.cells score

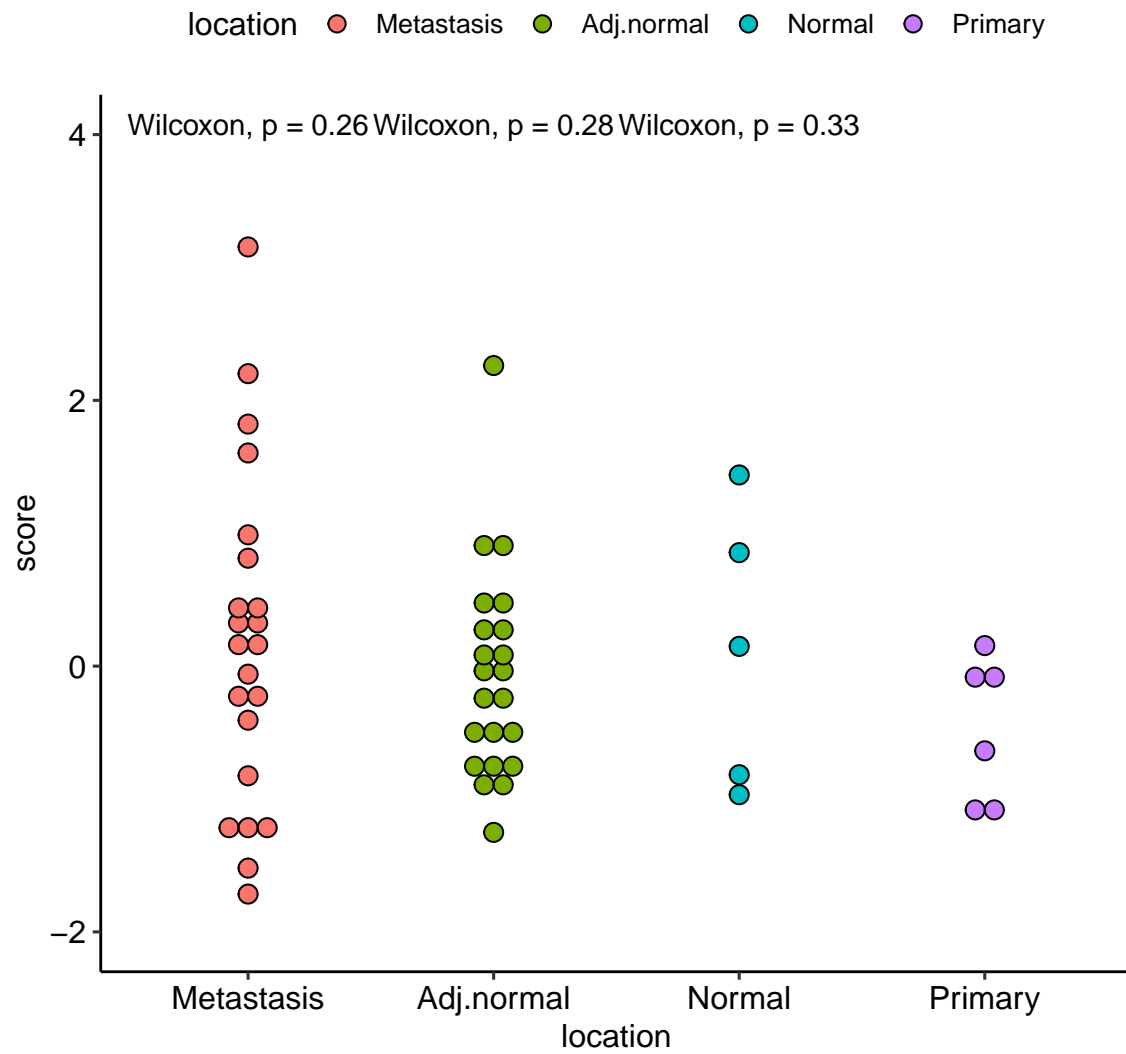
location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



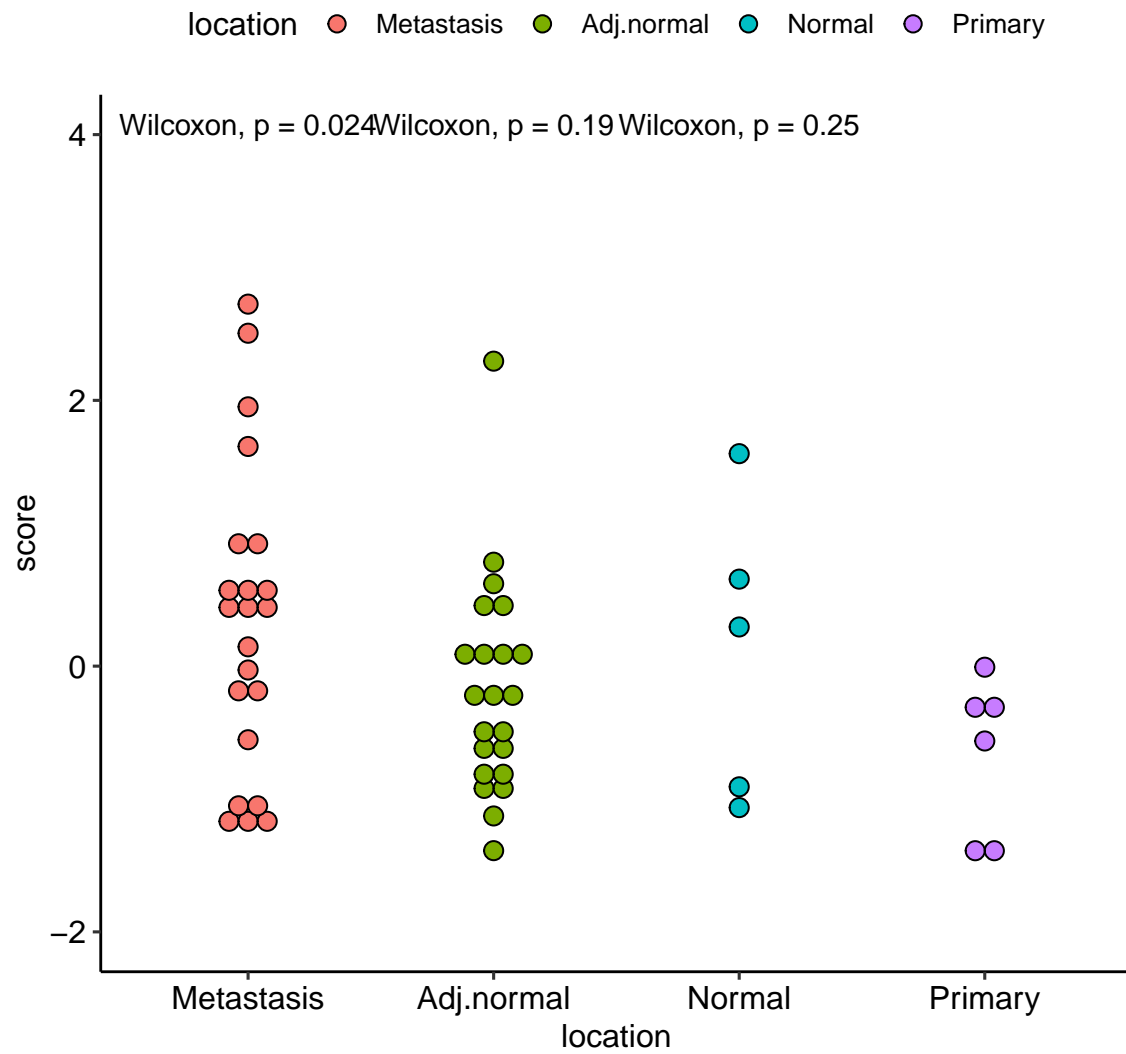
## CIBERSORT.T.cells.CD8 score



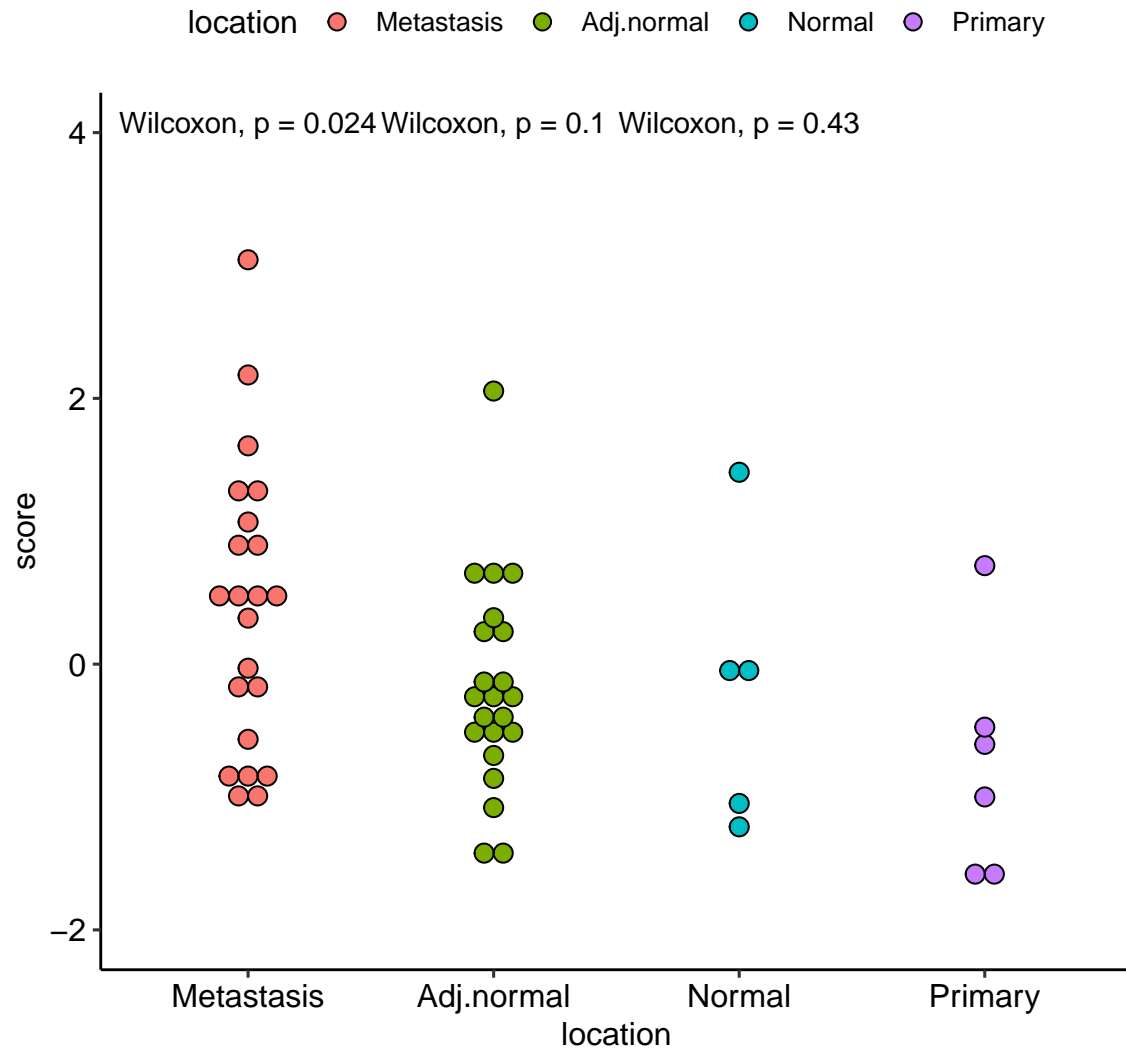
# CIBERSORT.T.cells.CD4.naive score



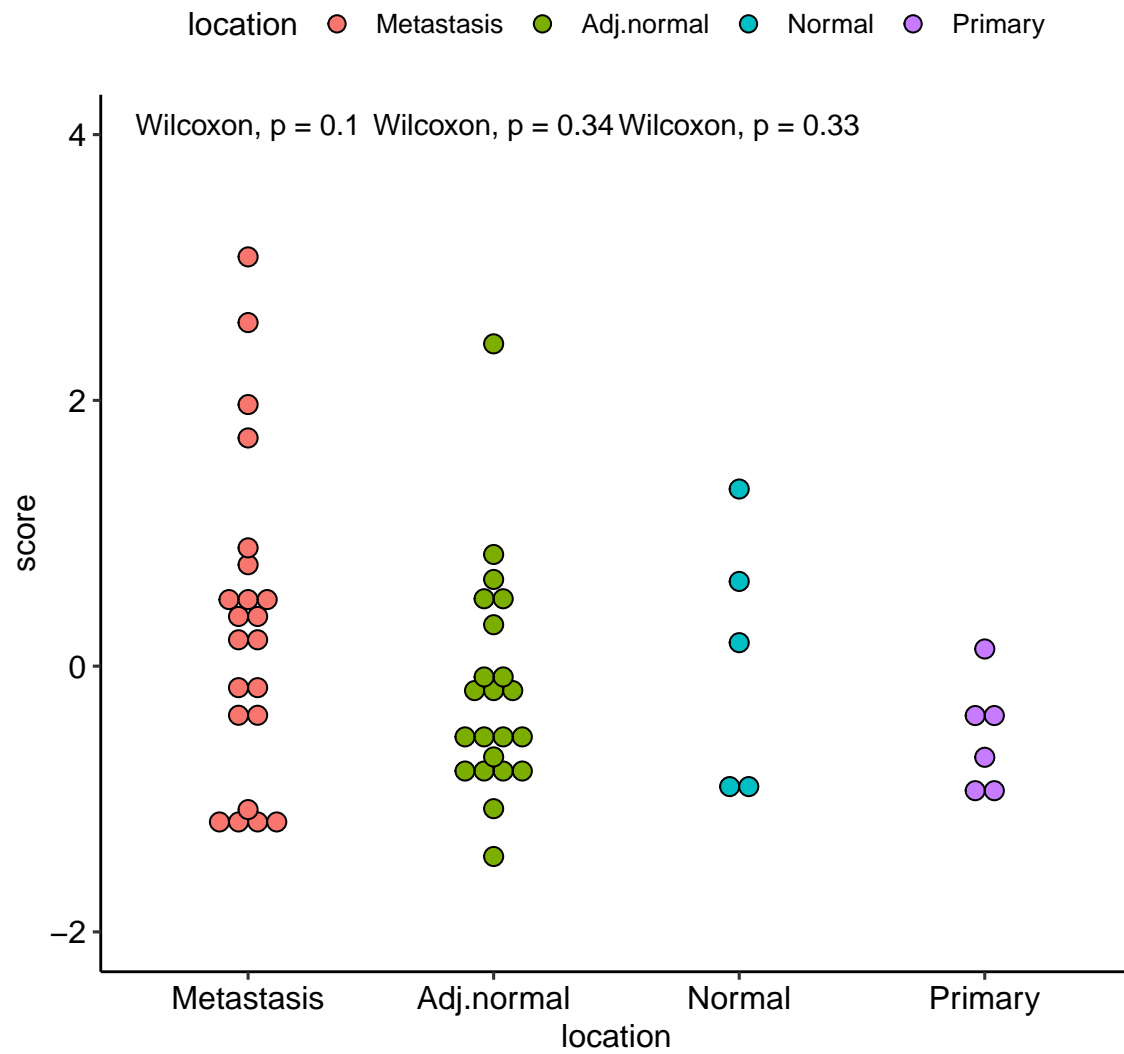
# CIBERSORT.T.cells.CD4.memory.resting score



# CIBERSORT.T.cells.CD4.memory.activated score

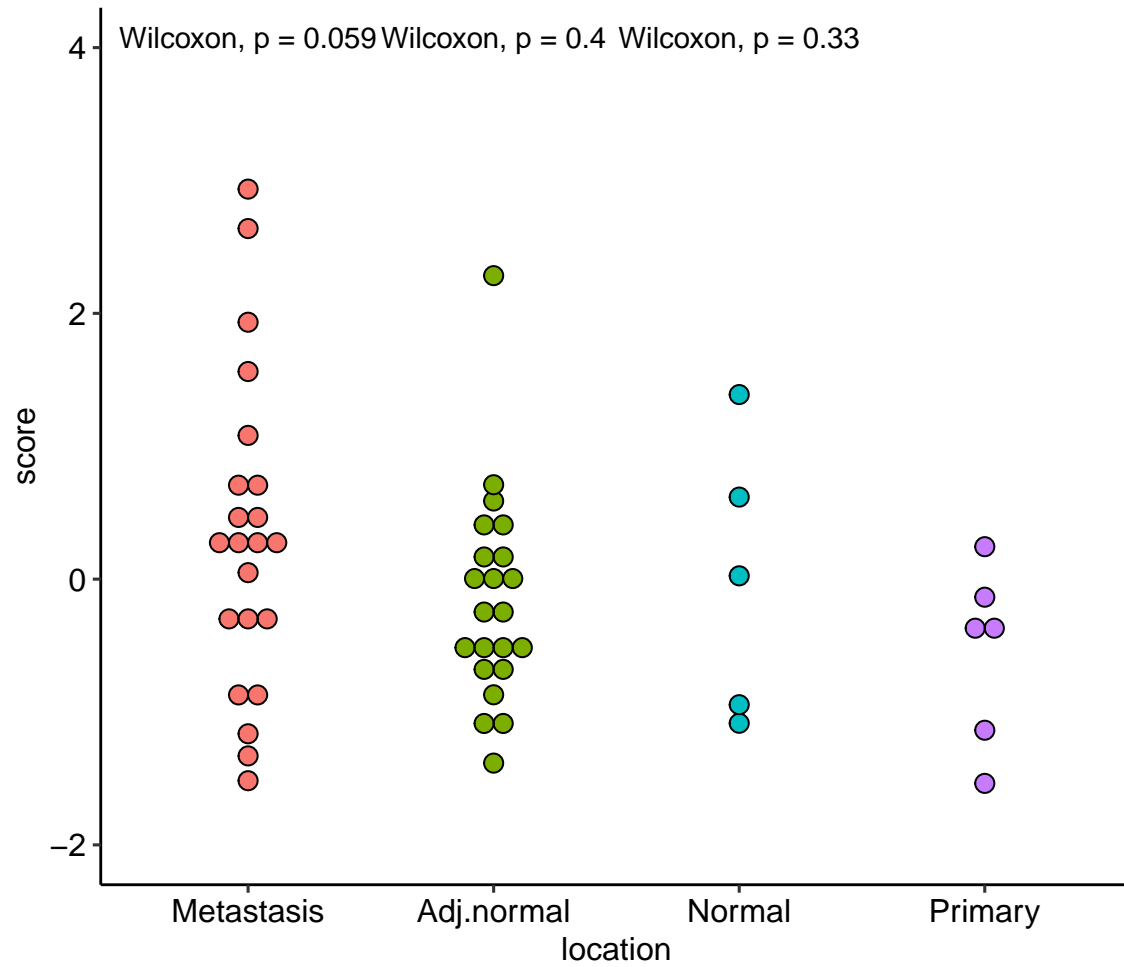


CIBERSORT.T.cells.follicular.helper score



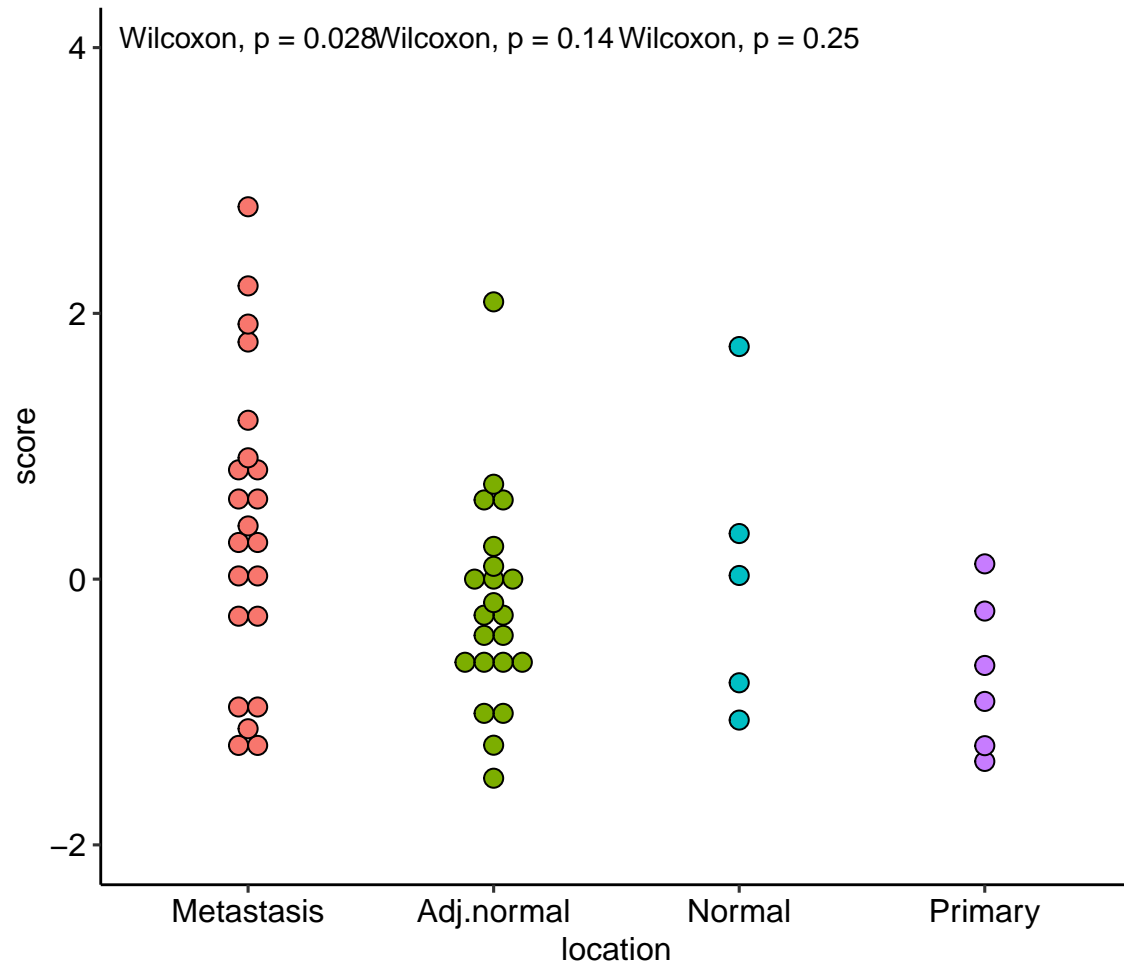
CIBERSORT.T.cells.regulatory..Tregs. score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



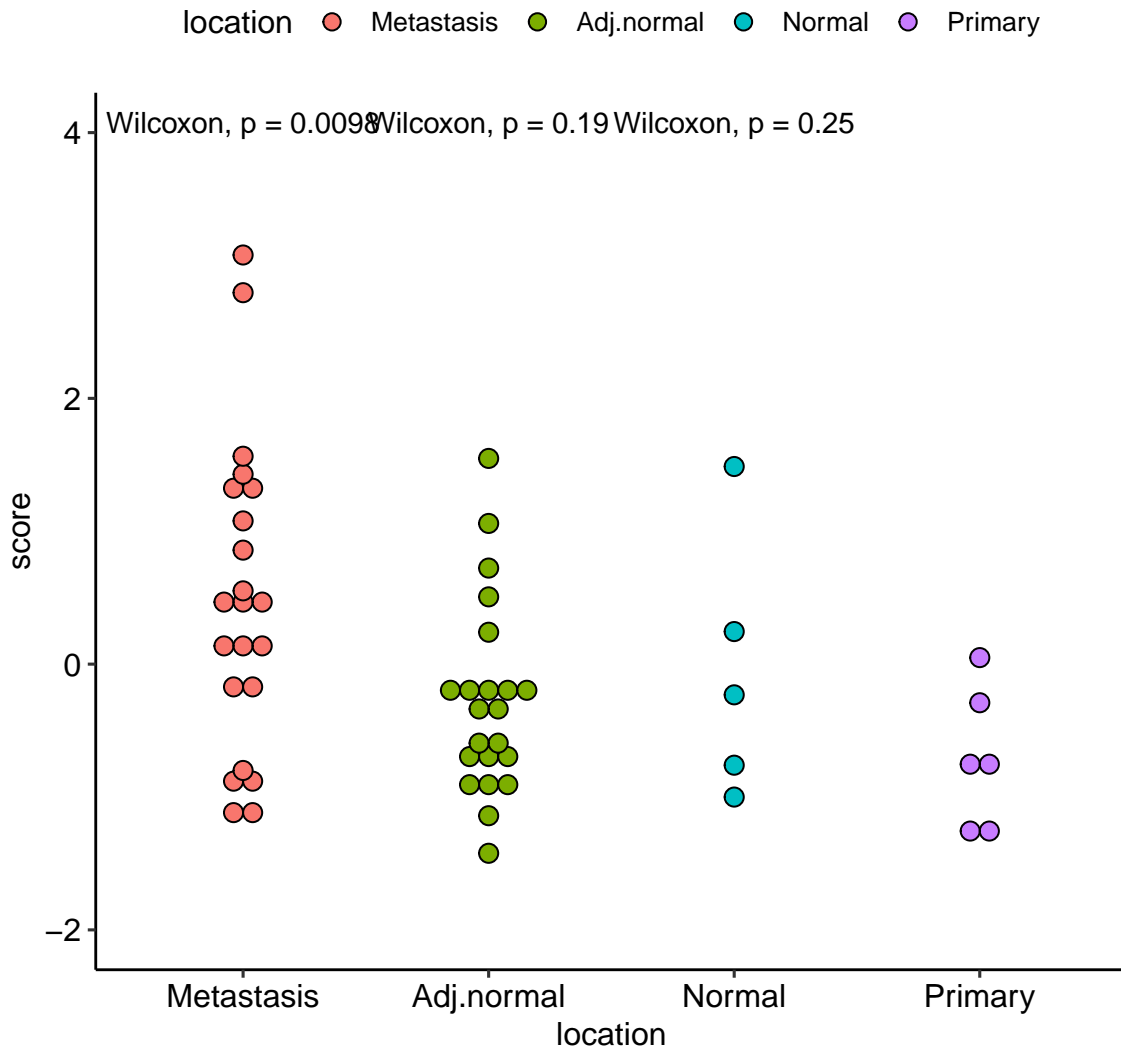
CIBERSORT.T.cells.gamma.delta score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

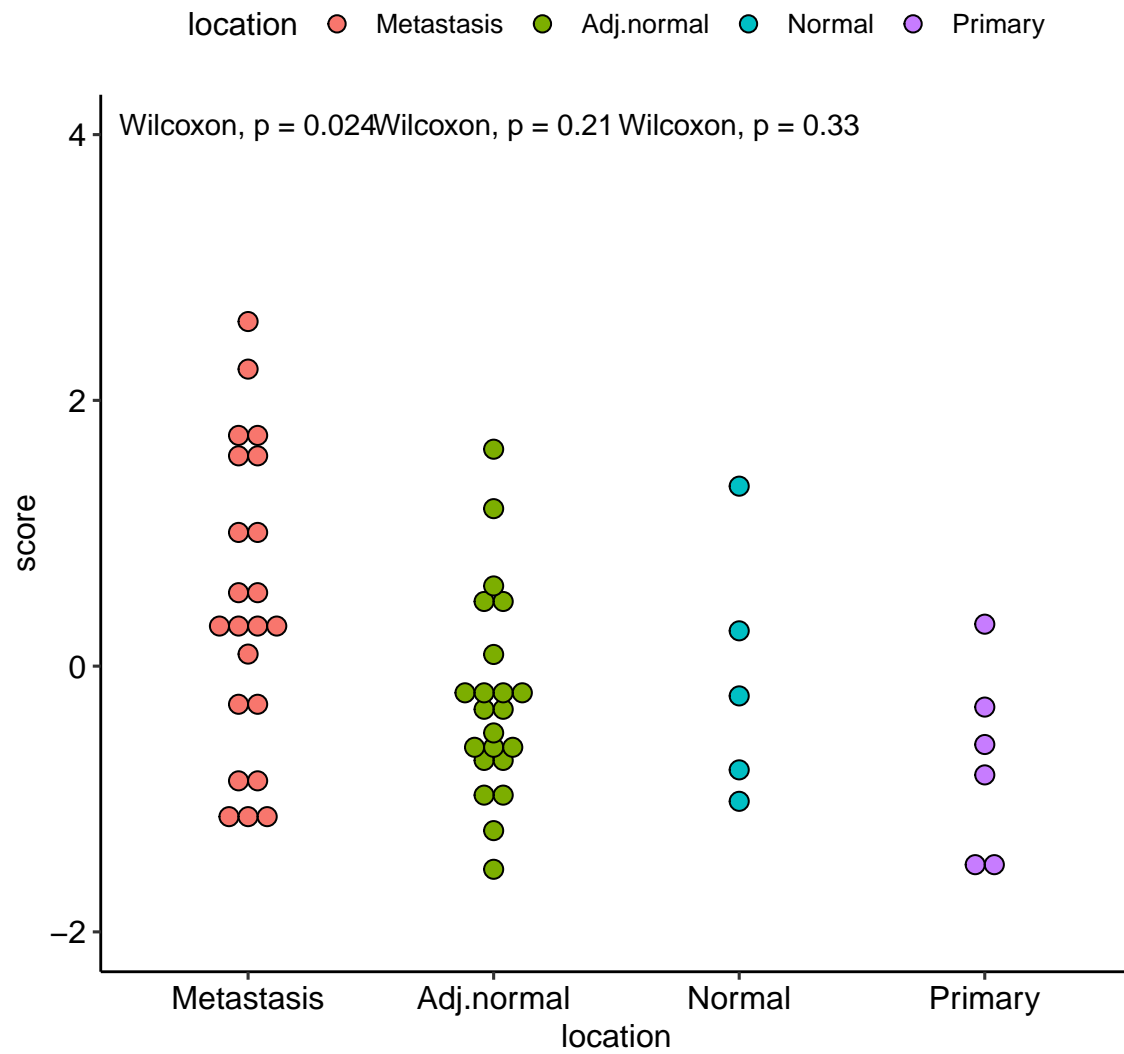




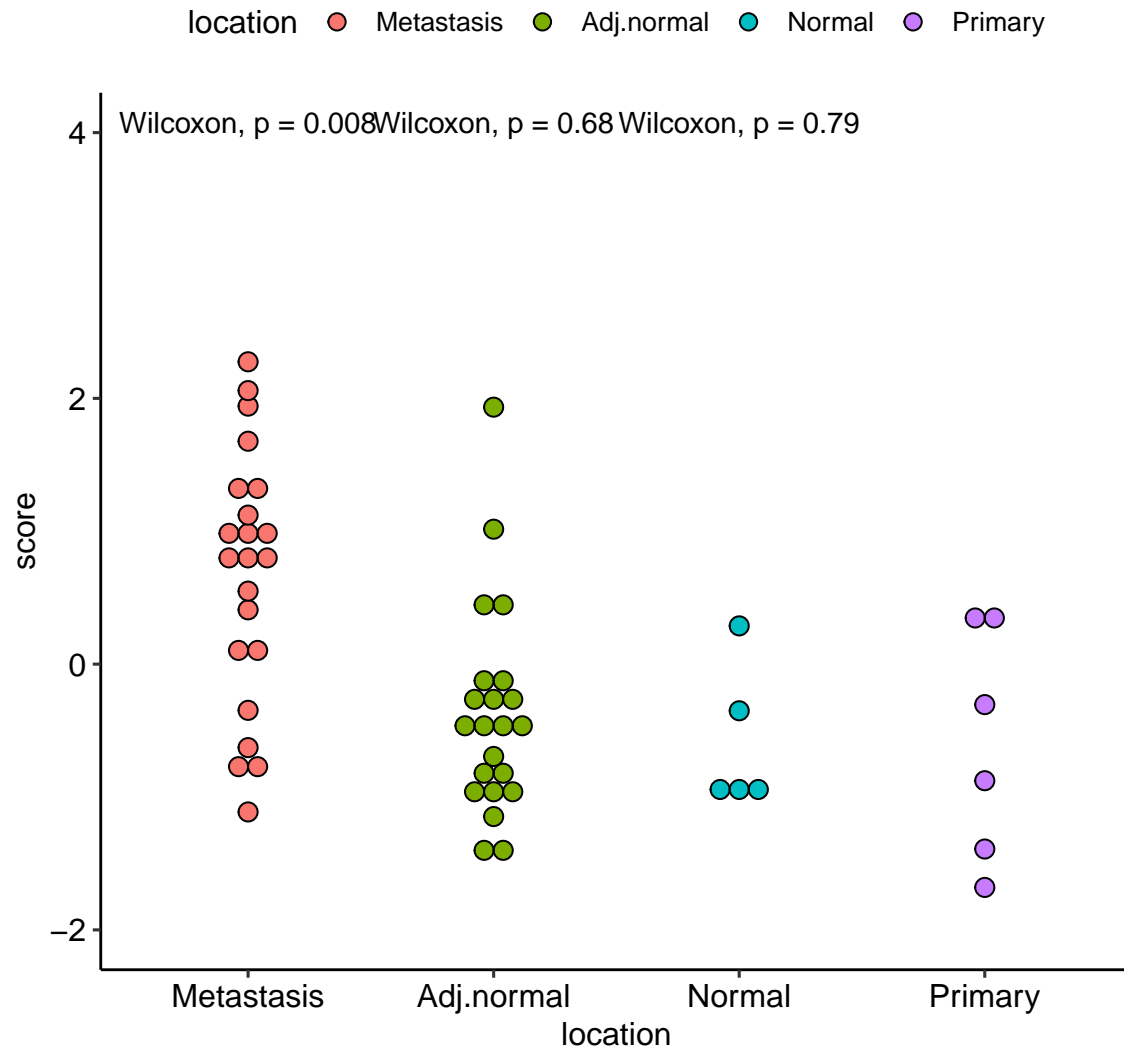
## CIBERSORT.NK.cells.resting score



## CIBERSORT.NK.cells.activated score



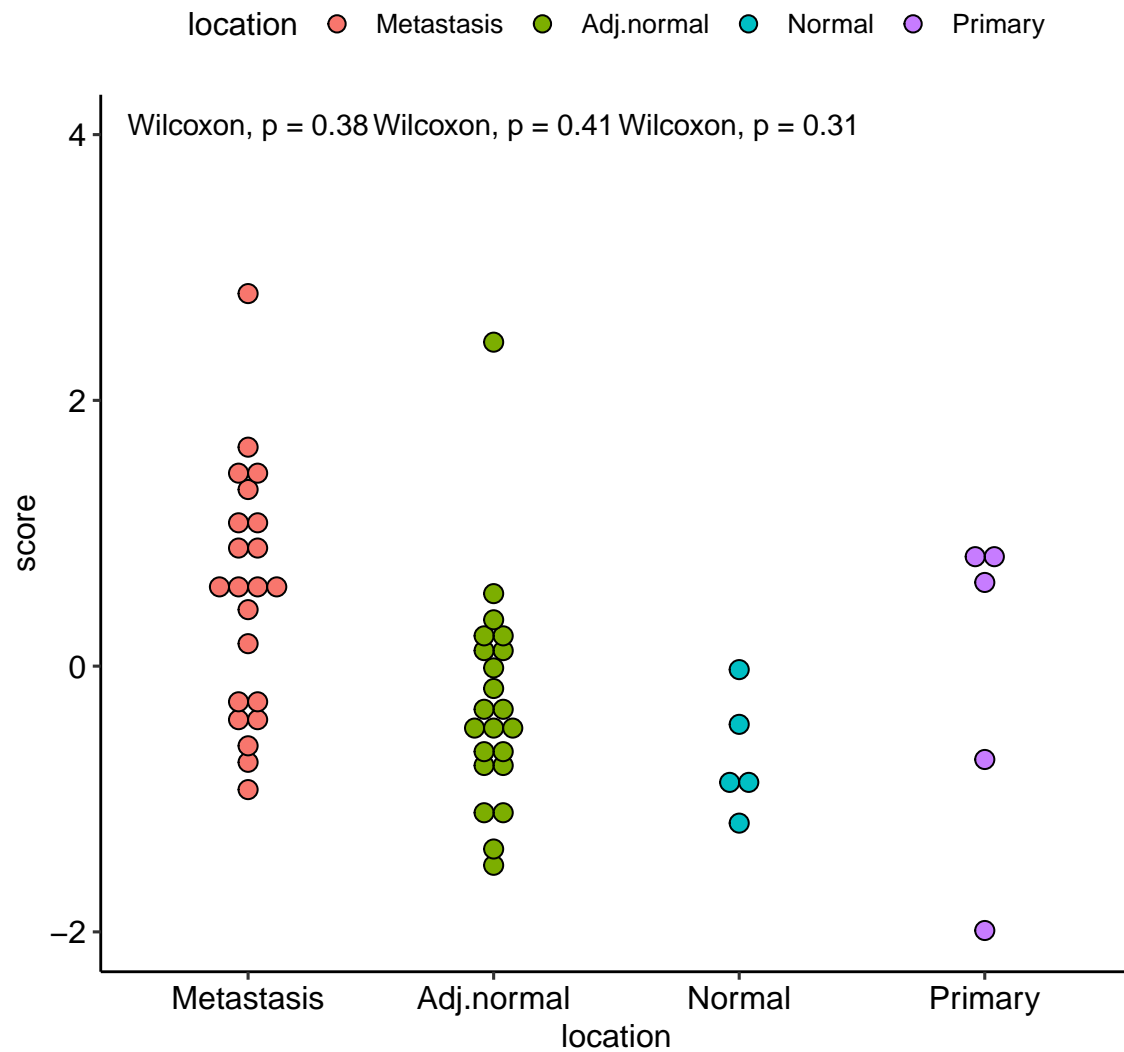
## CIBERSORT.Monocytes score



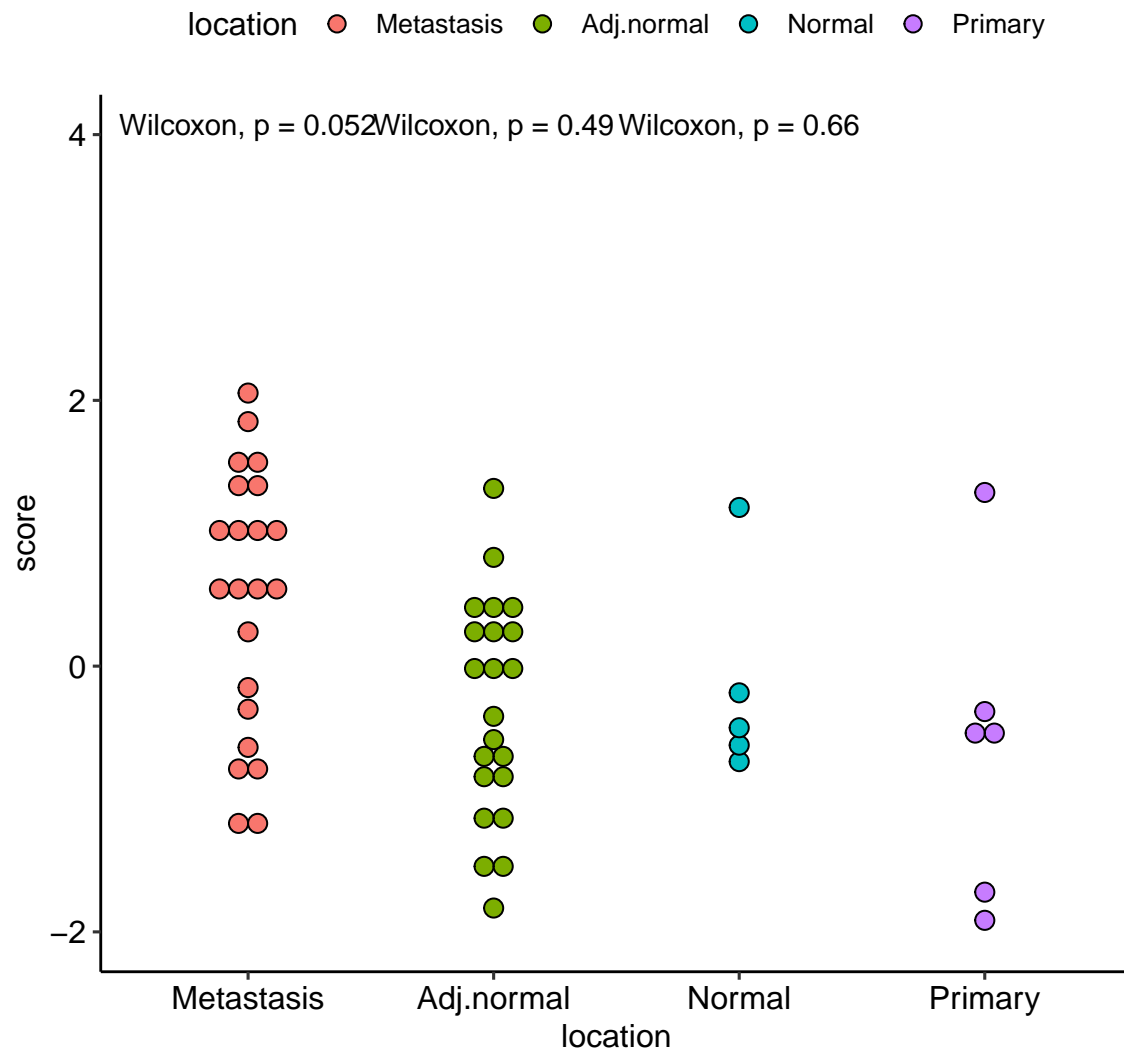
## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

## CIBERSORT.Macrophages.M0 score



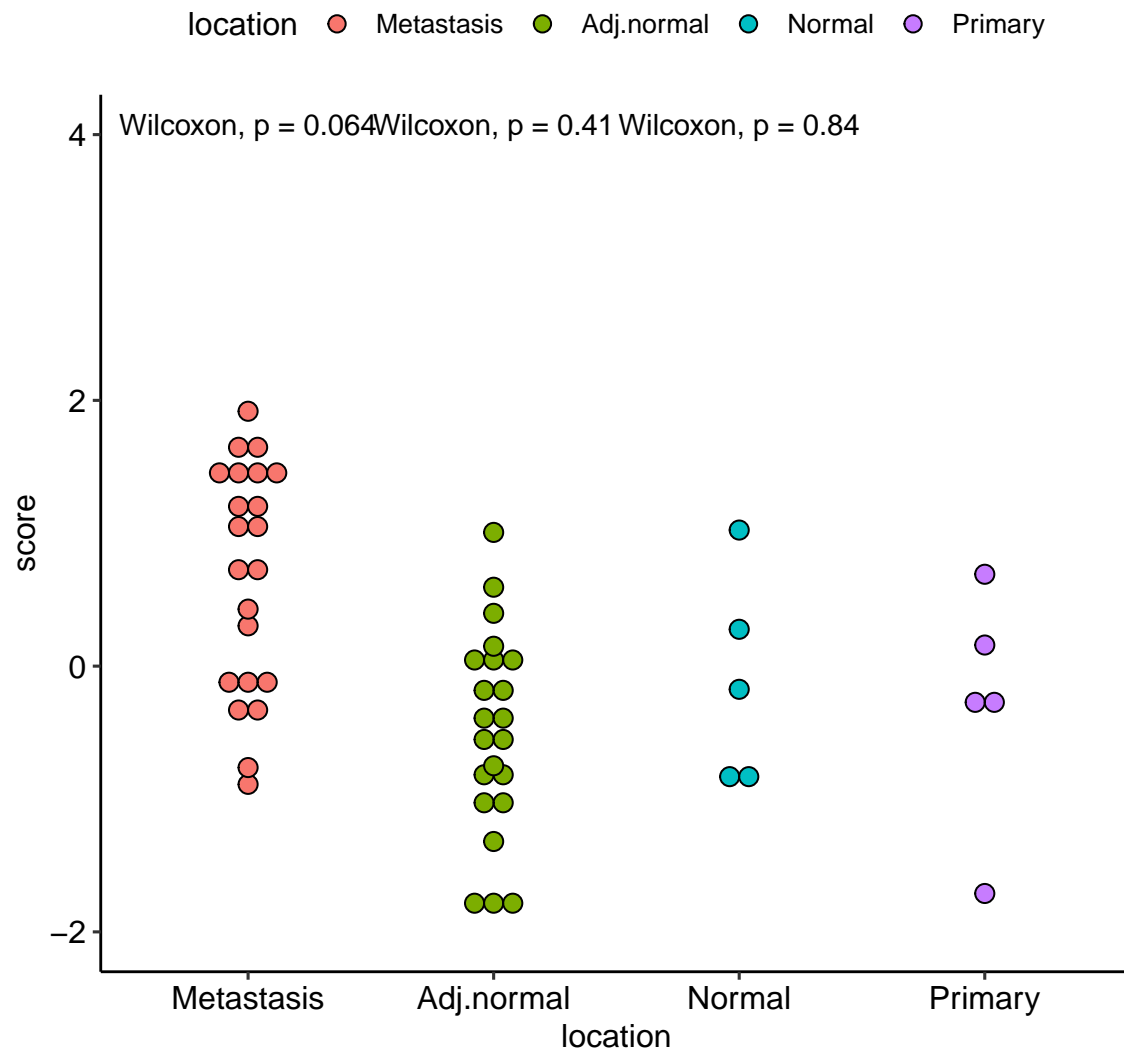
CIBERSORT.Macrophages.M1 score



```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

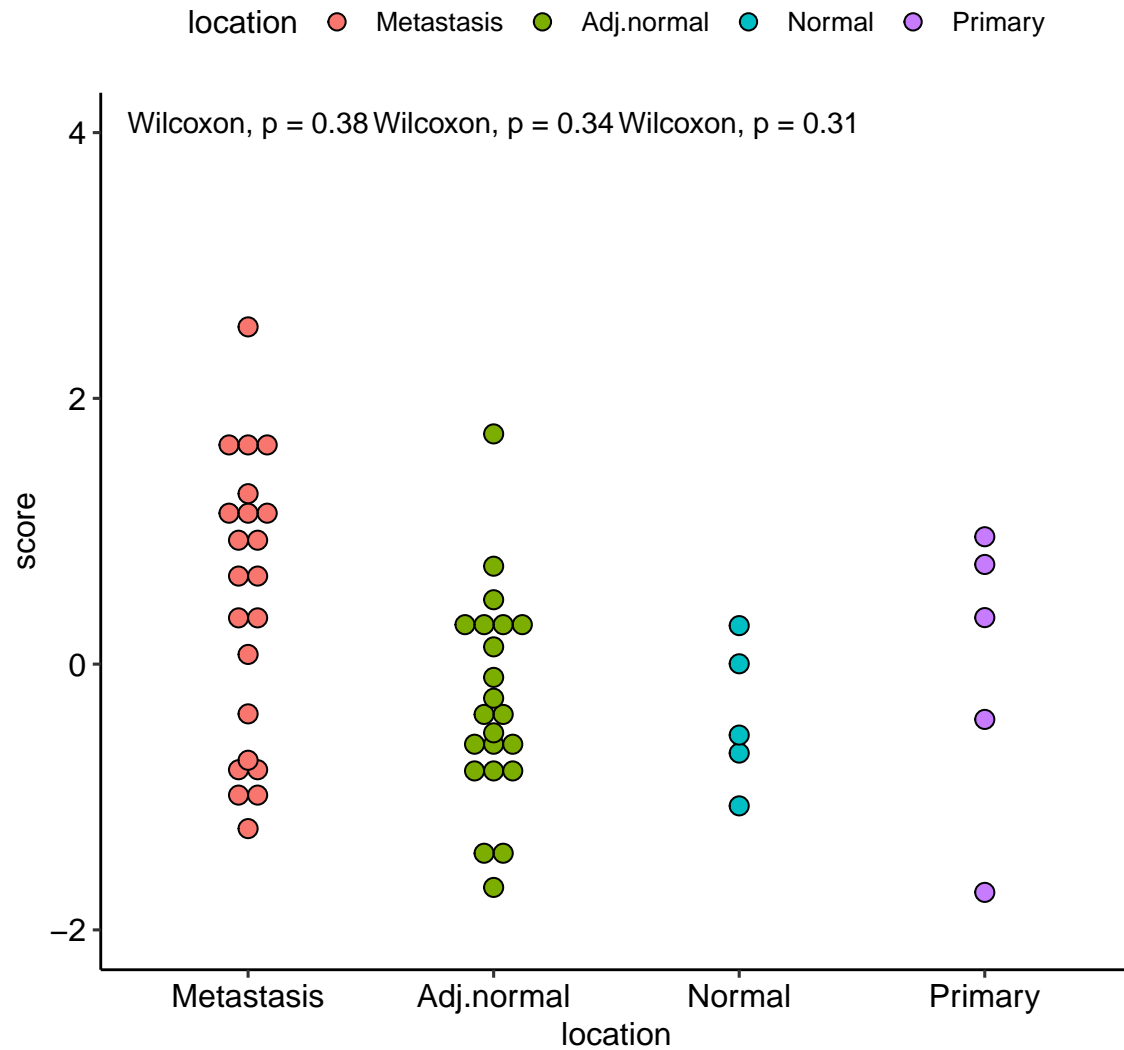
CIBERSORT.Macrophages.M2 score



```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```

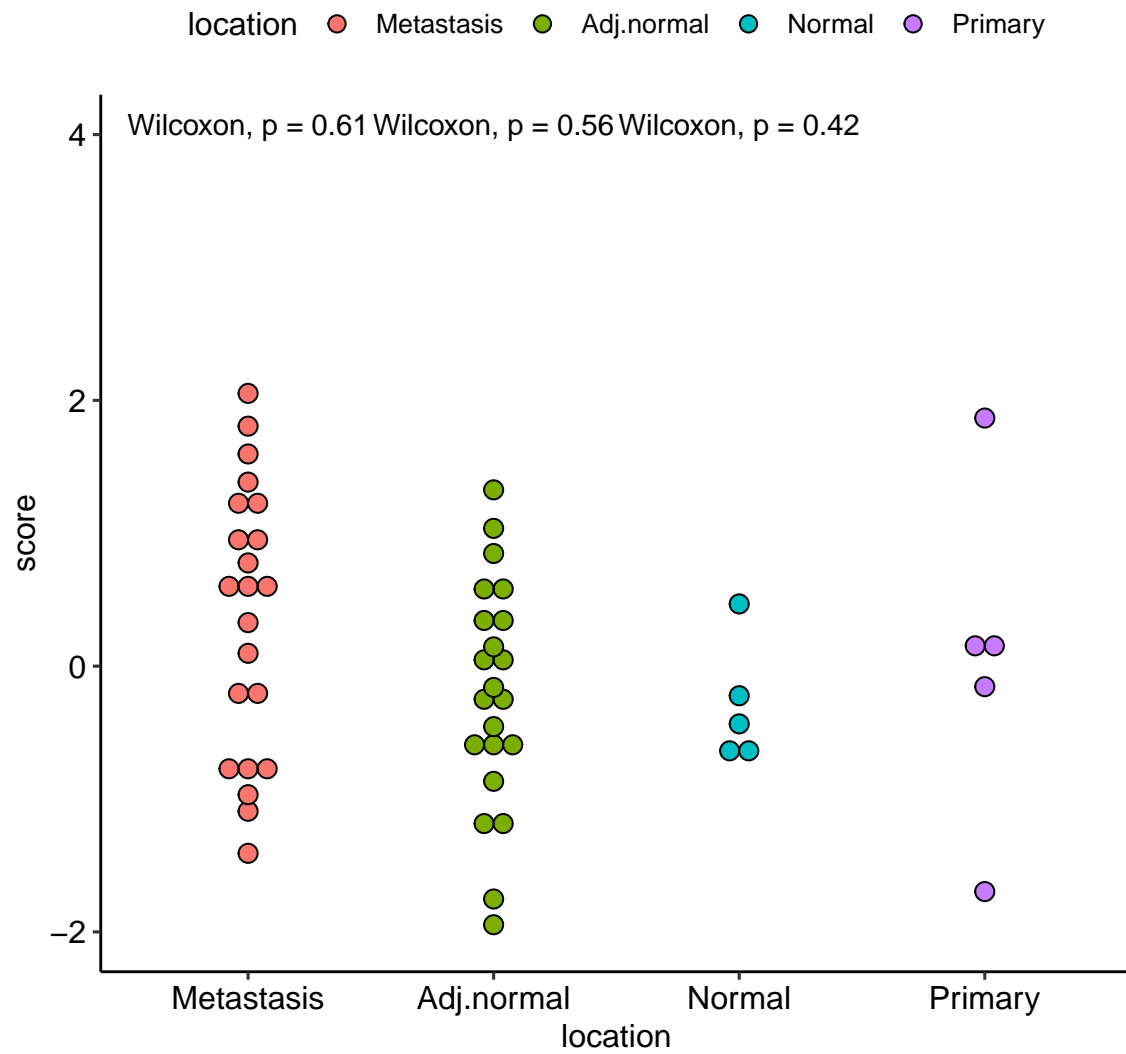
## CIBERSORT.Dendritic.cells.resting score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

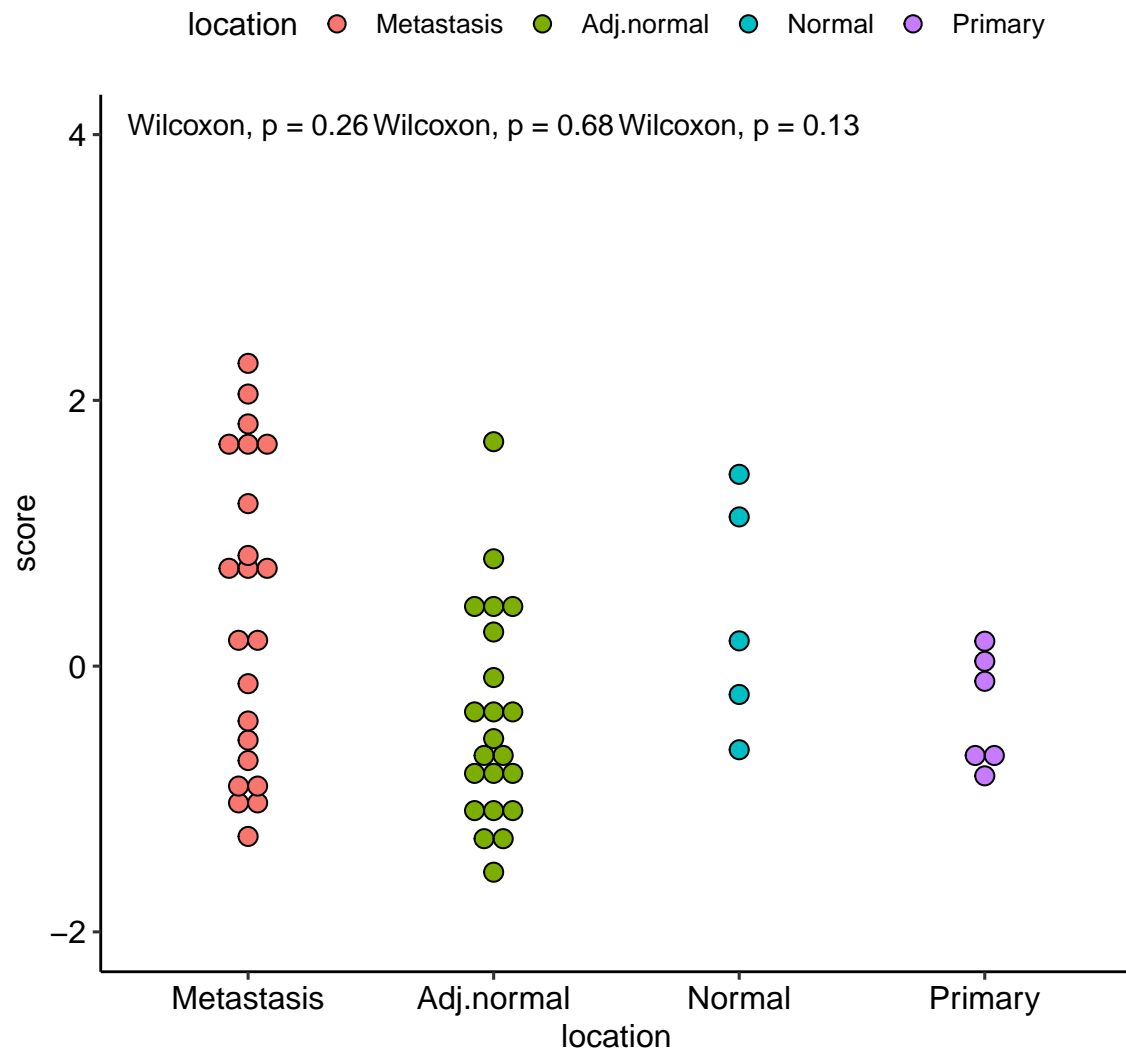
## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

# CIBERSORT.Dendritic.cells.activated score



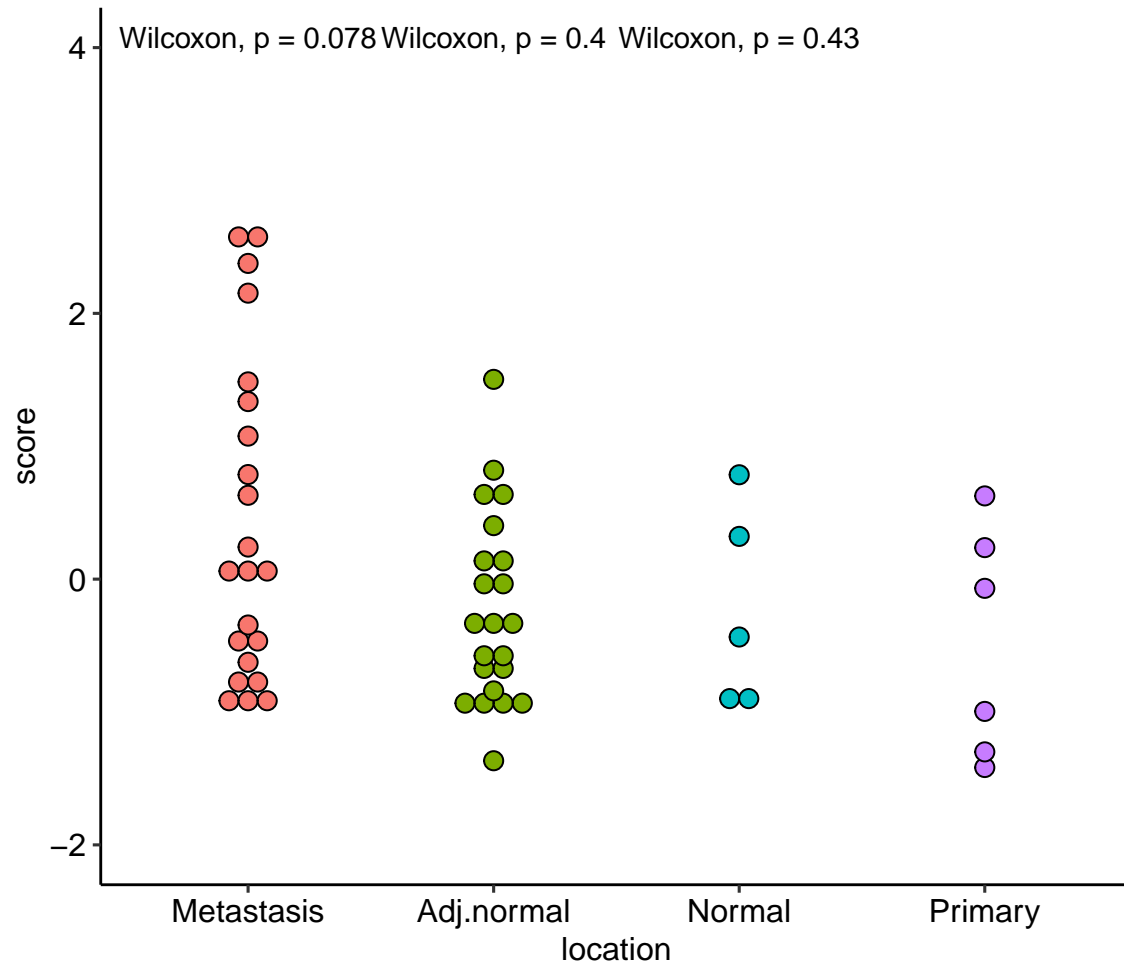


# CIBERSORT.Mast.cells.resting score

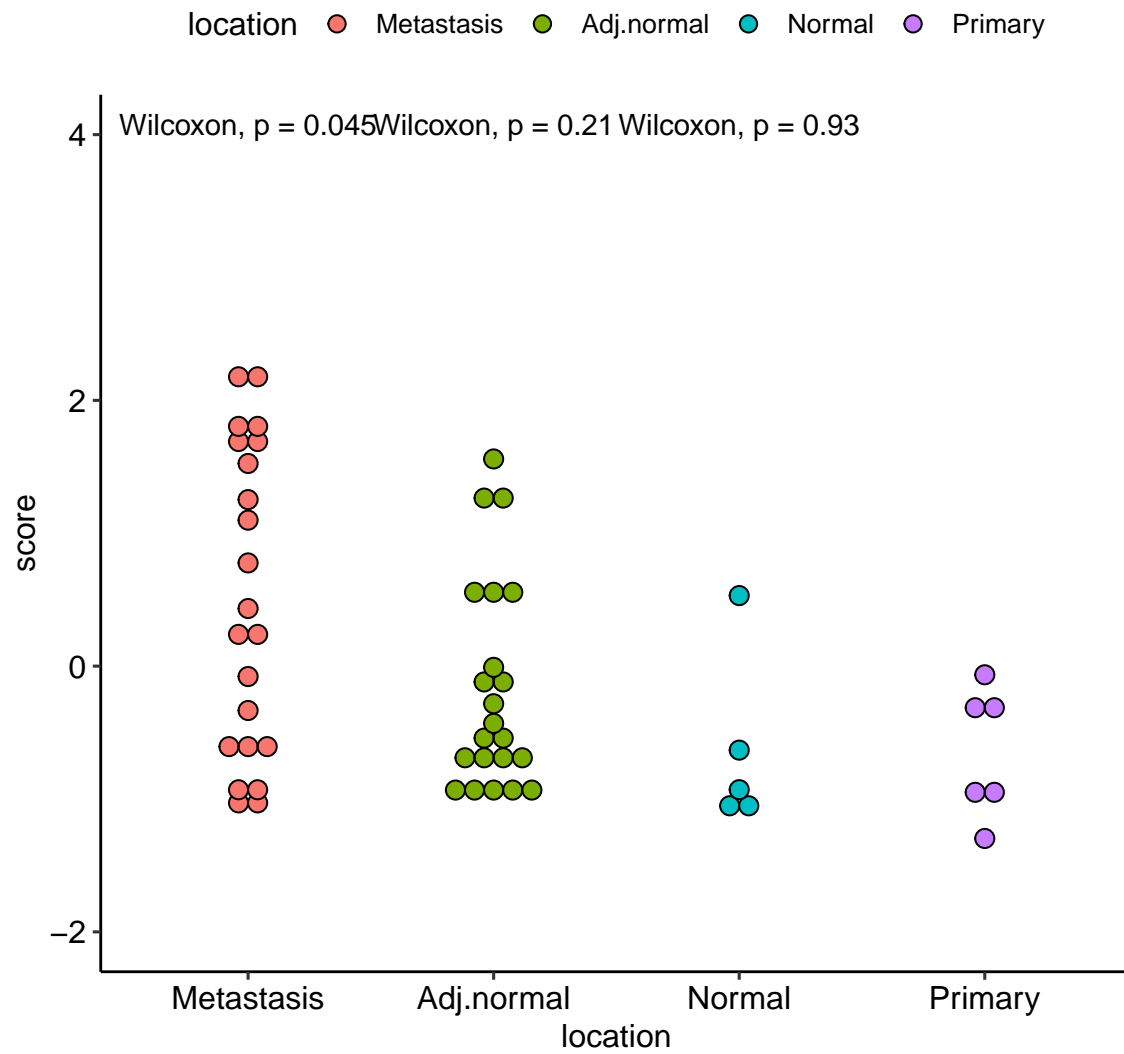


CIBERSORT.Mast.cells.activated score

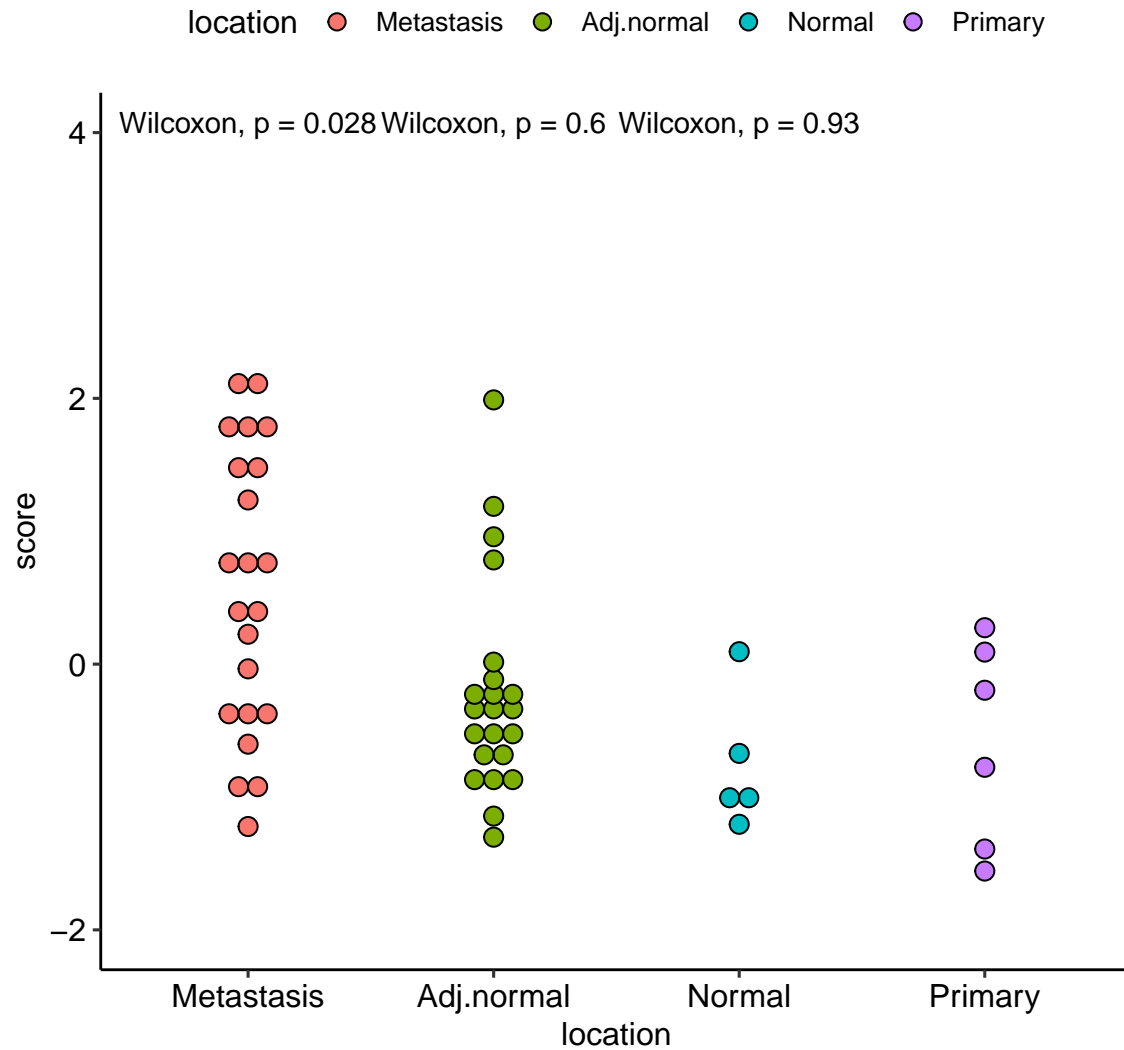
location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



CIBERSORT.Eosinophils score



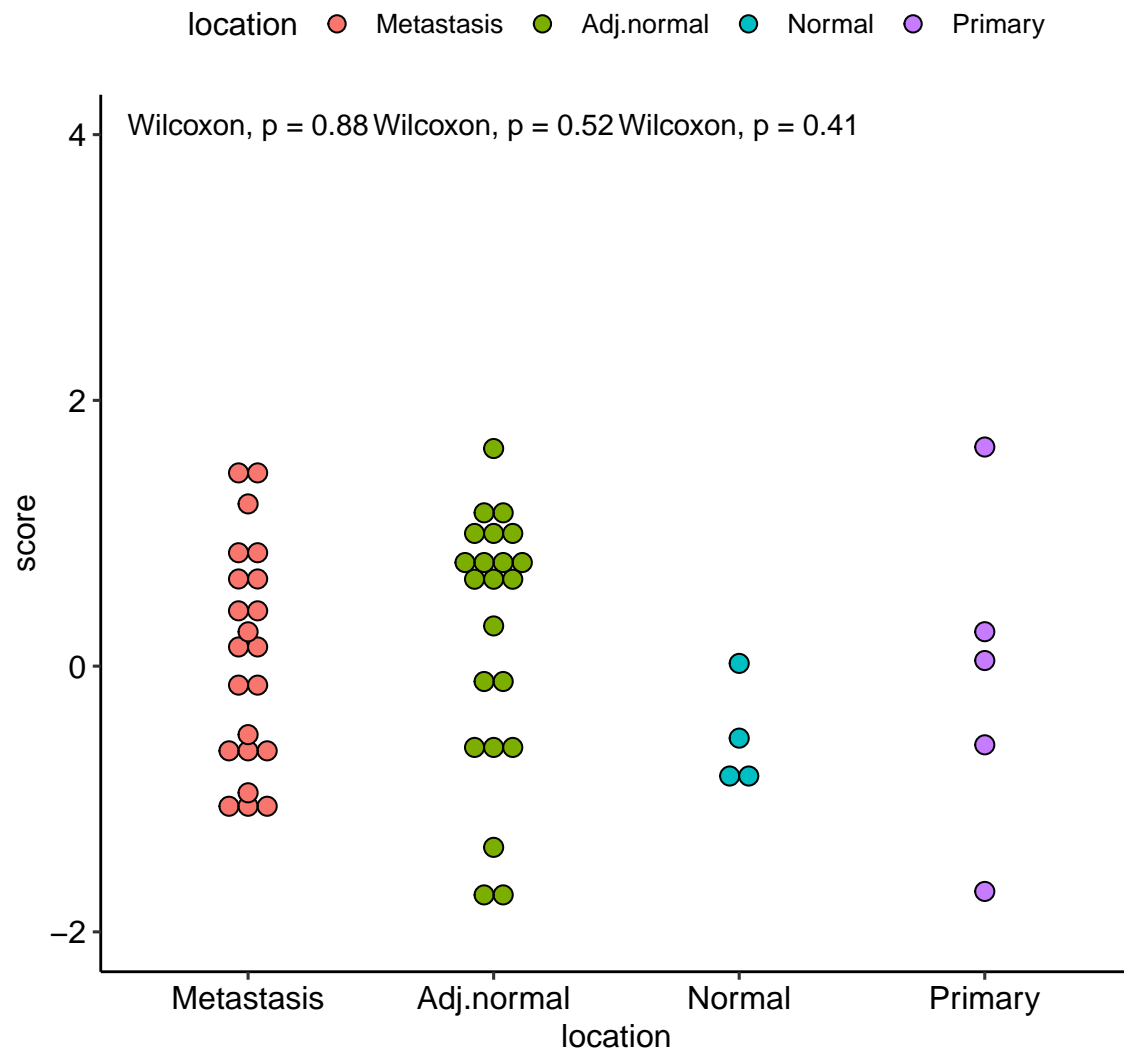
## CIBERSORT.Neutrophils score



## Warning: Removed 2 rows containing non-finite values (stat\_bindot).

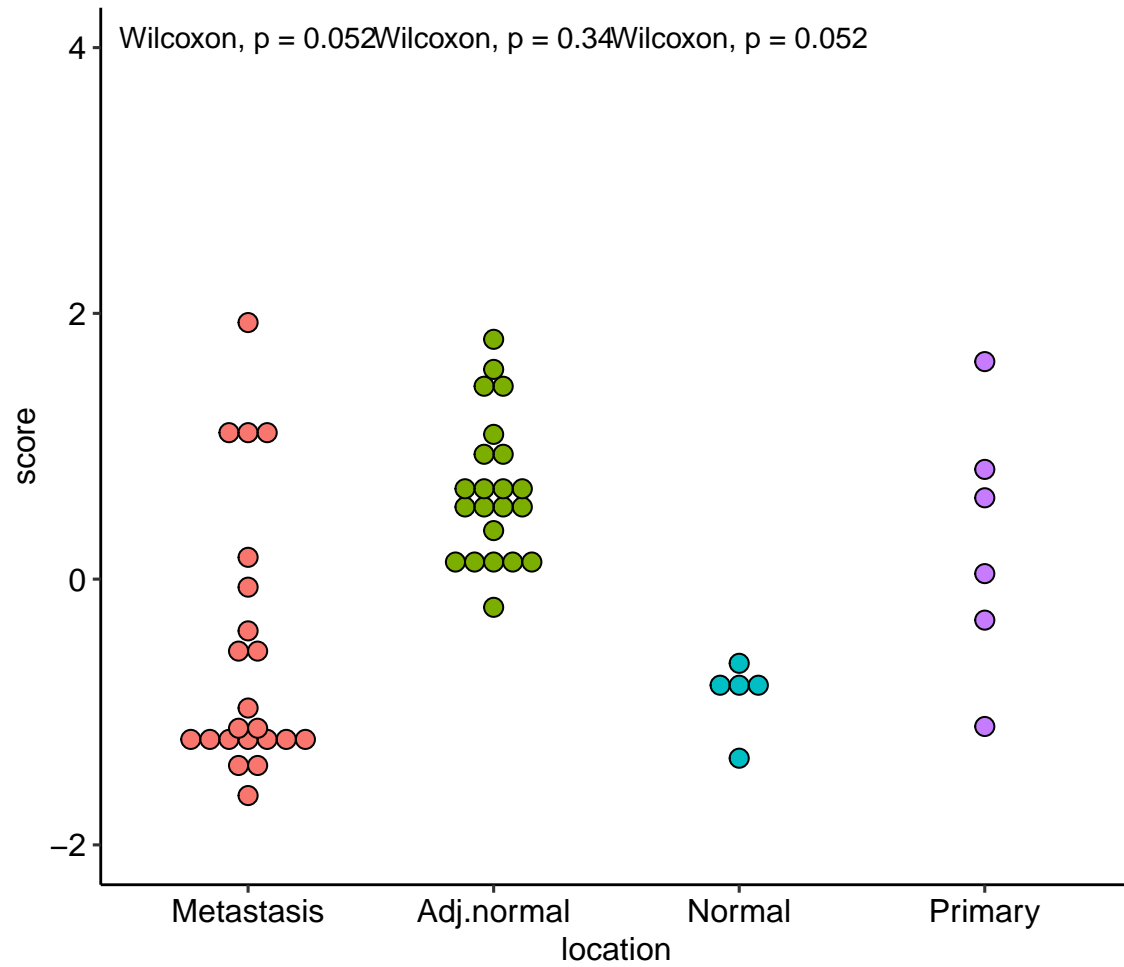
## Warning: Removed 2 rows containing non-finite values (stat\_compare\_means).

## Notta.BasalA score



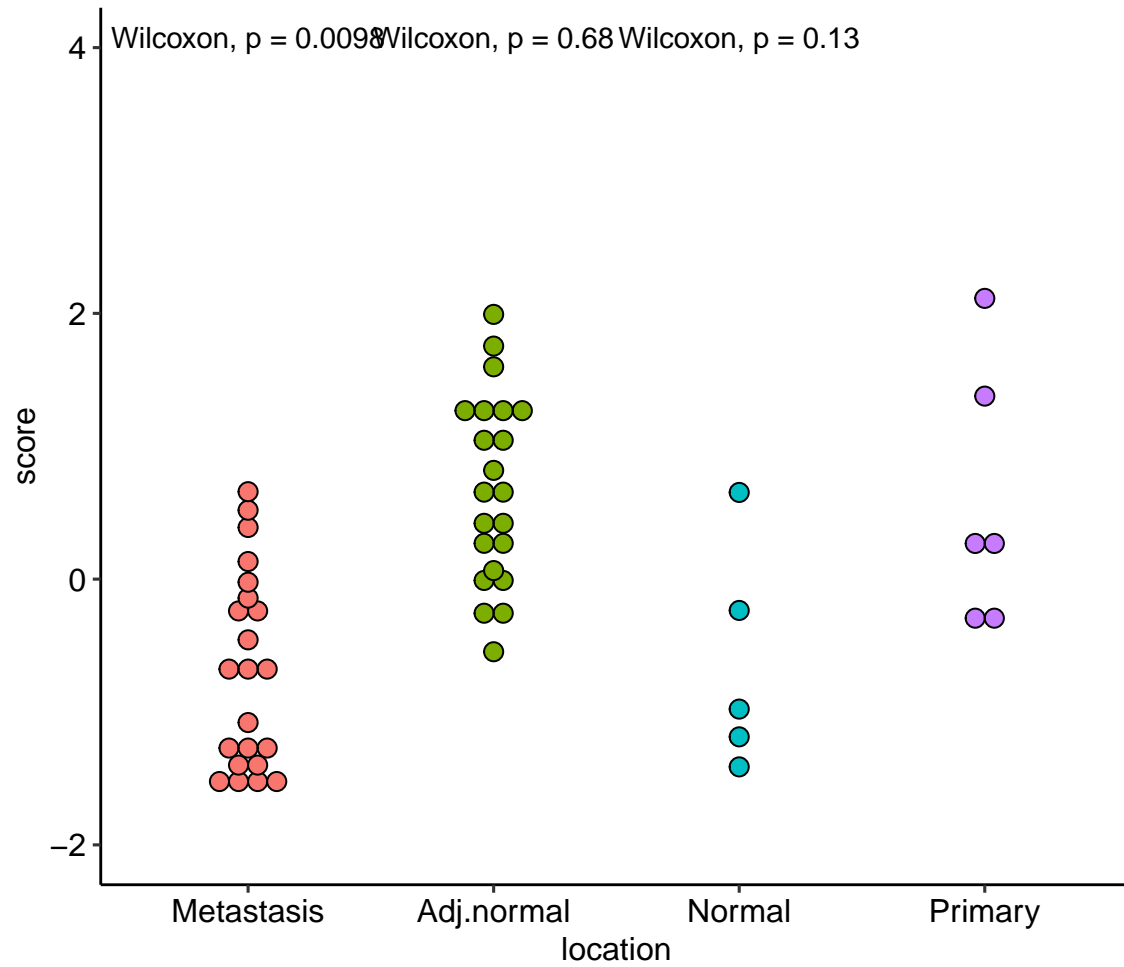
## Notta.BasalB score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

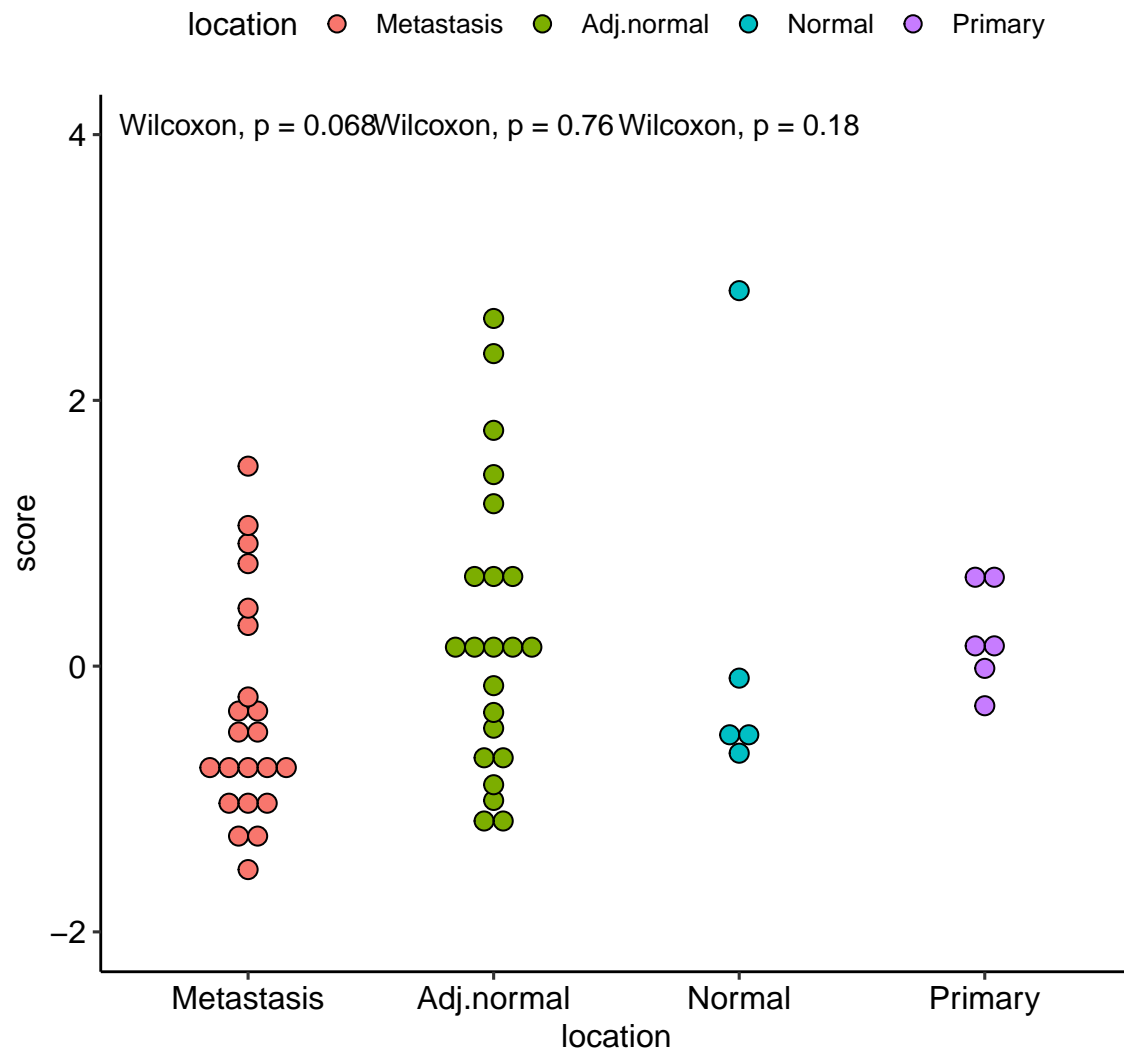


## Notta.ClassicalA score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



Notta.ClassicalB score

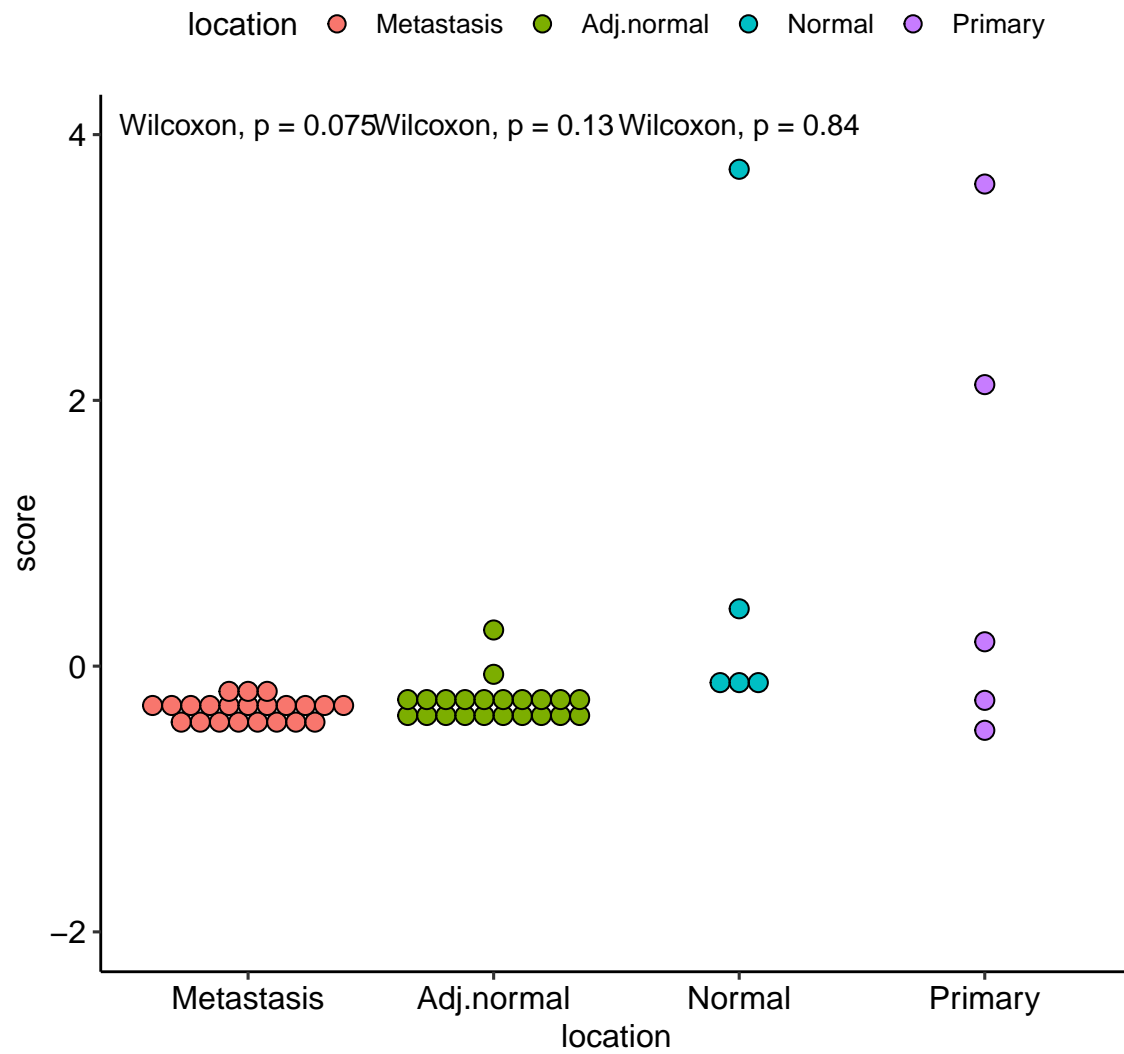


```
## Warning: Removed 1 rows containing non-finite values (stat_bindot).
```

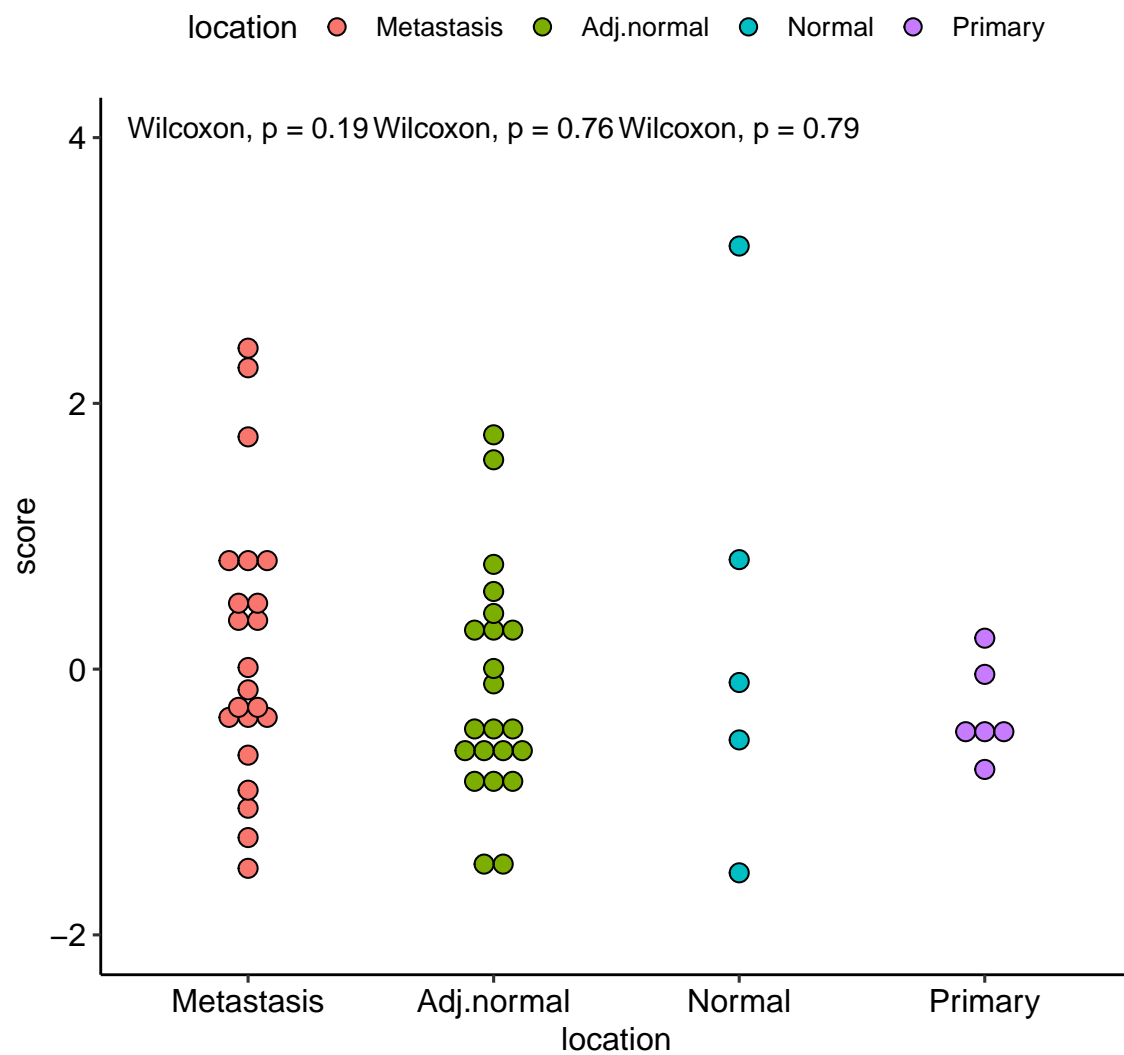
```
## Warning: Removed 1 rows containing non-finite values (stat_compare_means).
```



ADEX\_unique score

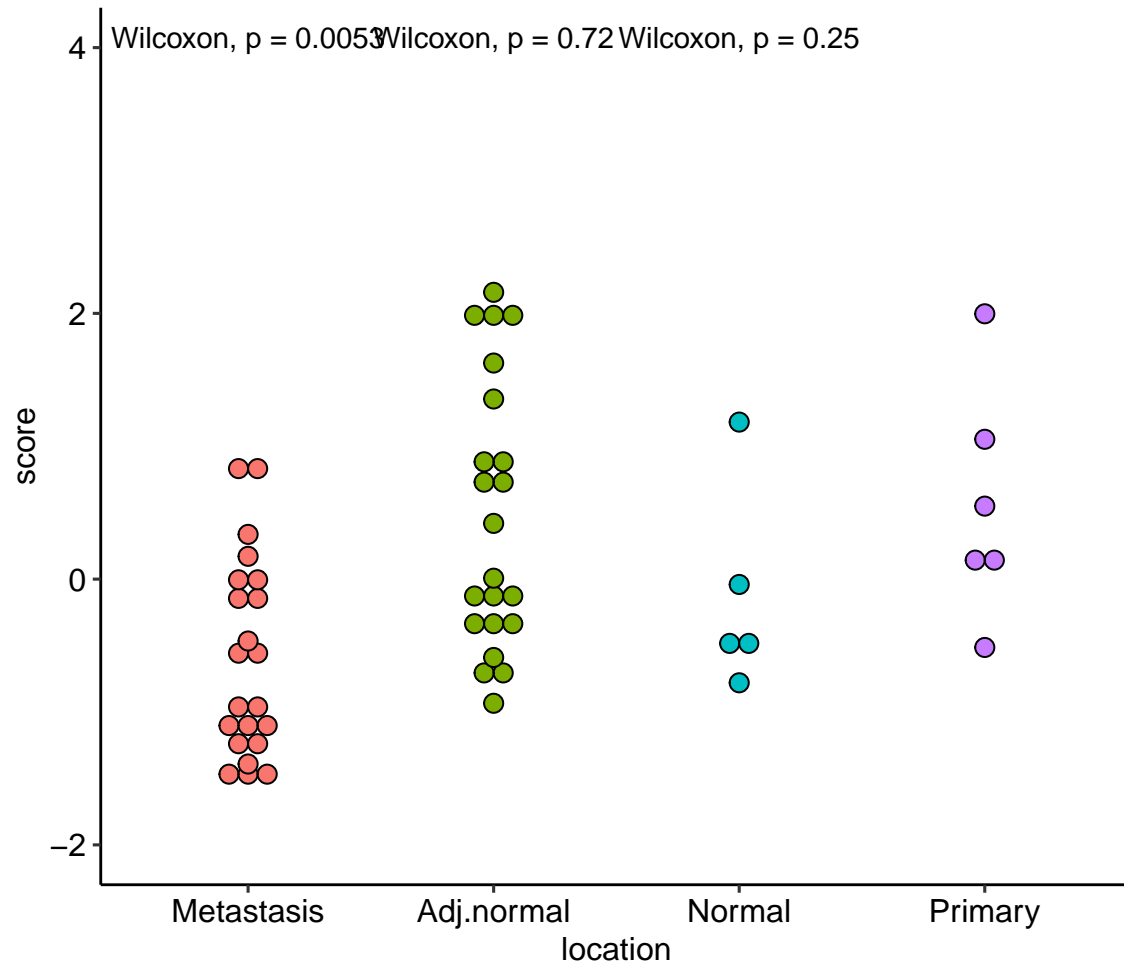


# Immunogenic\_unique score



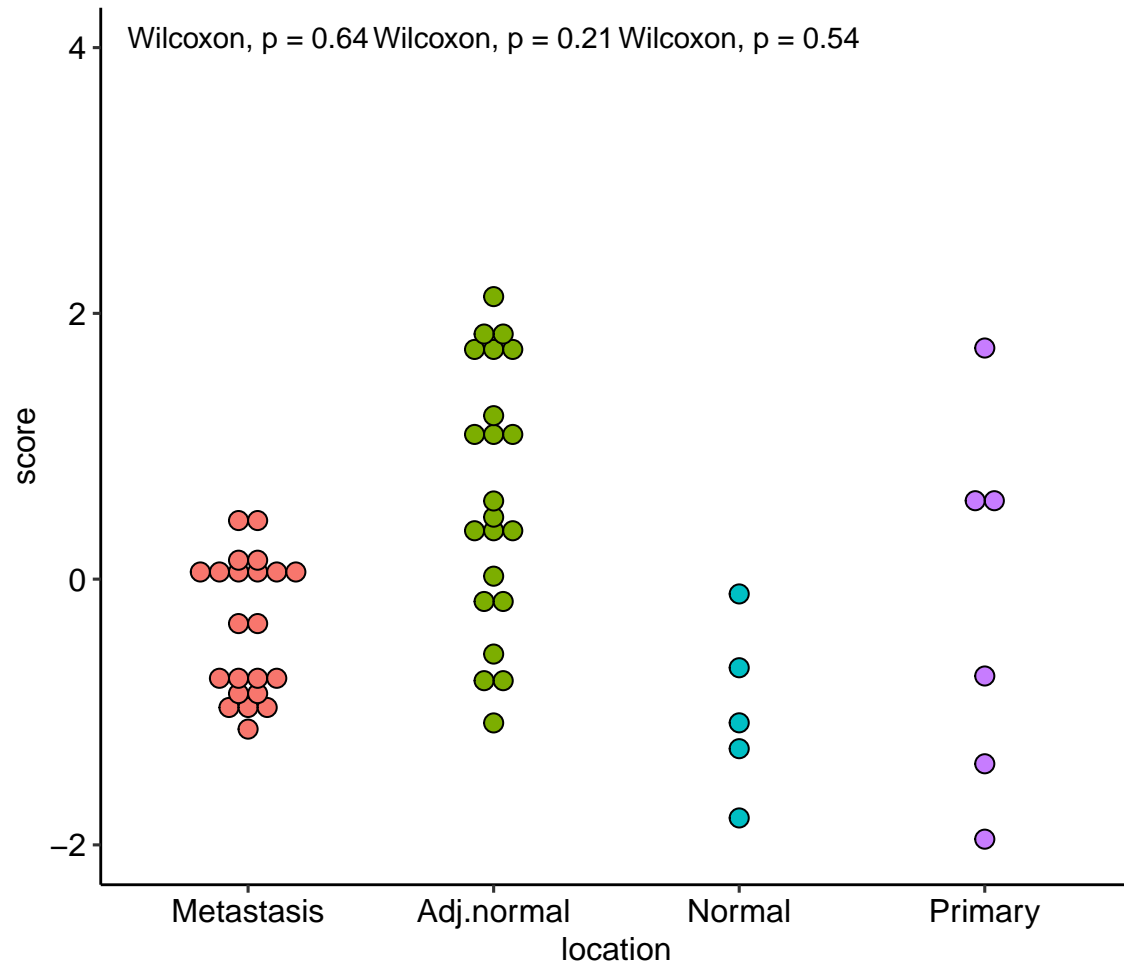
## Progenitor\_unique score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



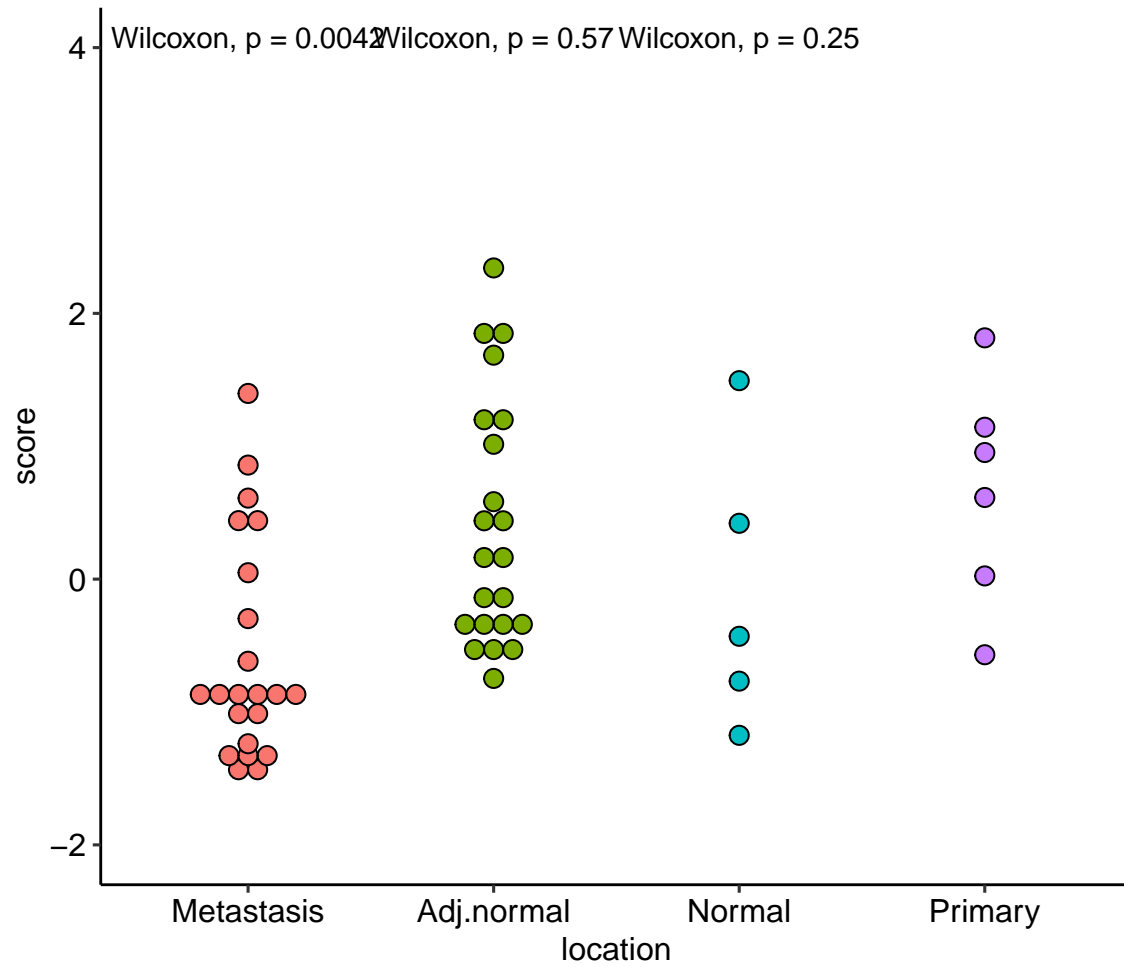
Squamous\_unique score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

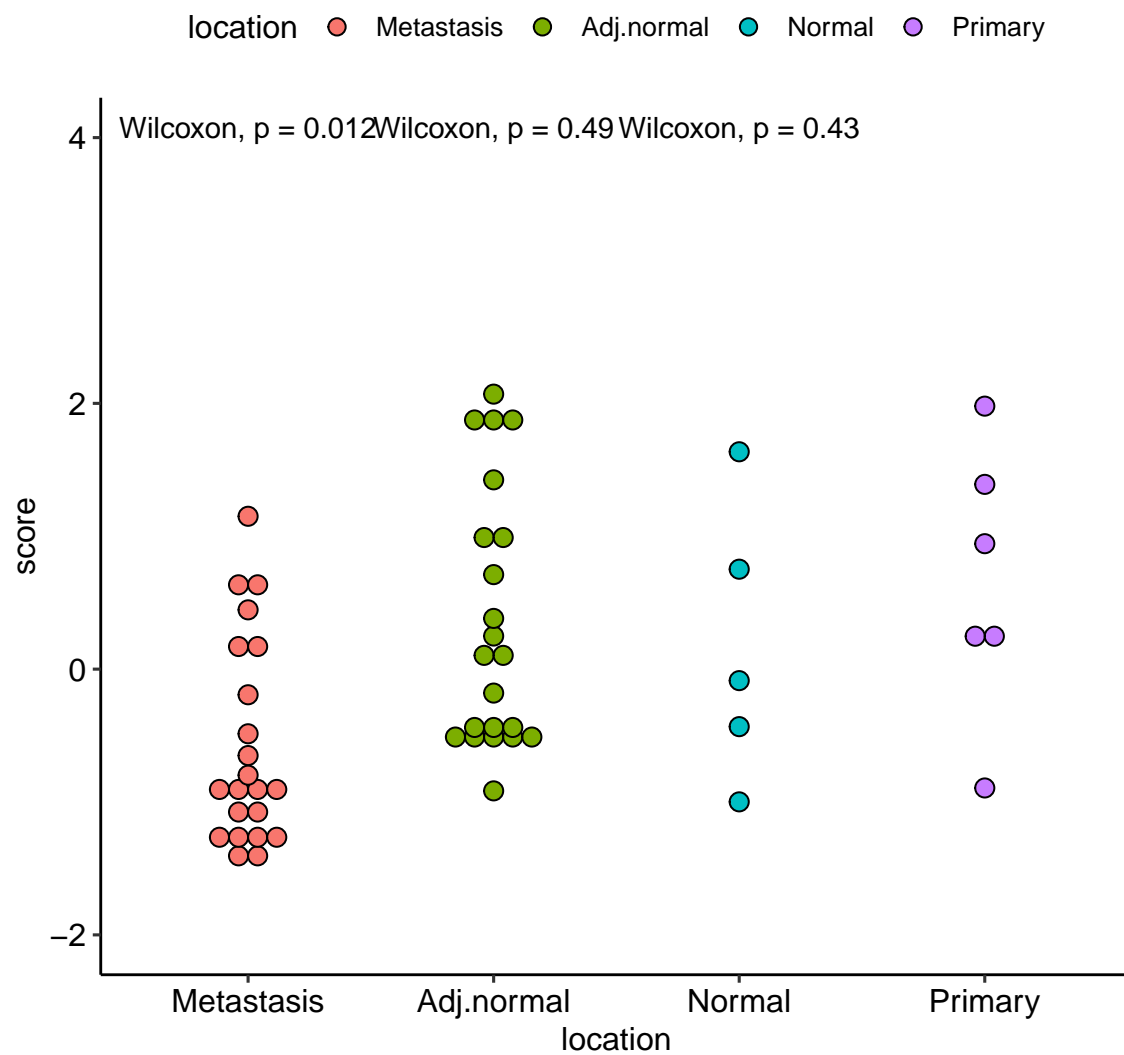


## molgrad\_PDX score

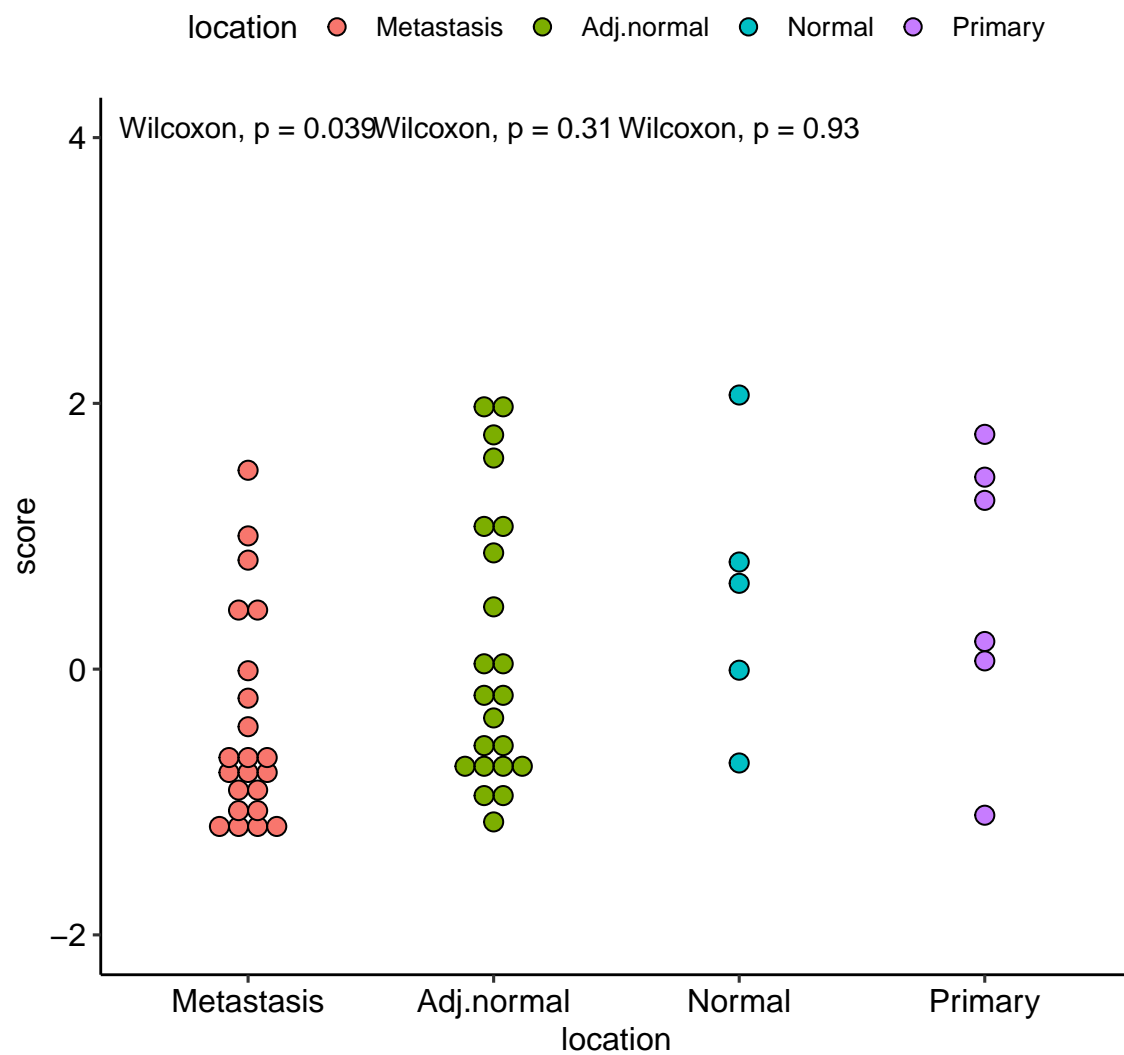
location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



## molgrad\_Puleo score

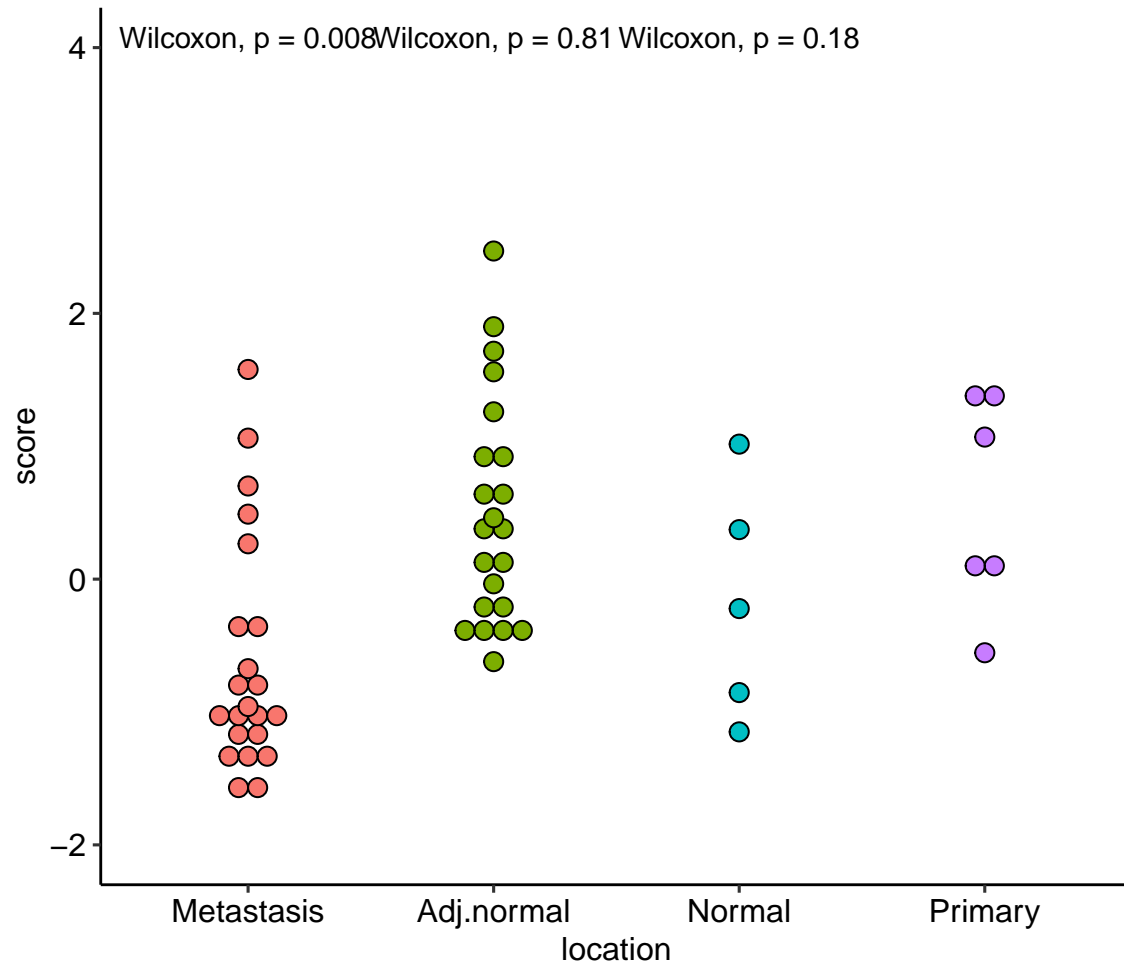


# molgrad\_ICGCarray score



## molgrad\_ICGCrnaseq score

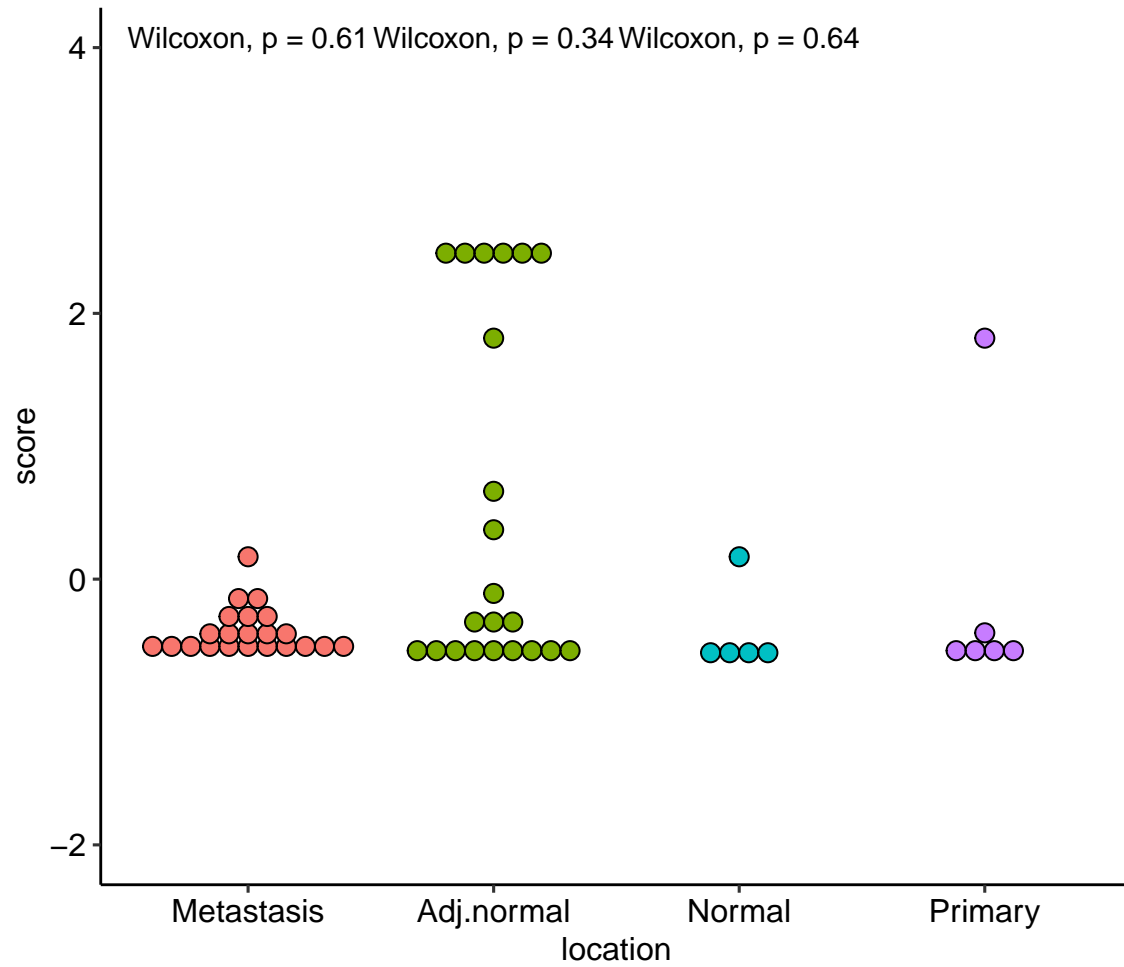
location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary





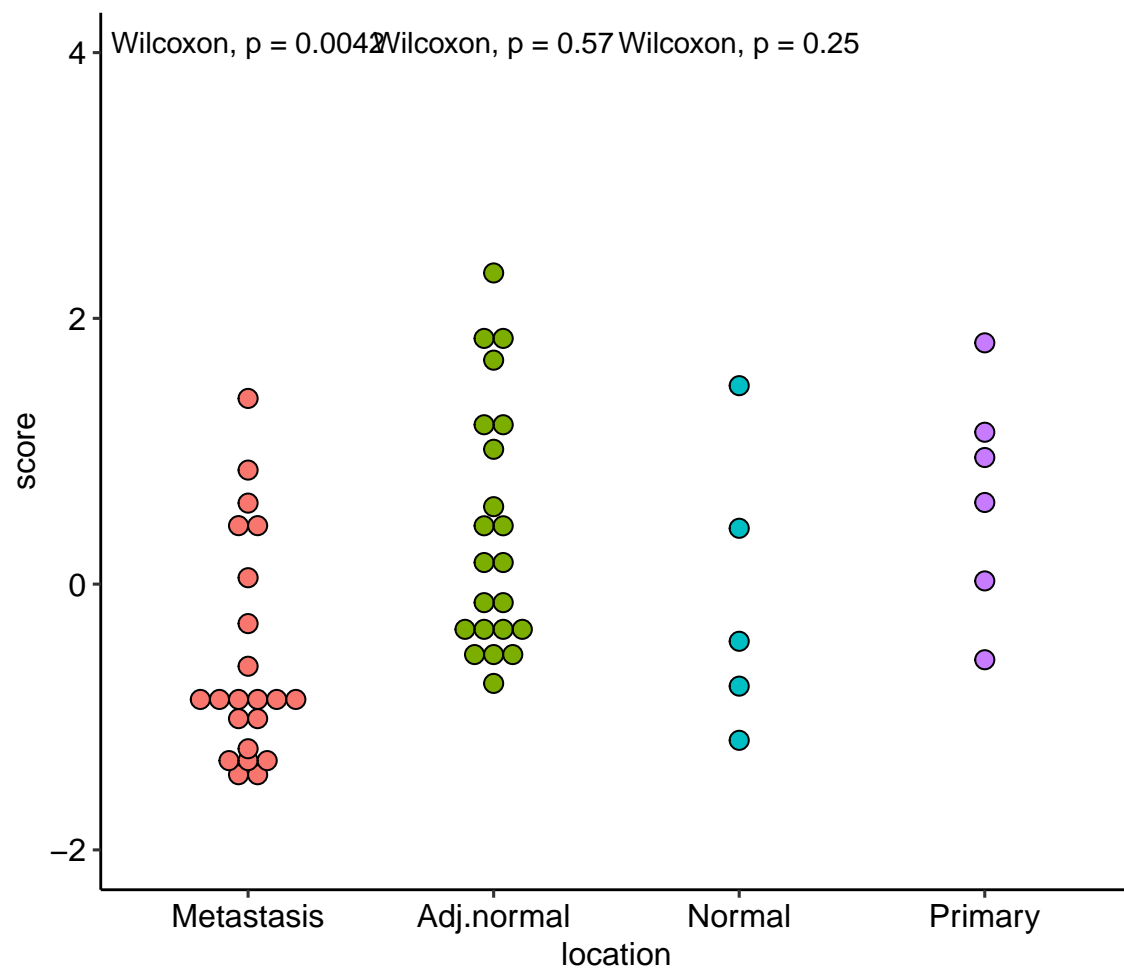
purIST score

location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary

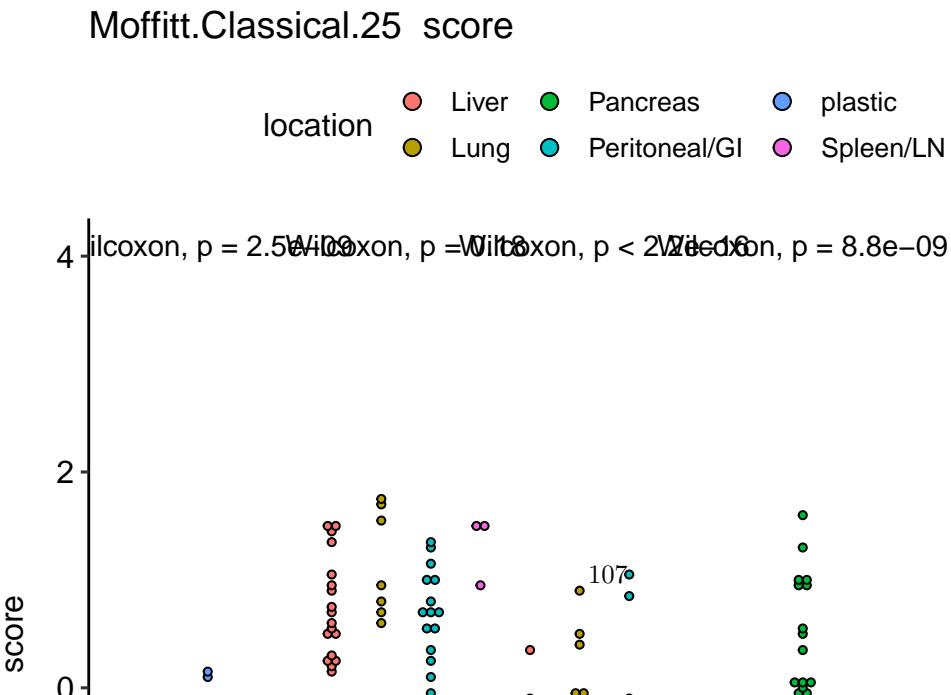
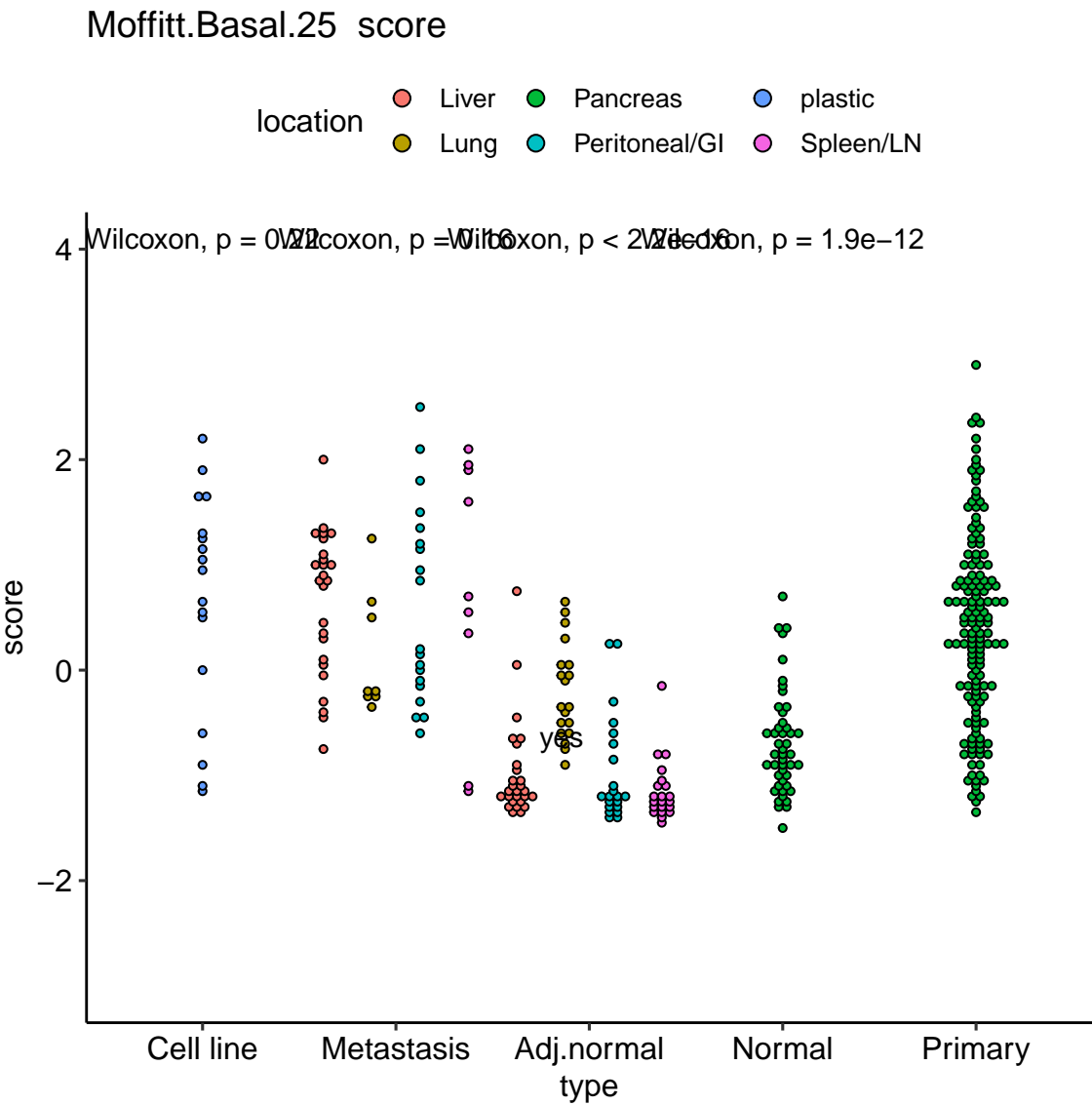


molgrad\_scaled score

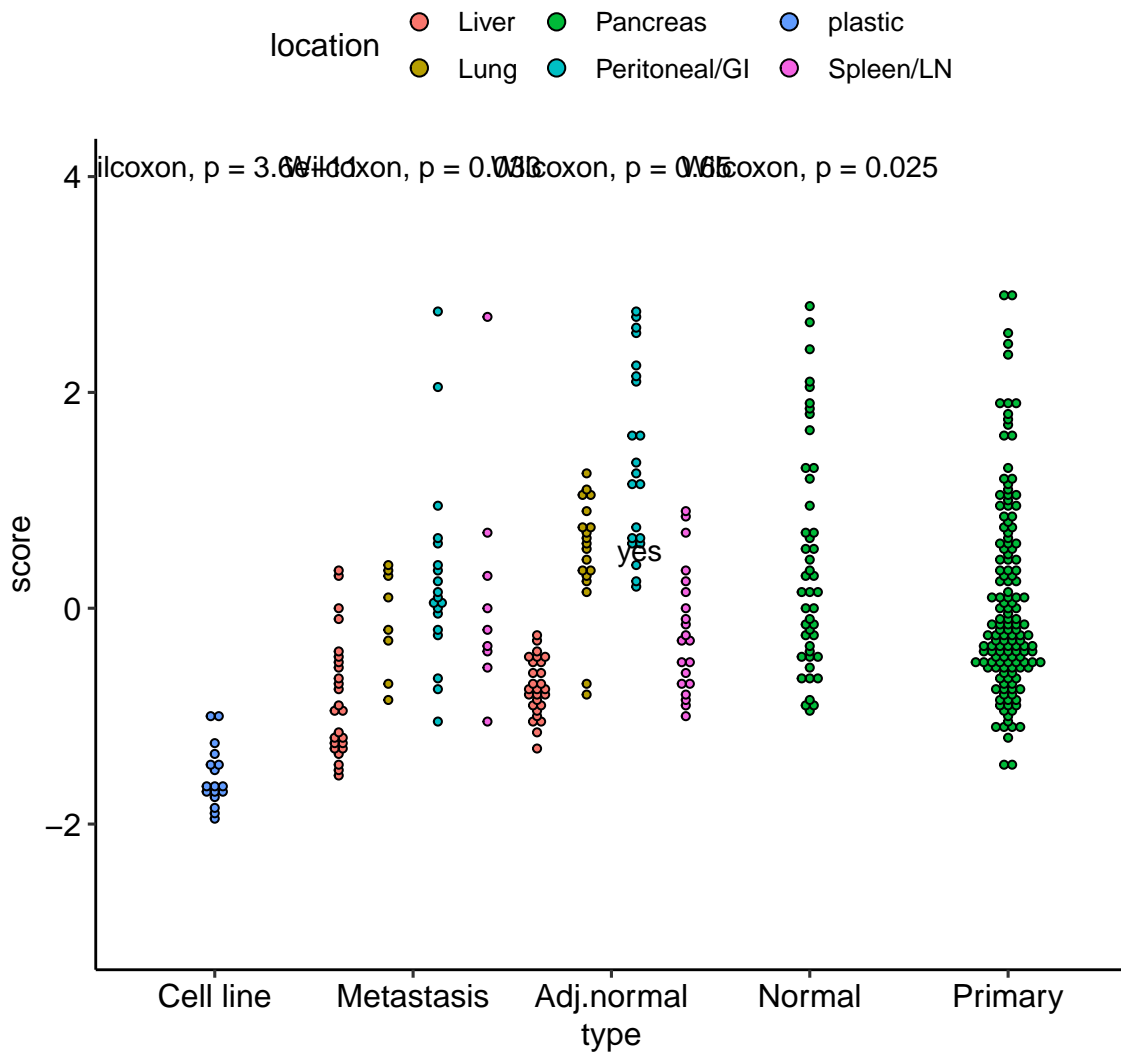
location   ● Metastasis   ● Adj.normal   ● Normal   ● Primary



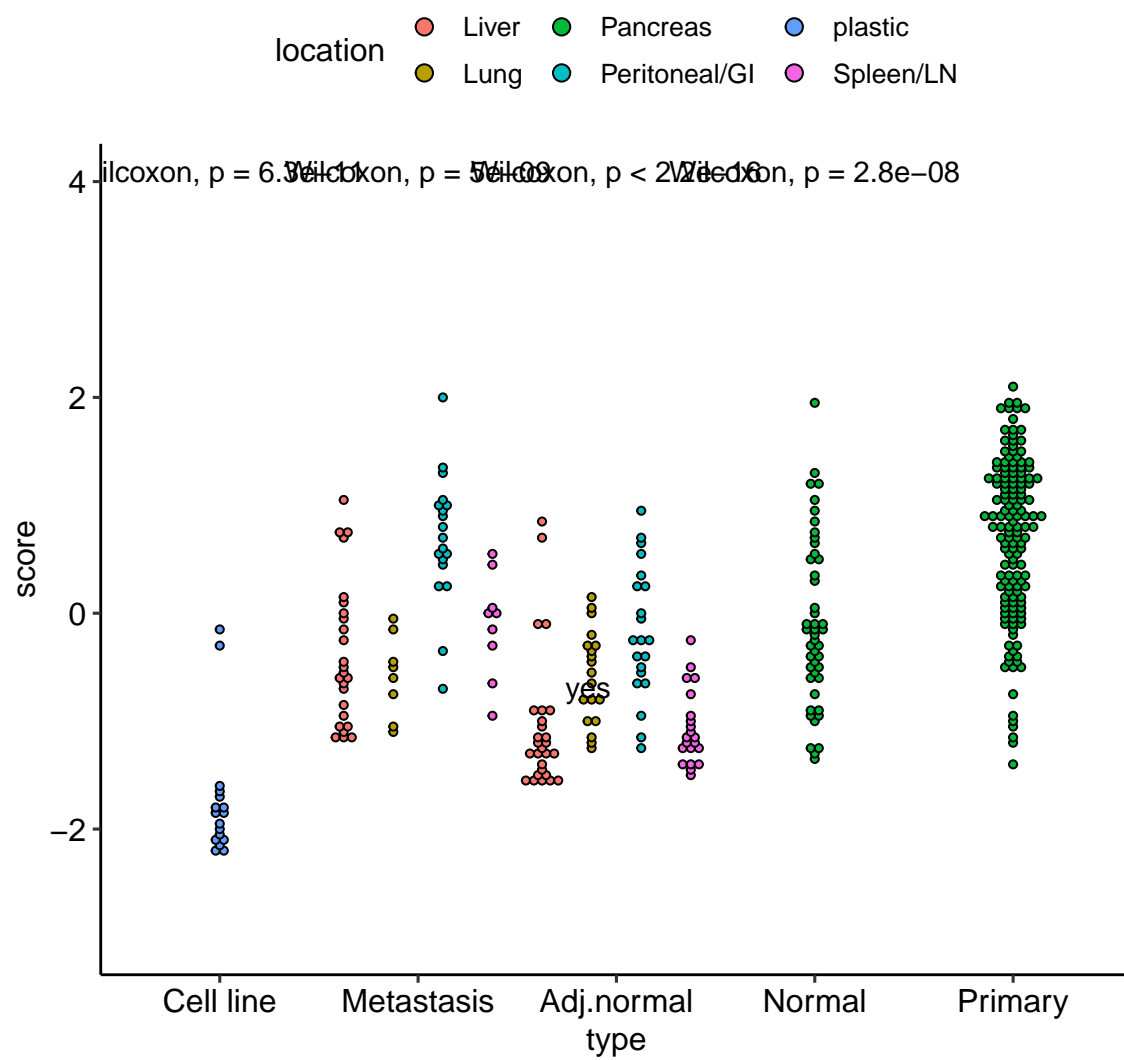
Simplified patient matched sample plots, all samples



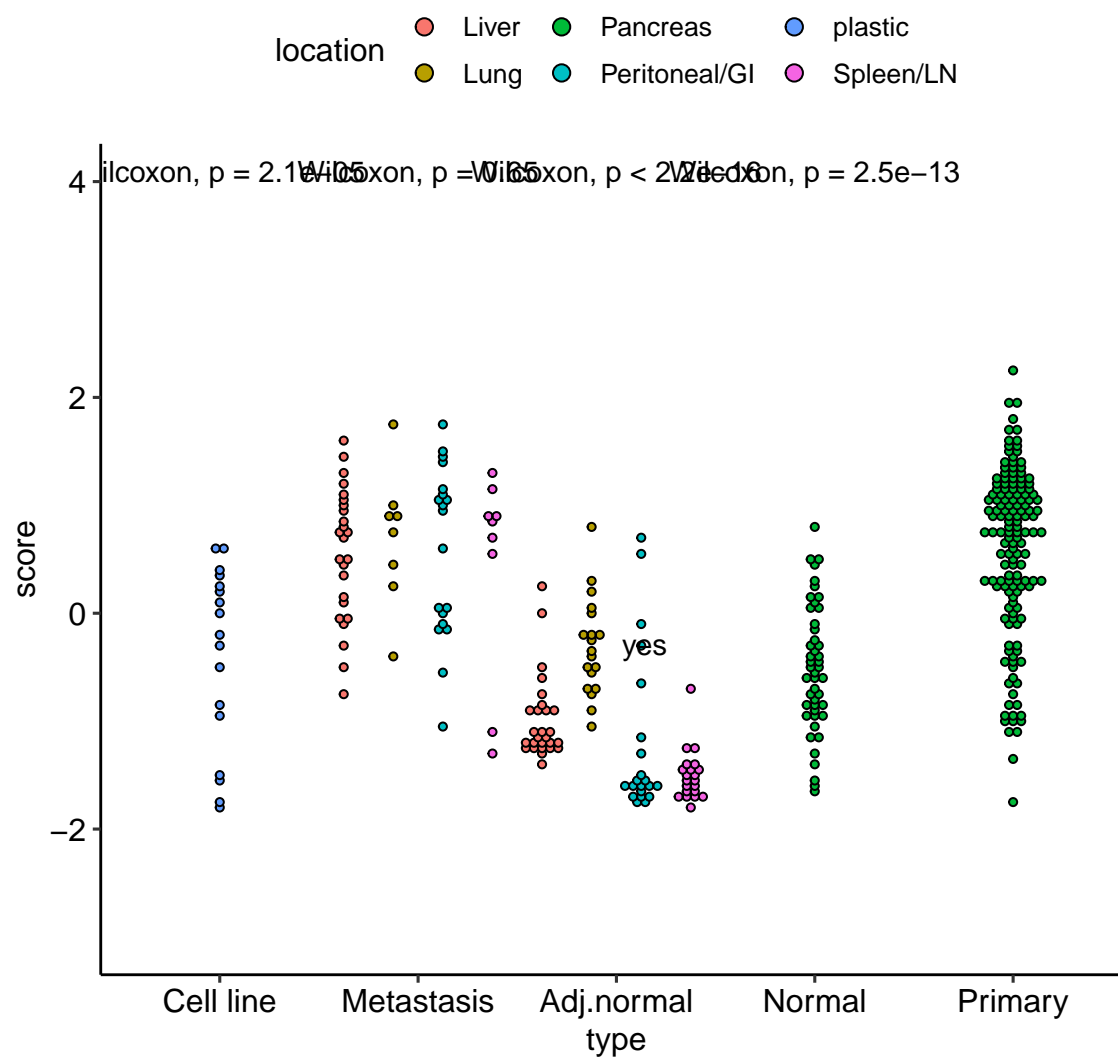
# Moffitt.Normal.25 score



# Moffitt.Activated.25 score



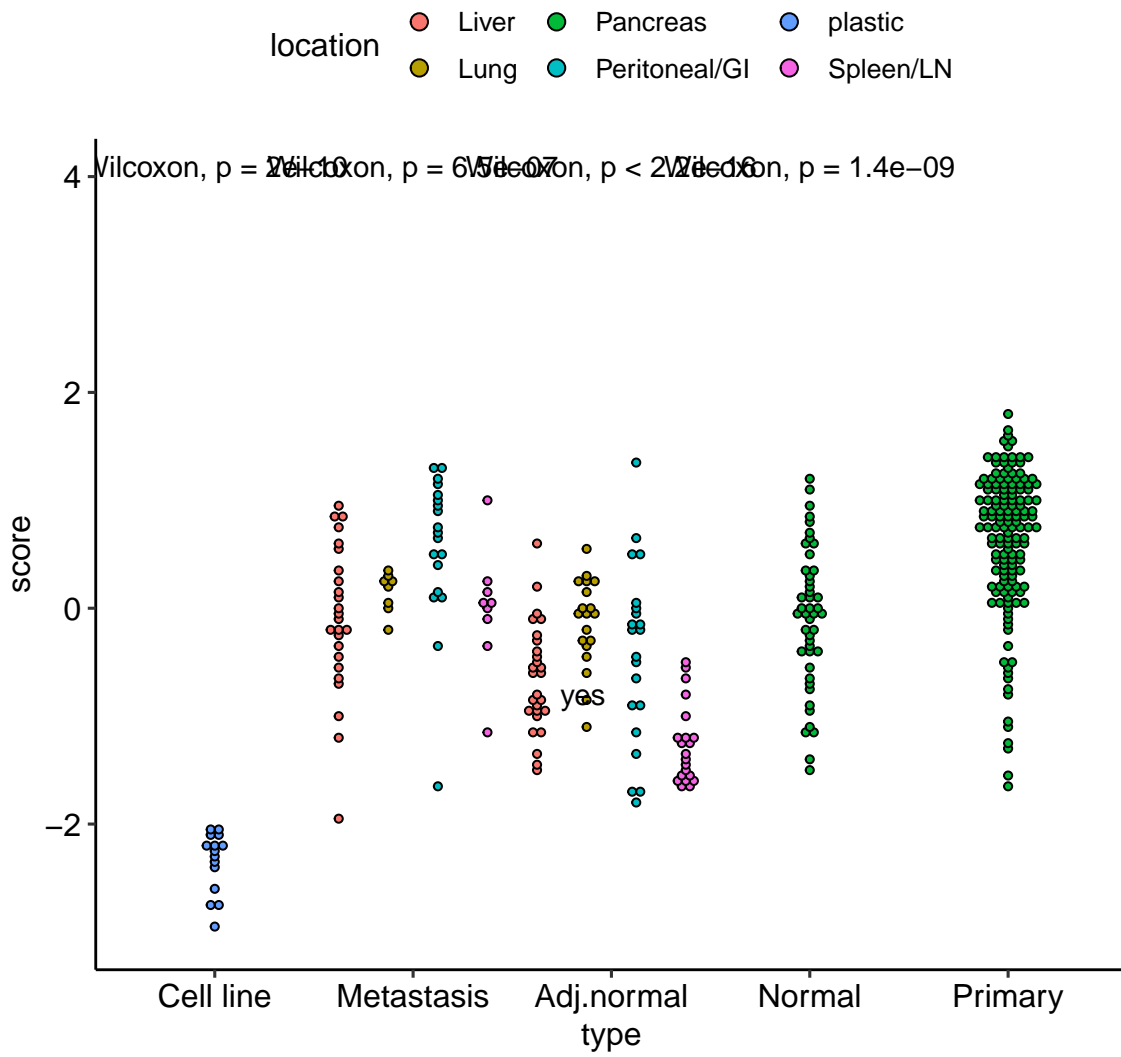
## Moffitt.Tumor score



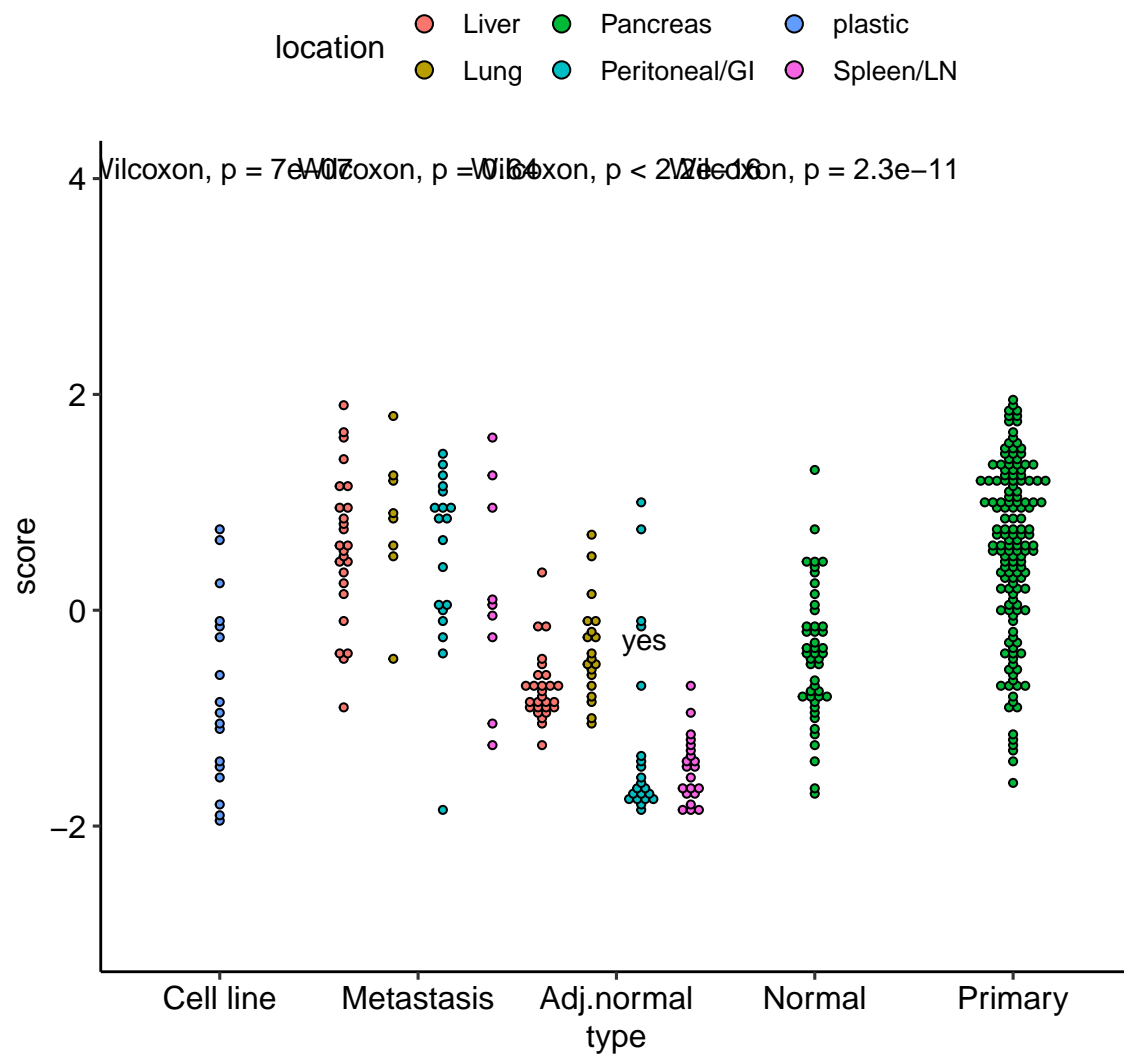
## Warning: Removed 2 rows containing non-finite values (stat\_bindot).

## Warning: Removed 2 rows containing non-finite values (stat\_compare\_means).

## Puleo.Centroids score

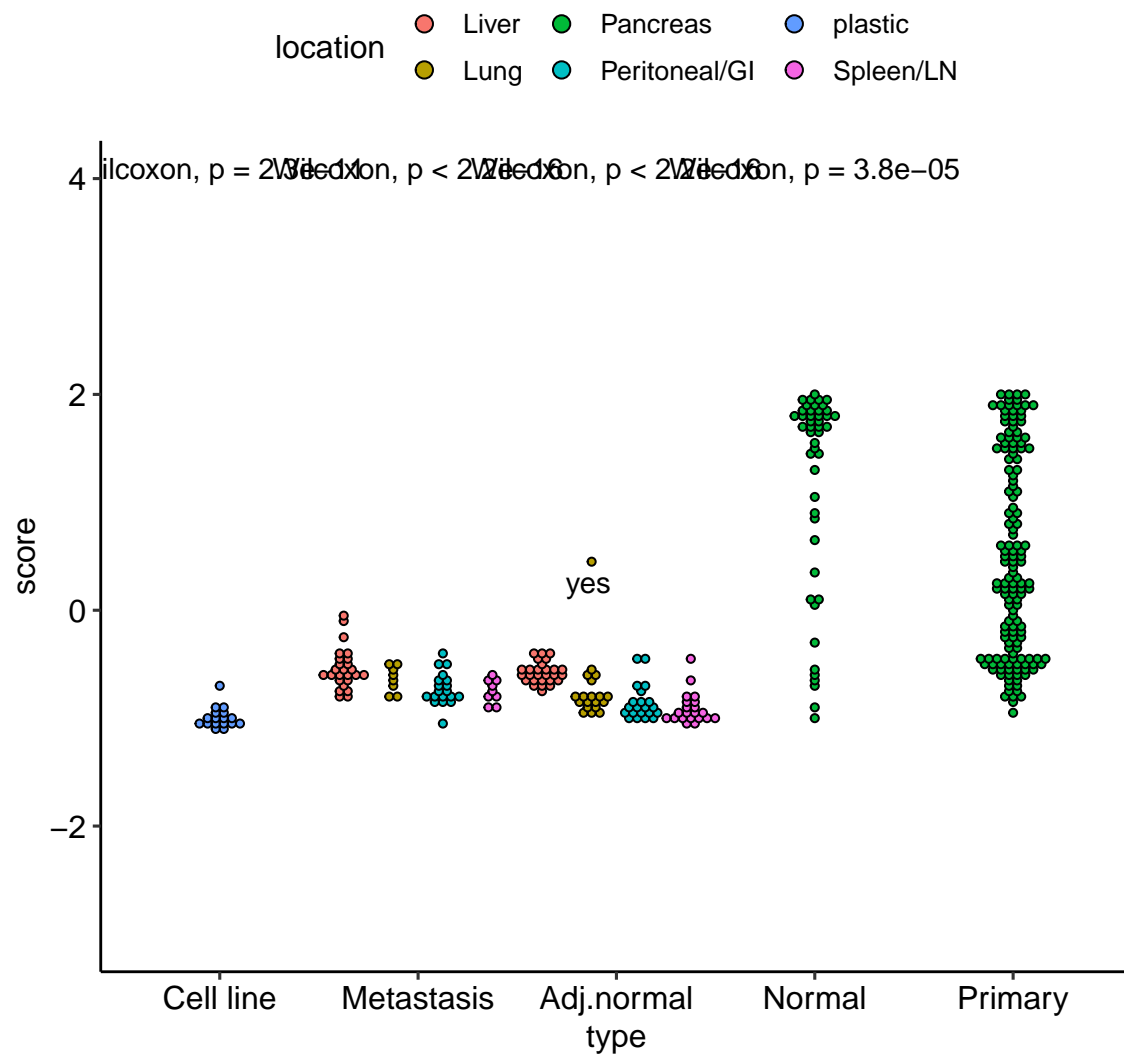


## Collisson.Classical score

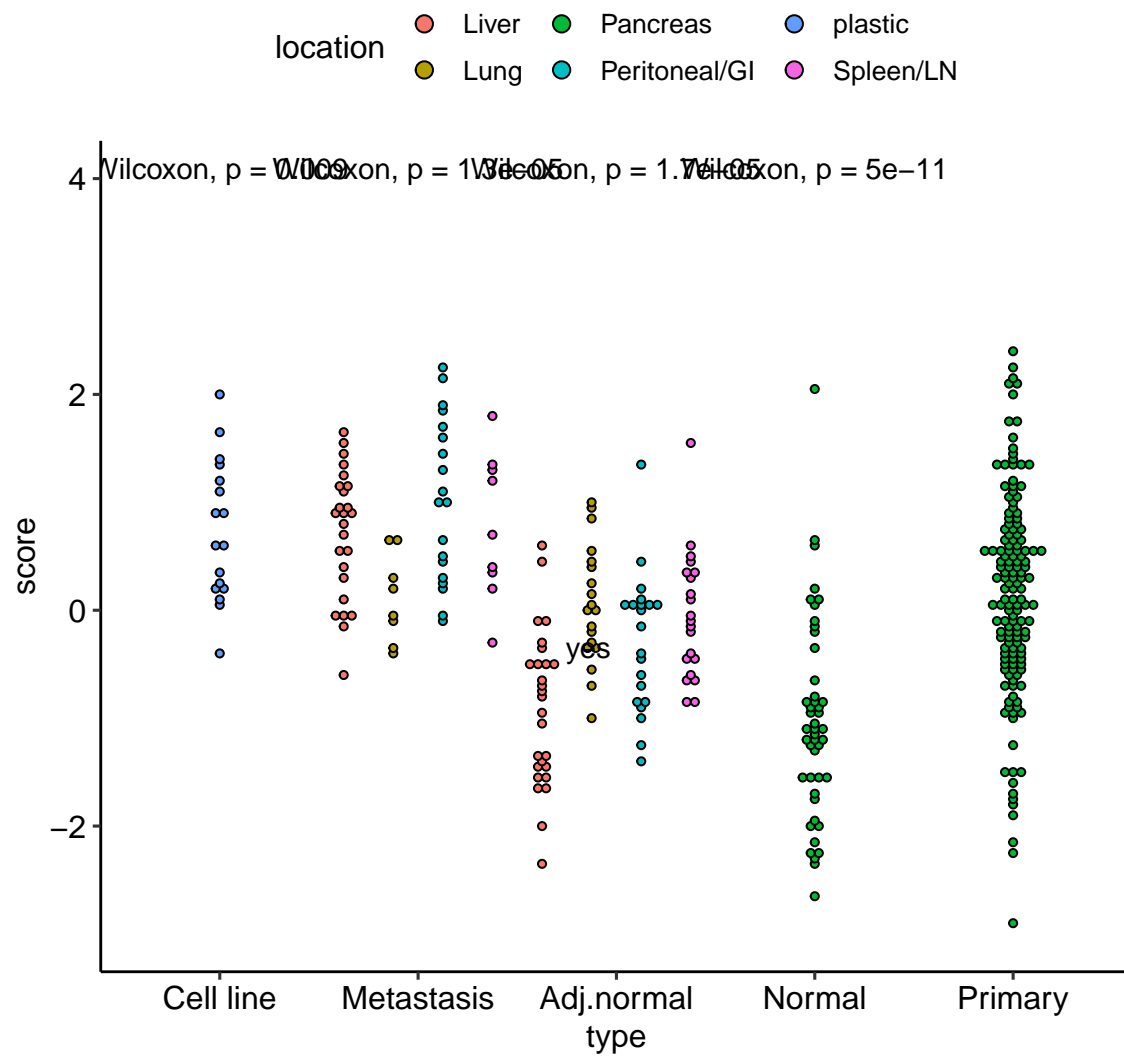




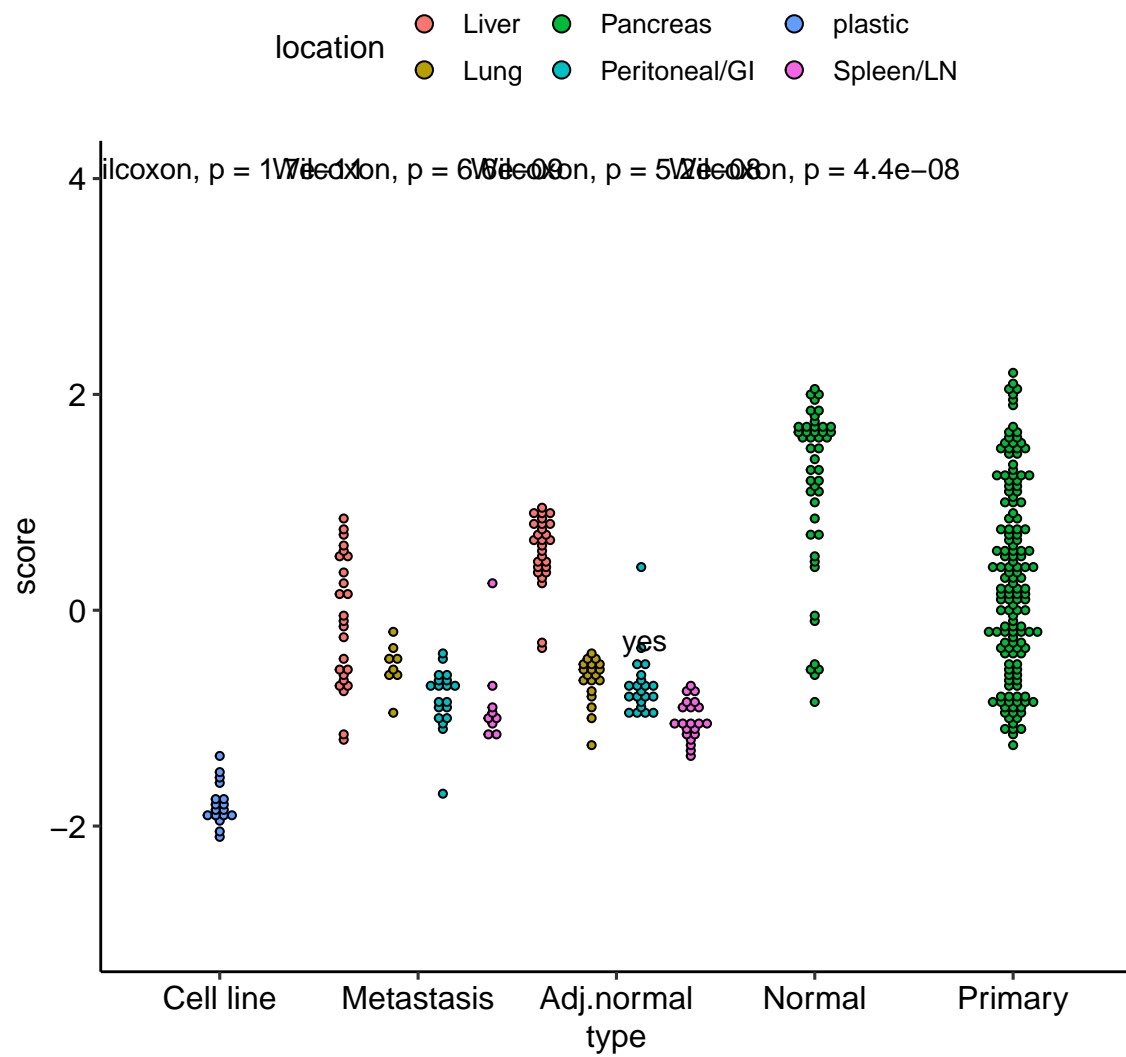
## Collisson.Exocrine score



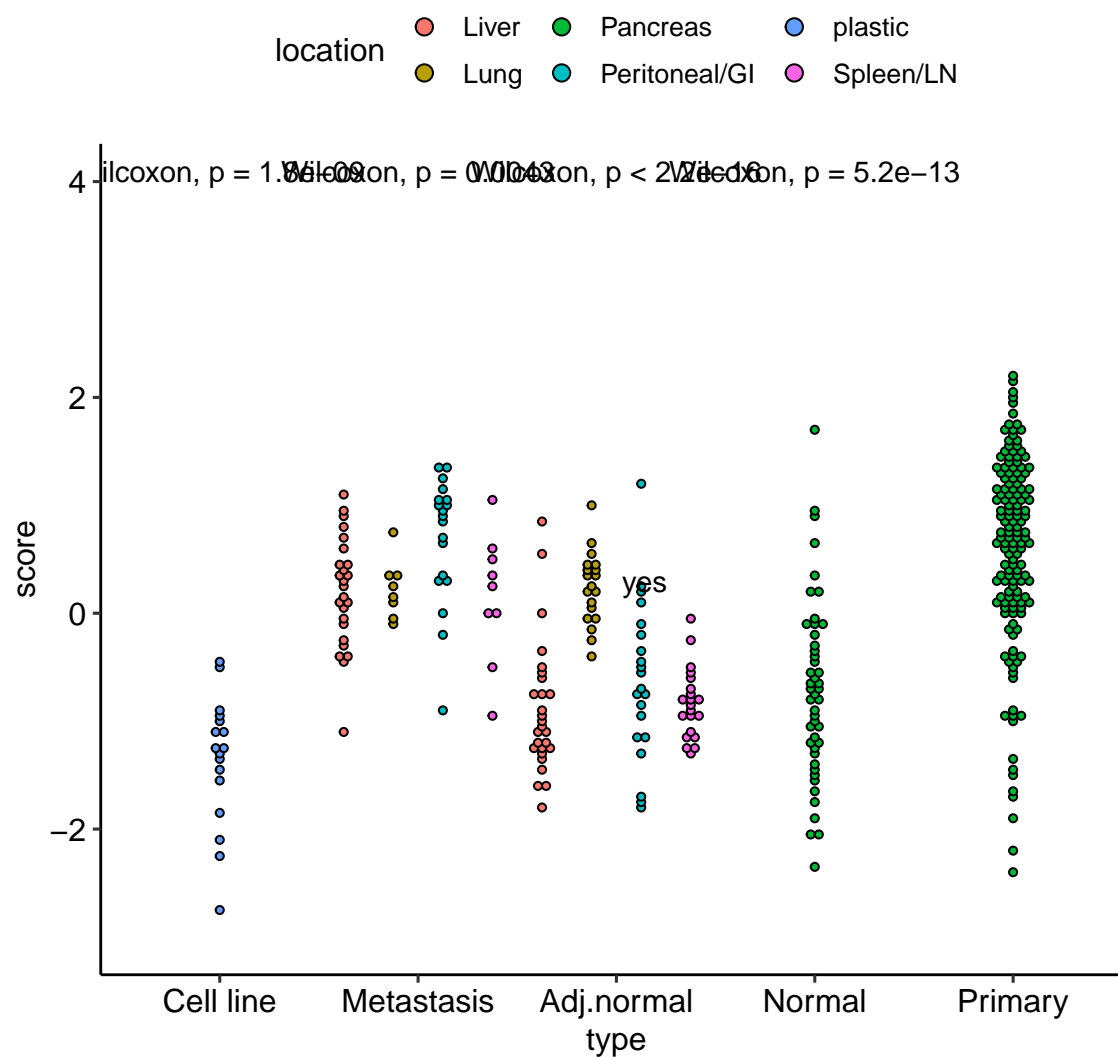
## Collisson.QM score



# ICGC.ADEX.Up score



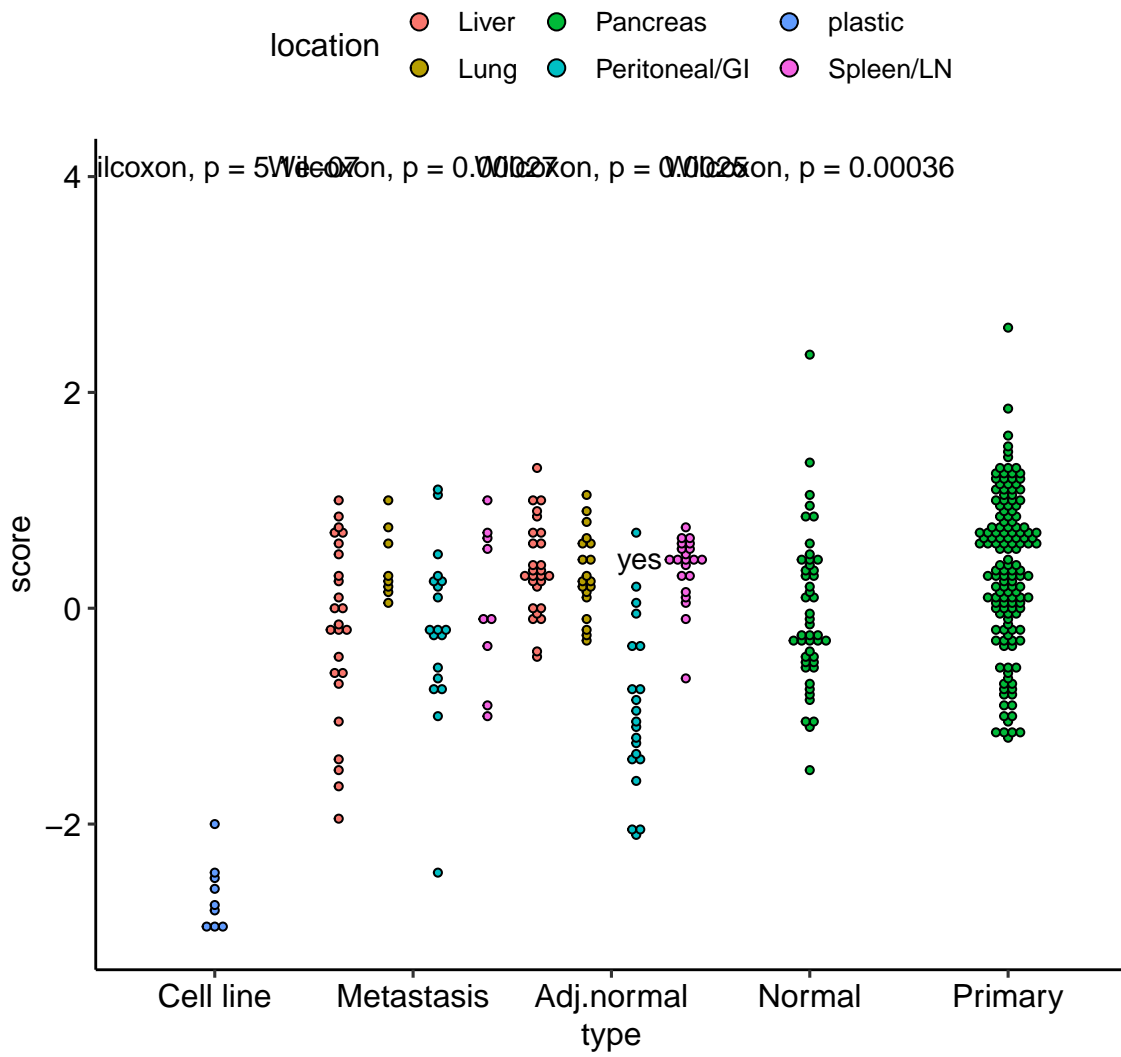
## ICGC.ADEX.Down score



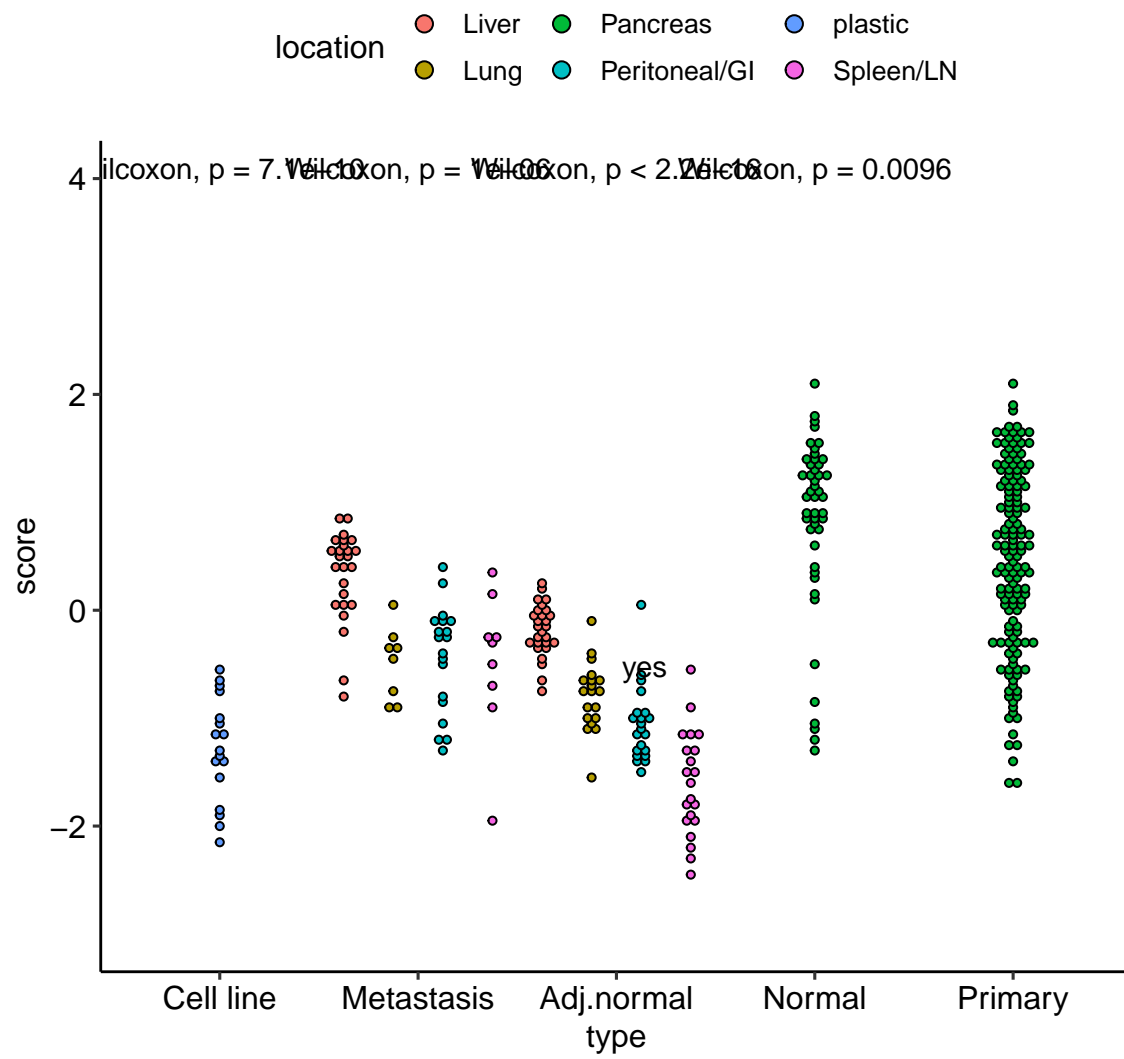
## Warning: Removed 8 rows containing non-finite values (stat\_bindot).

## Warning: Removed 8 rows containing non-finite values (stat\_compare\_means).

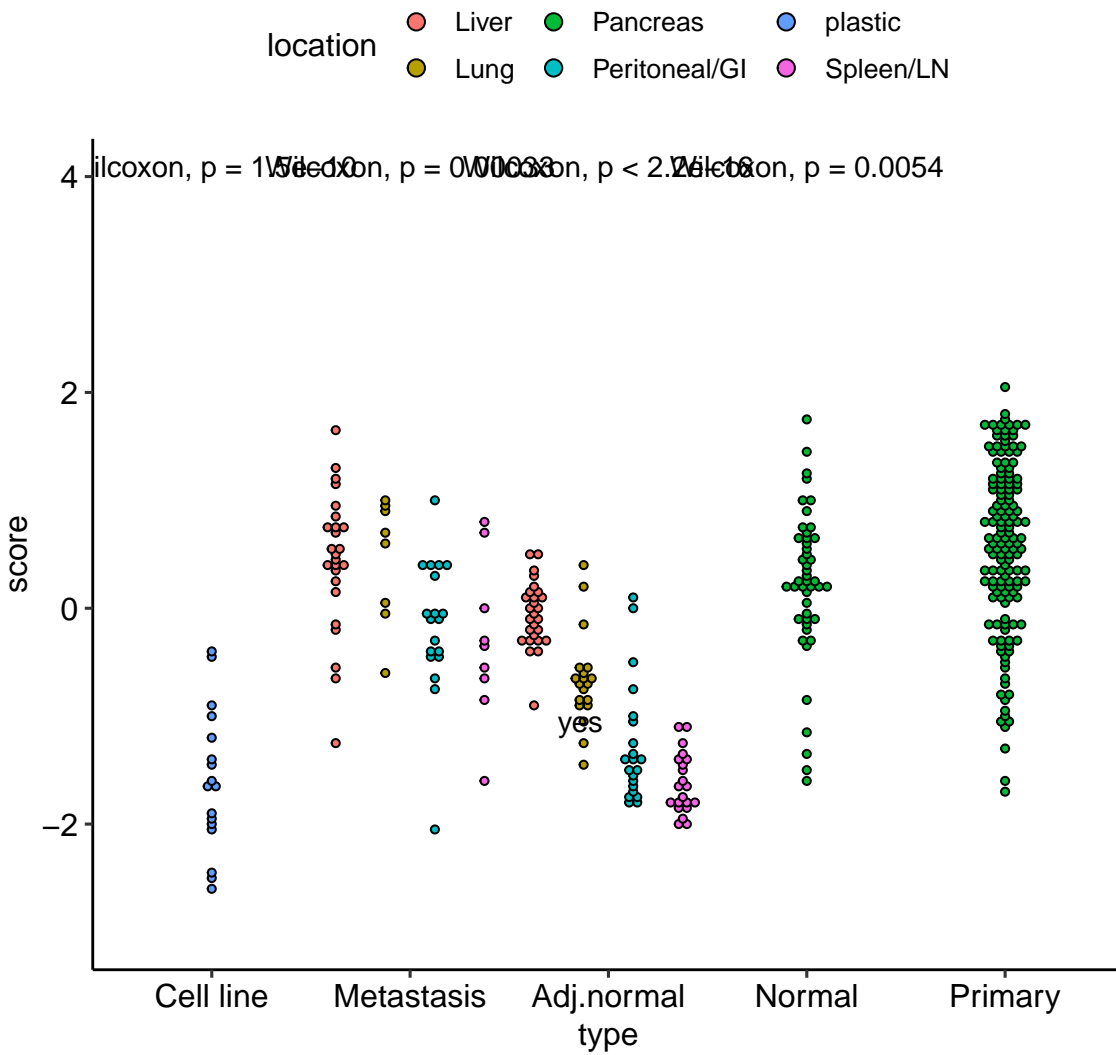
# ICGC.Immunogenic.Up score



# ICGC.Immunogenic.Down score



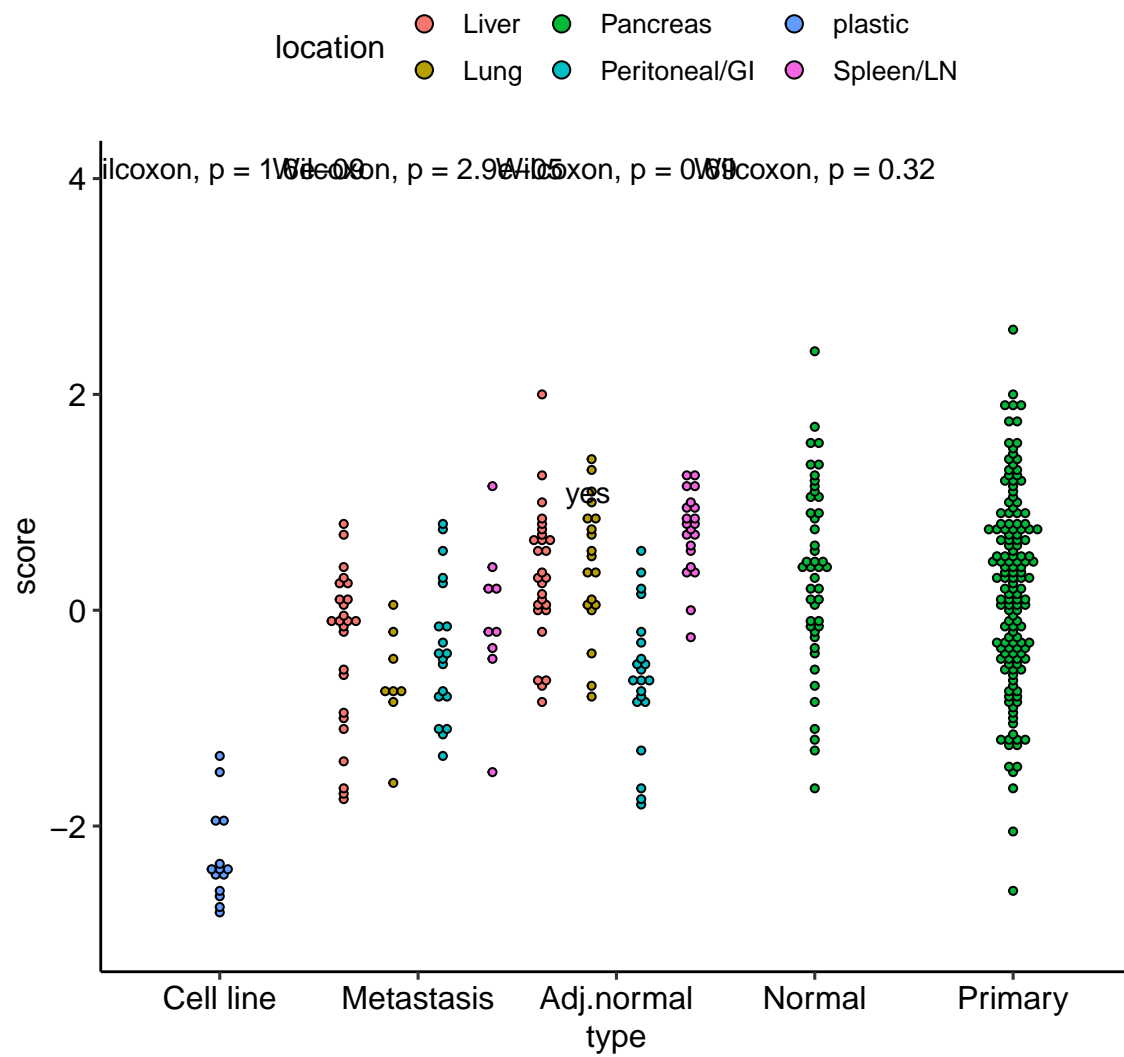
## ICGC.Progenitor.Up score



## Warning: Removed 3 rows containing non-finite values (stat\_bindot).

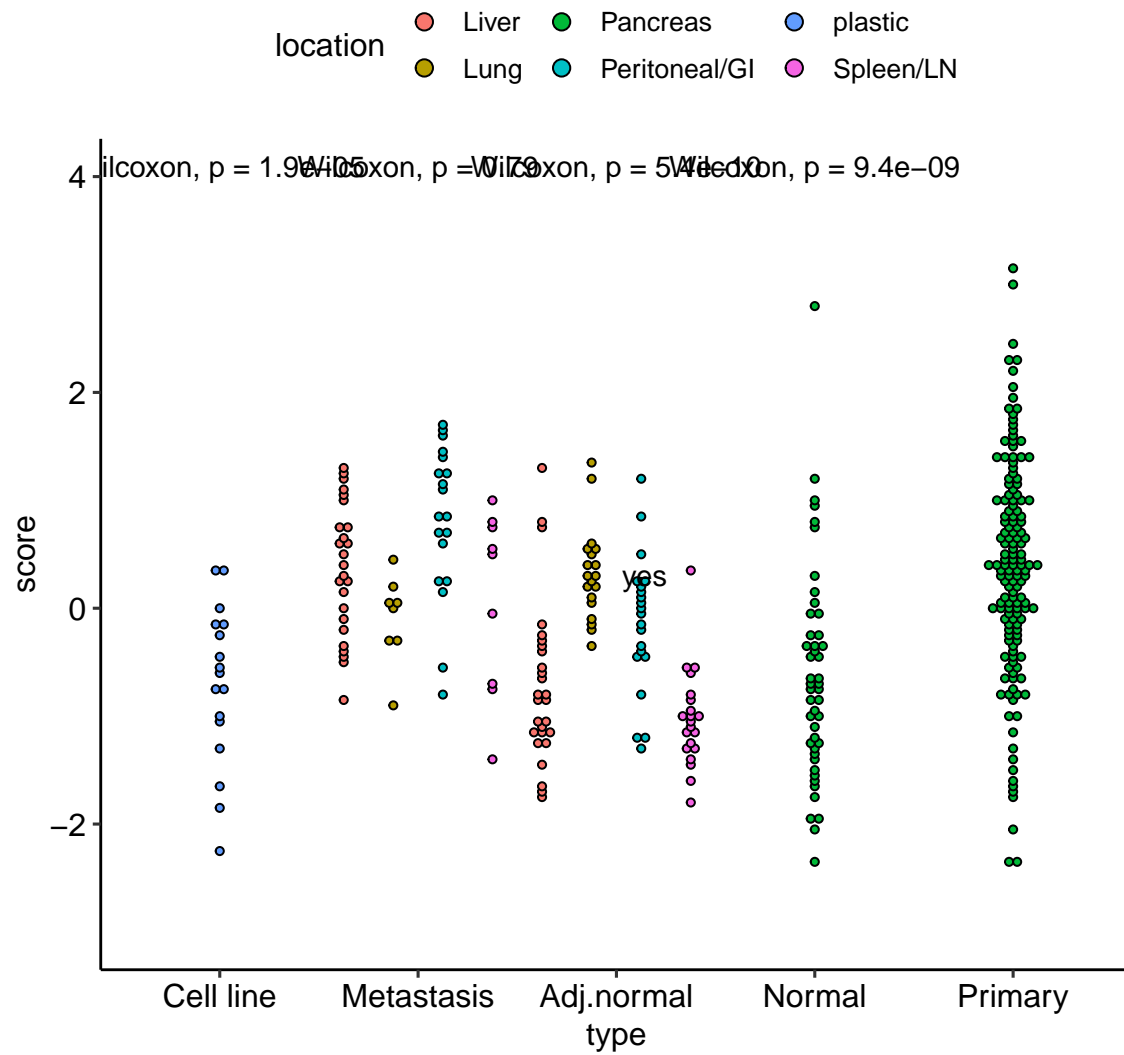
## Warning: Removed 3 rows containing non-finite values (stat\_compare\_means).

# ICGC.Progenitor.Down score





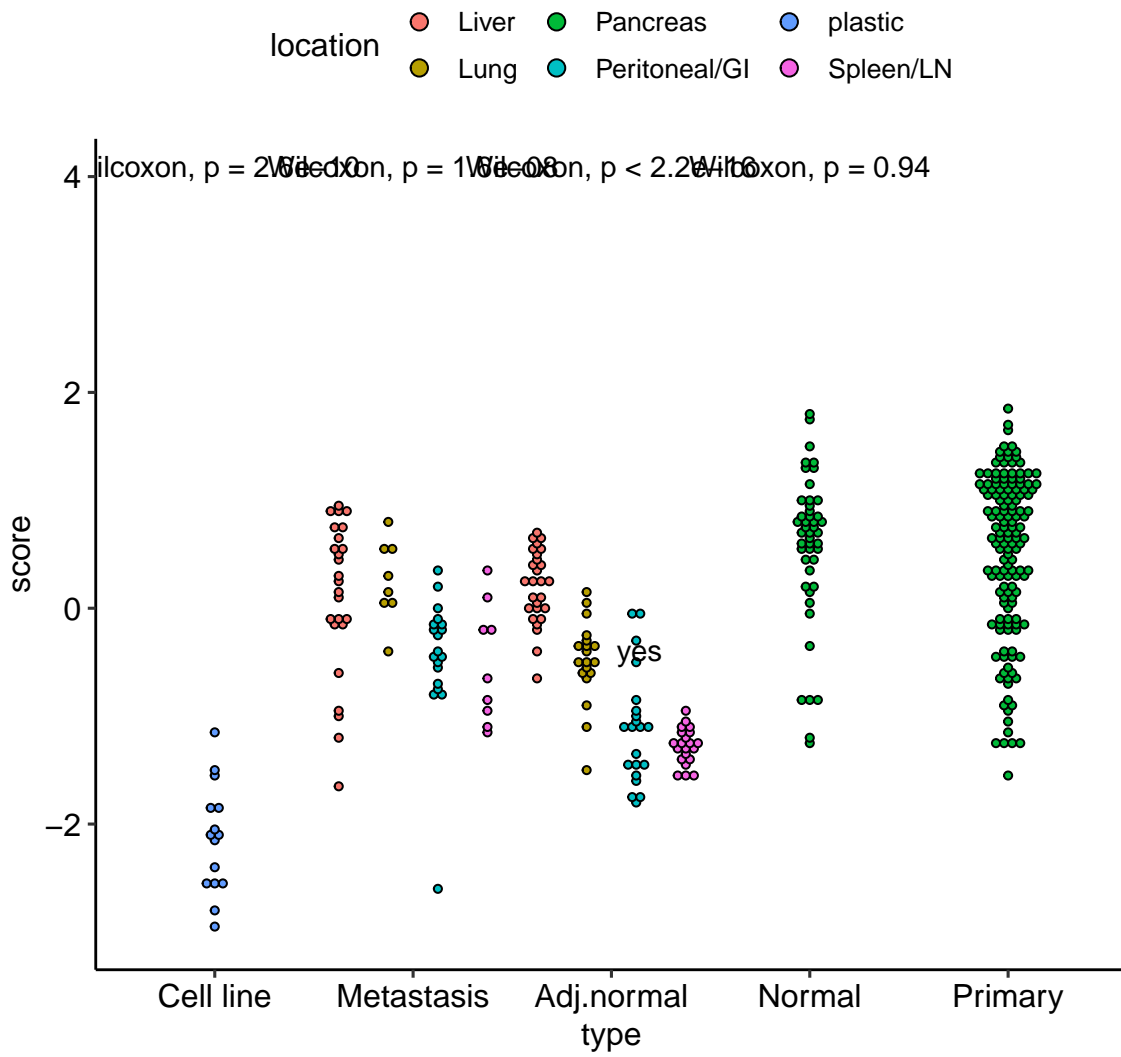
## ICGC.Squamous.Up score



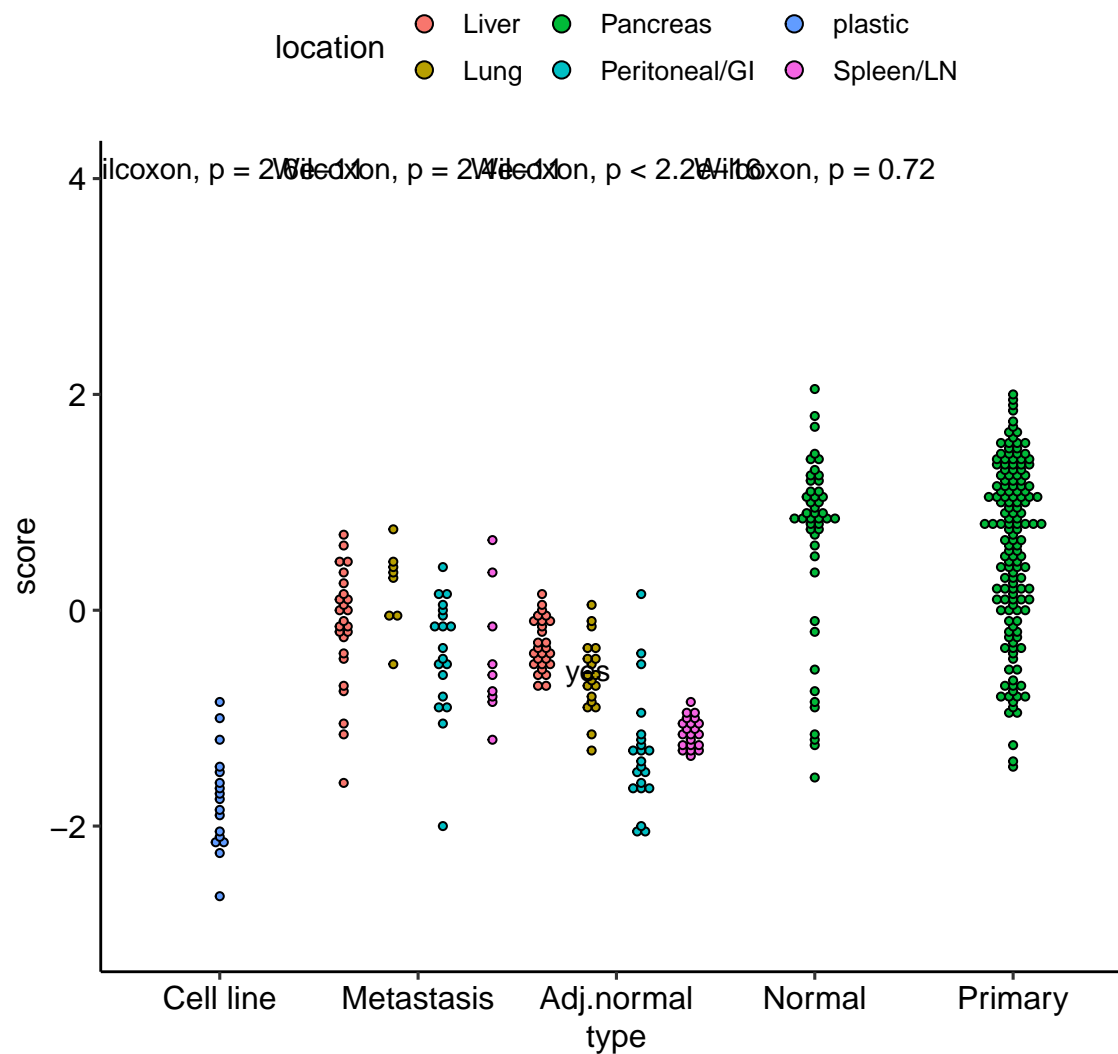
## Warning: Removed 2 rows containing non-finite values (stat\_bindot).

## Warning: Removed 2 rows containing non-finite values (stat\_compare\_means).

# ICGC.Squamous.Down score



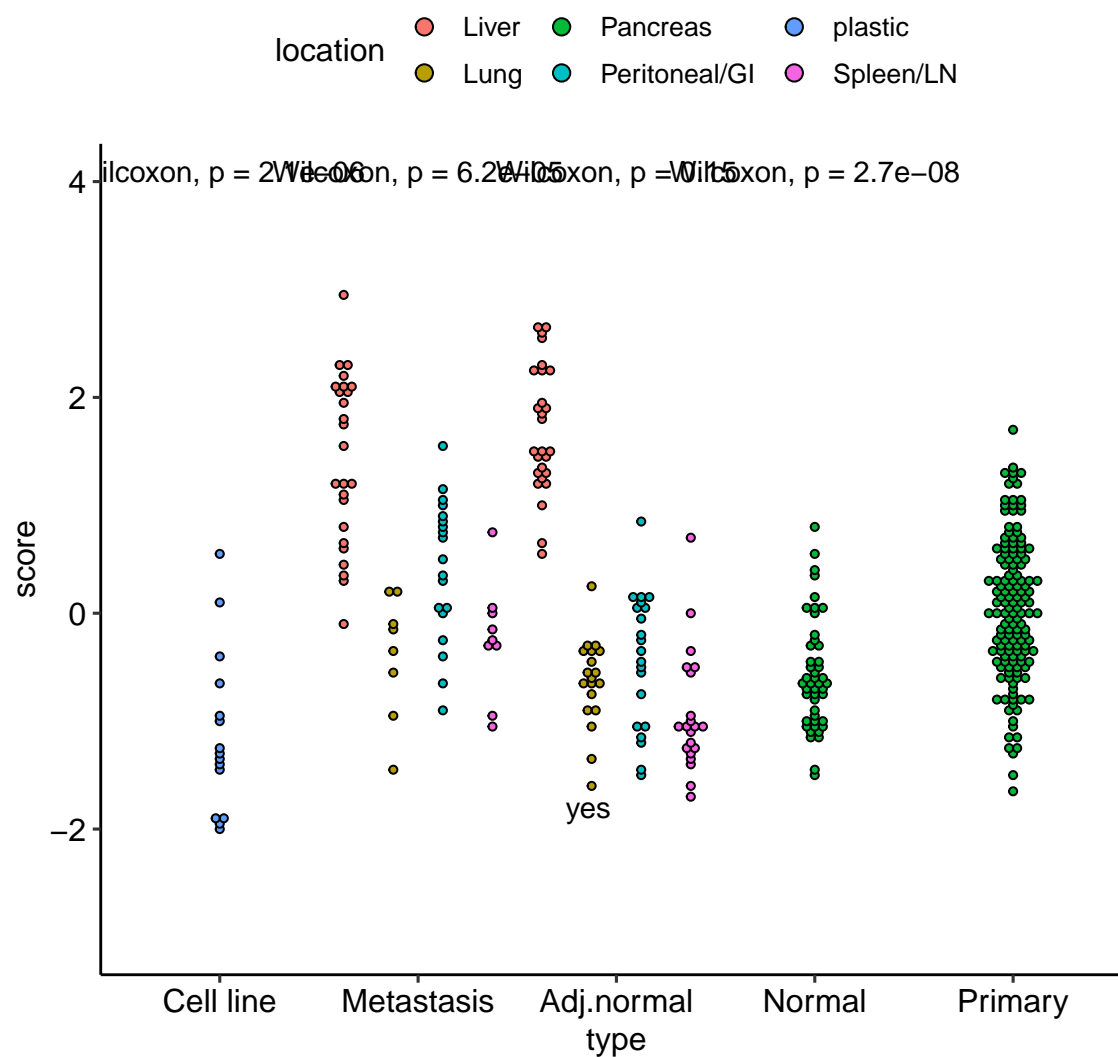
## ICGC.SAM score



## Warning: Removed 2 rows containing non-finite values (stat\_bindot).

## Warning: Removed 2 rows containing non-finite values (stat\_compare\_means).

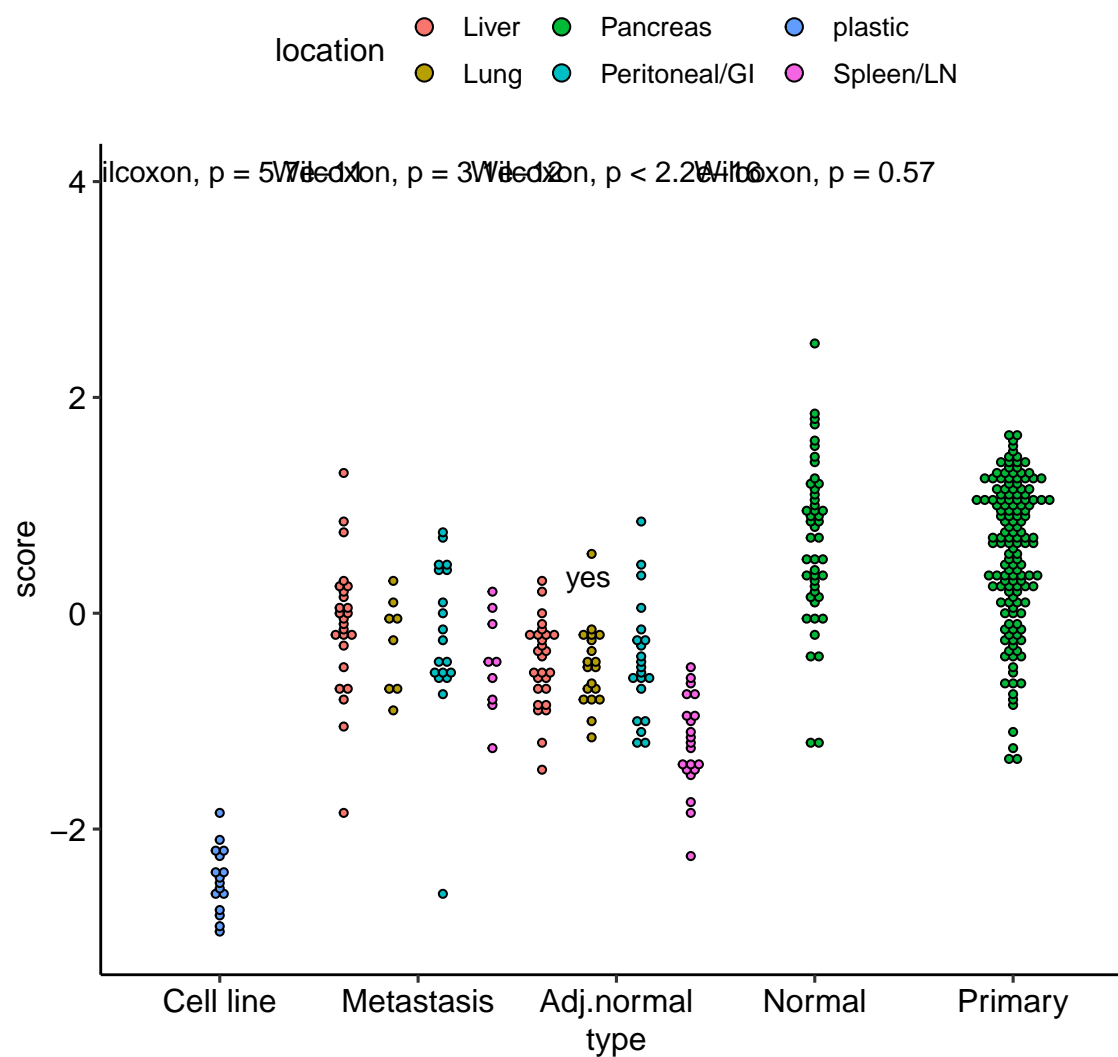
## Scarpa\_PanNET score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

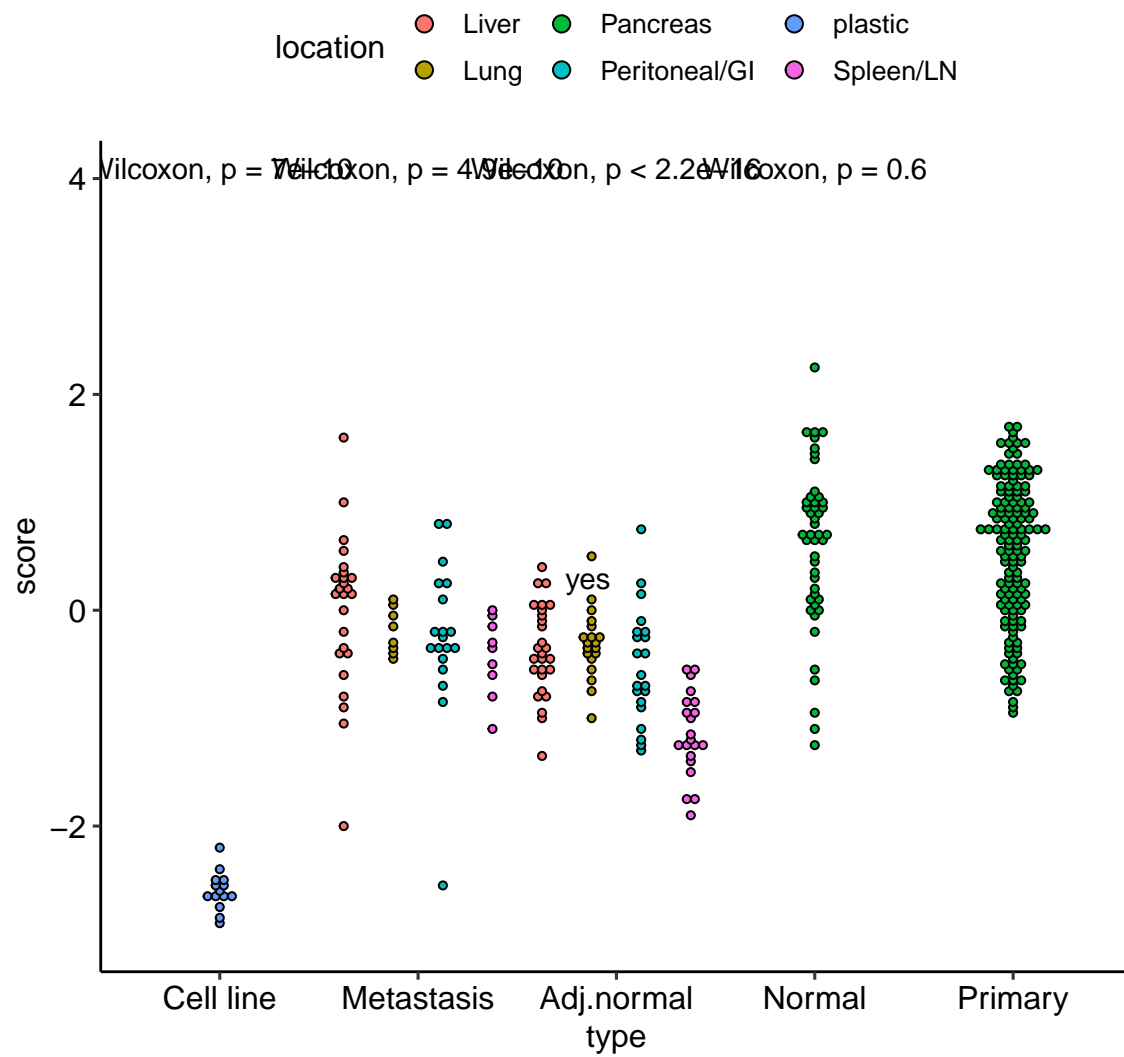
## Moffitt.Top5s score



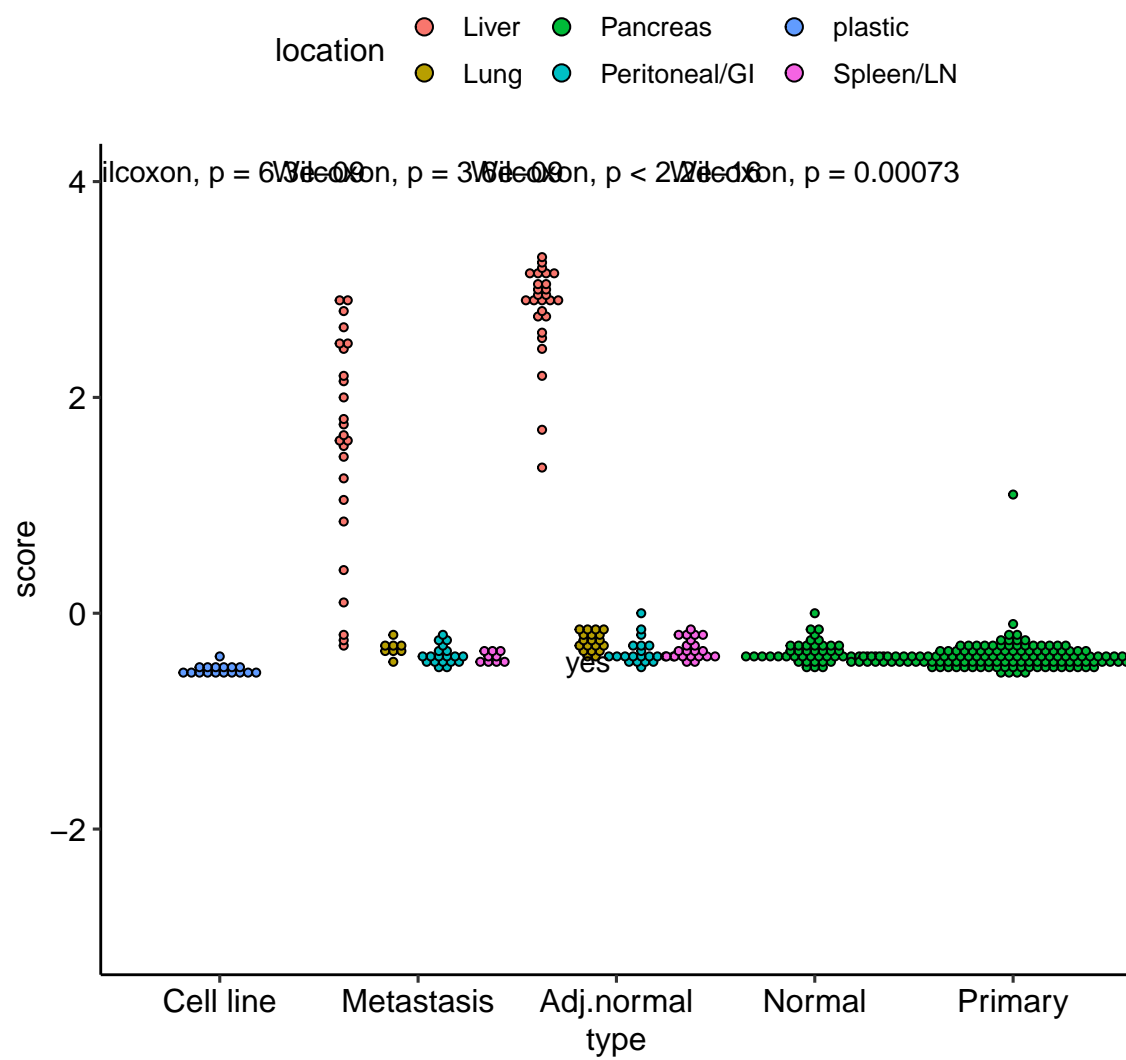
## Warning: Removed 3 rows containing non-finite values (stat\_bindot).

## Warning: Removed 3 rows containing non-finite values (stat\_compare\_means).

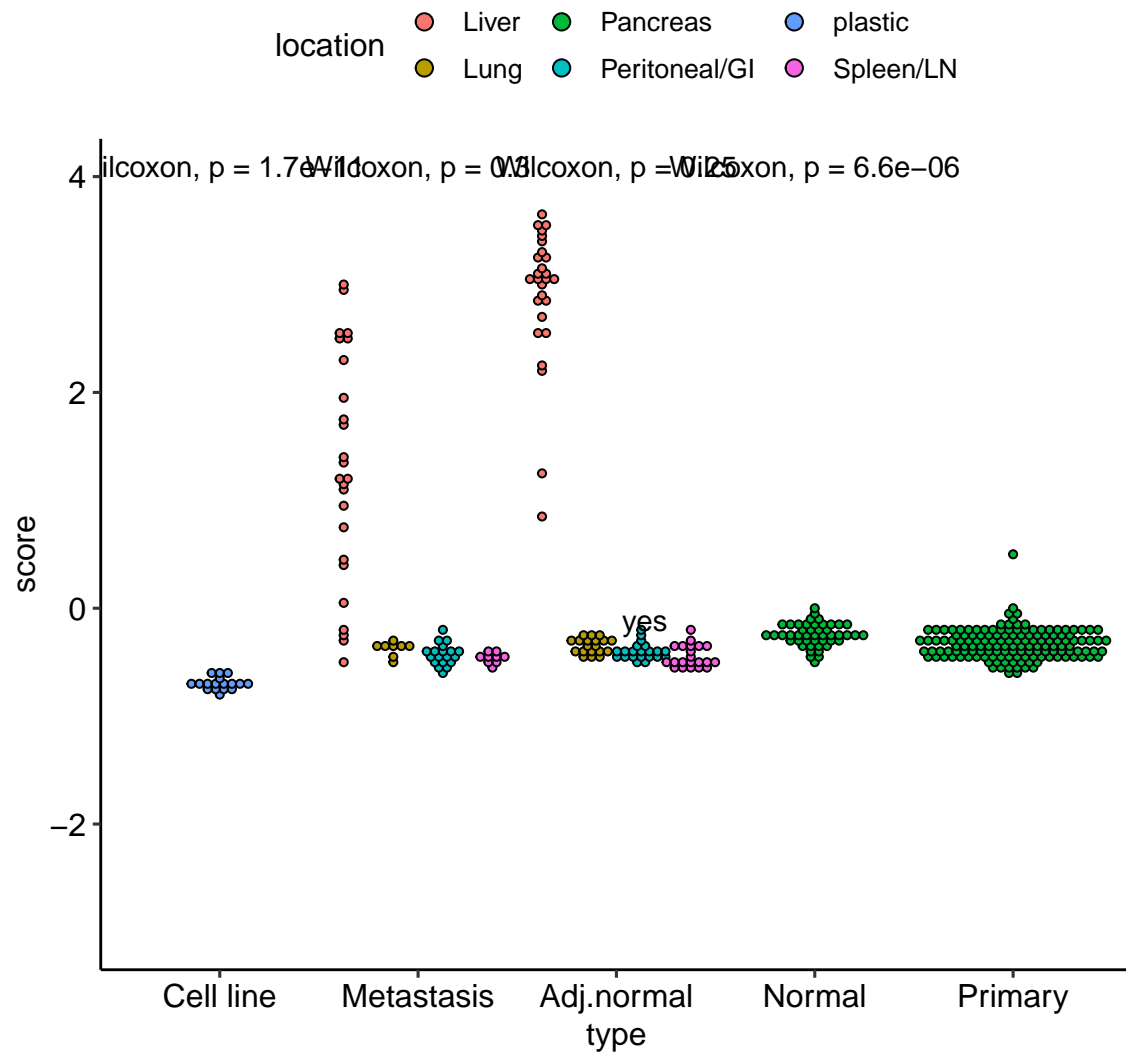
## Moffitt.Top25s score



# Moffitt.F1\_Liver.top25 score

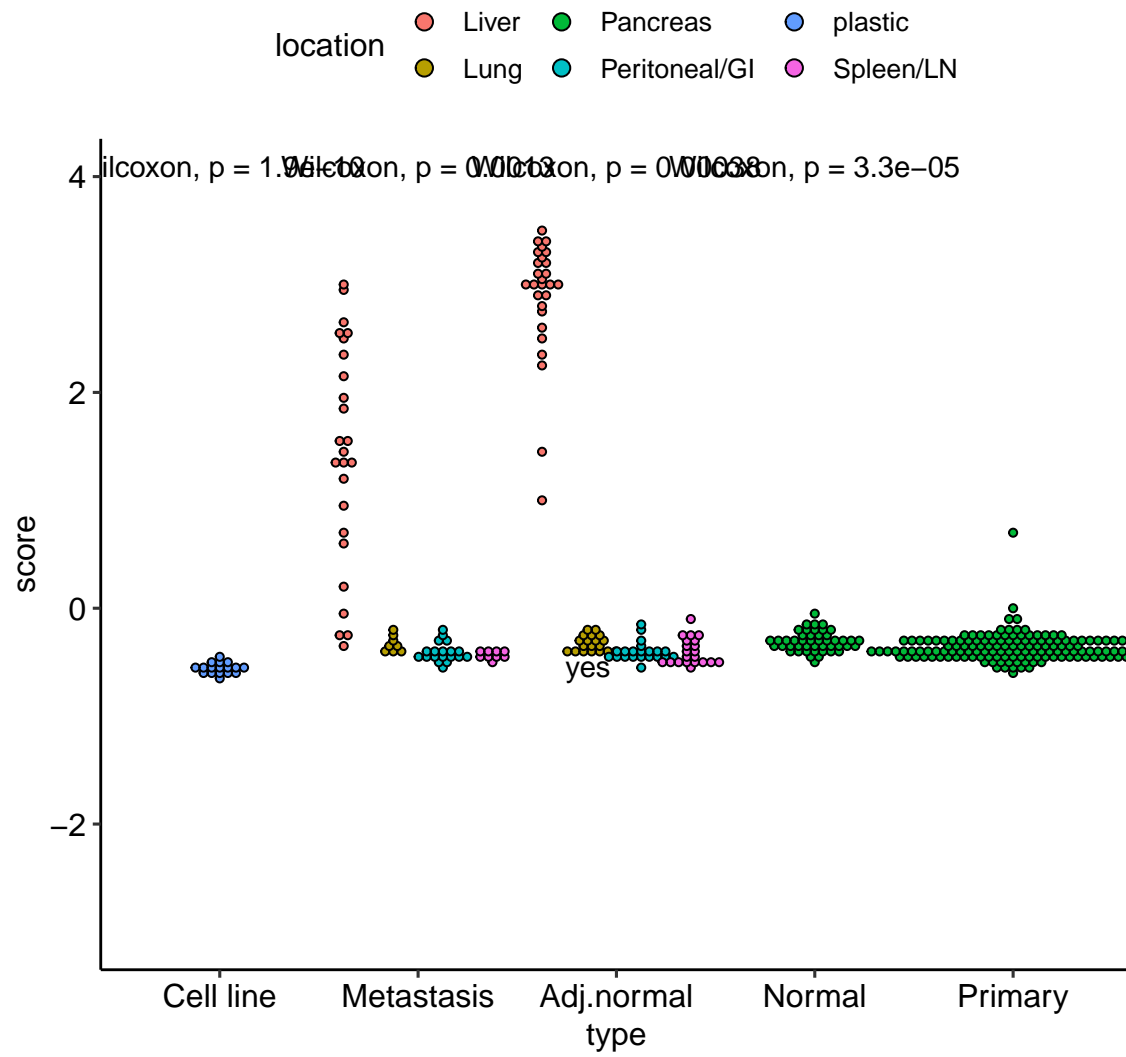


# Moffitt.F1\_Liver.top250 score

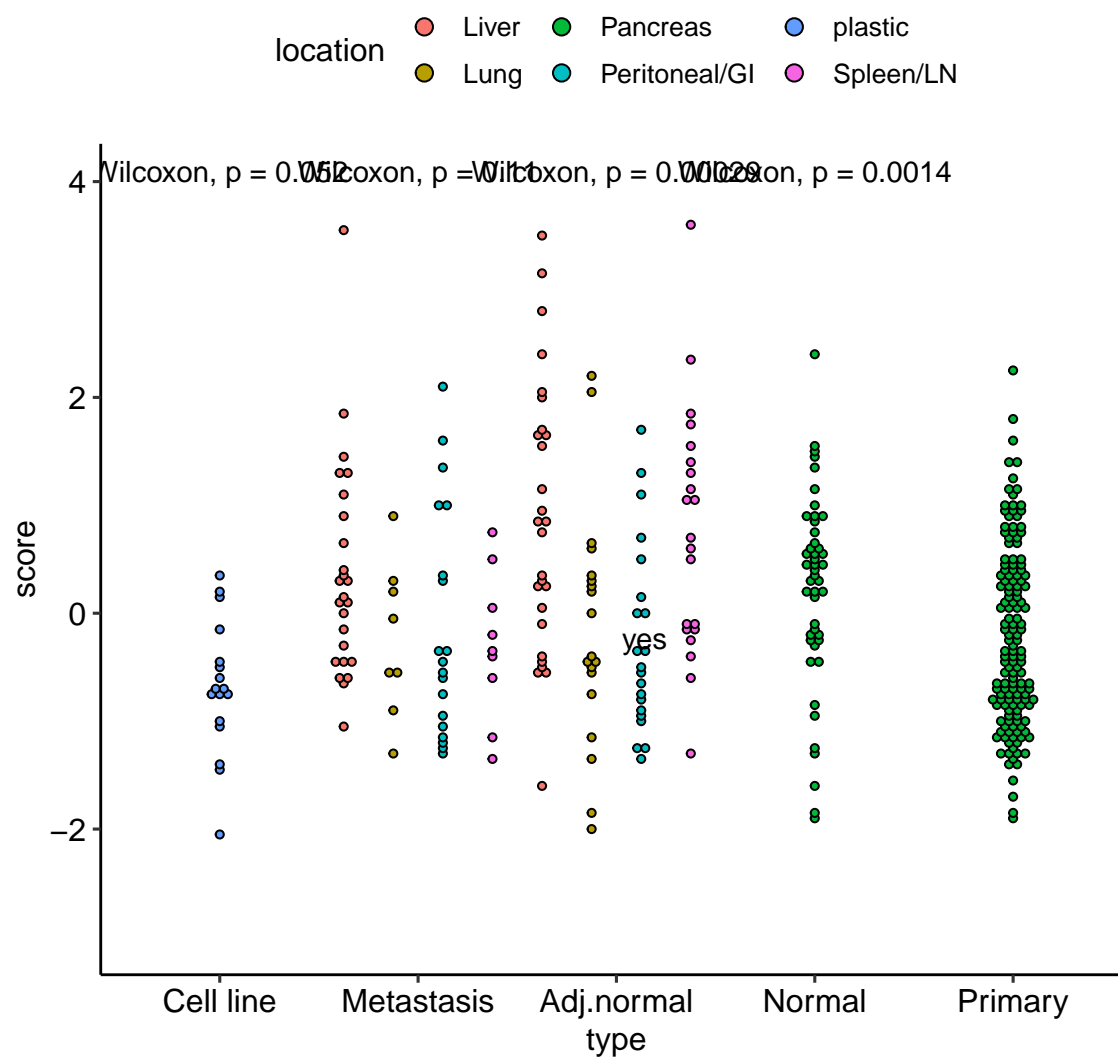




# Moffitt.F1\_Liver.top100 score



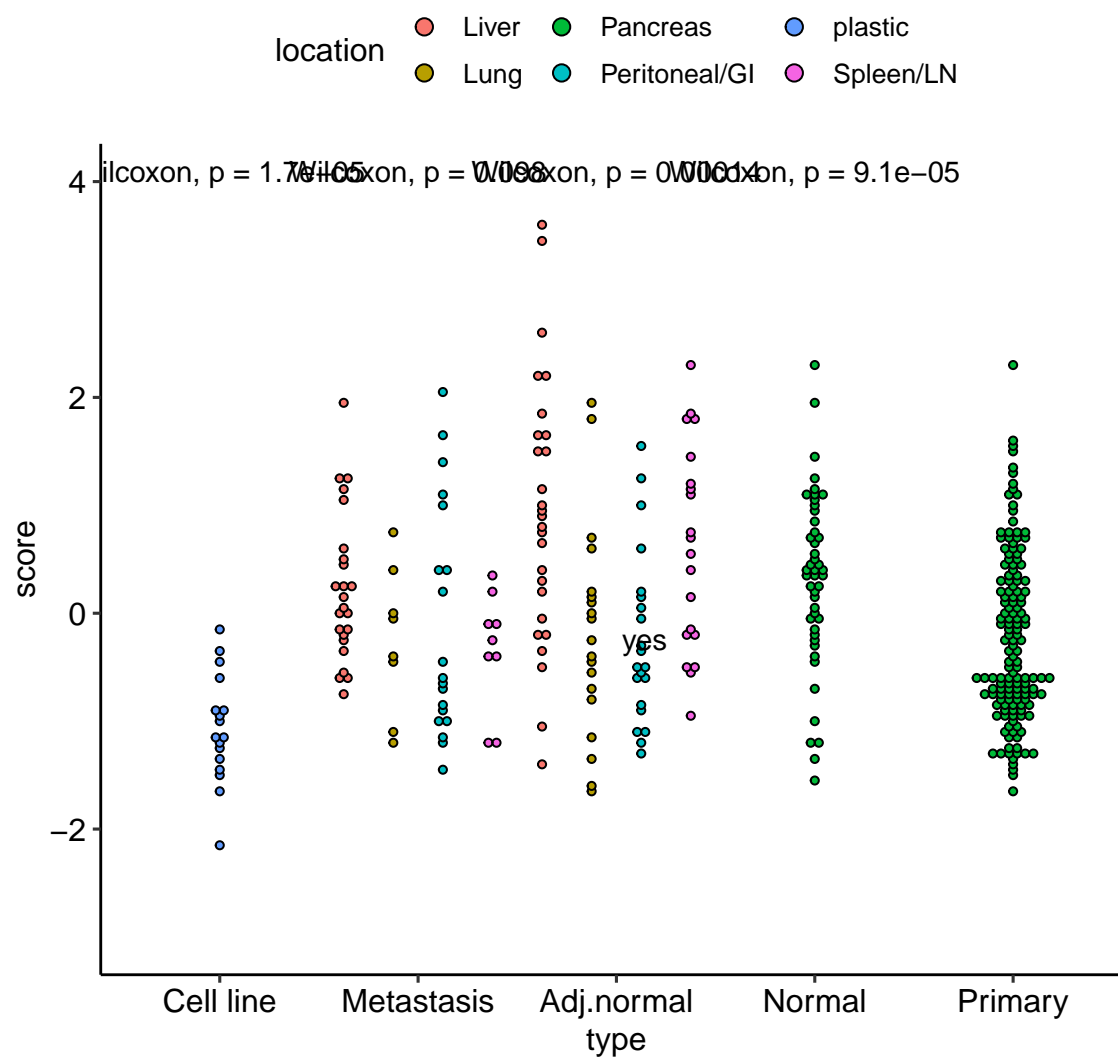
## Moffitt.F2.top25 score



## Warning: Removed 2 rows containing non-finite values (stat\_bindot).

## Warning: Removed 2 rows containing non-finite values (stat\_compare\_means).

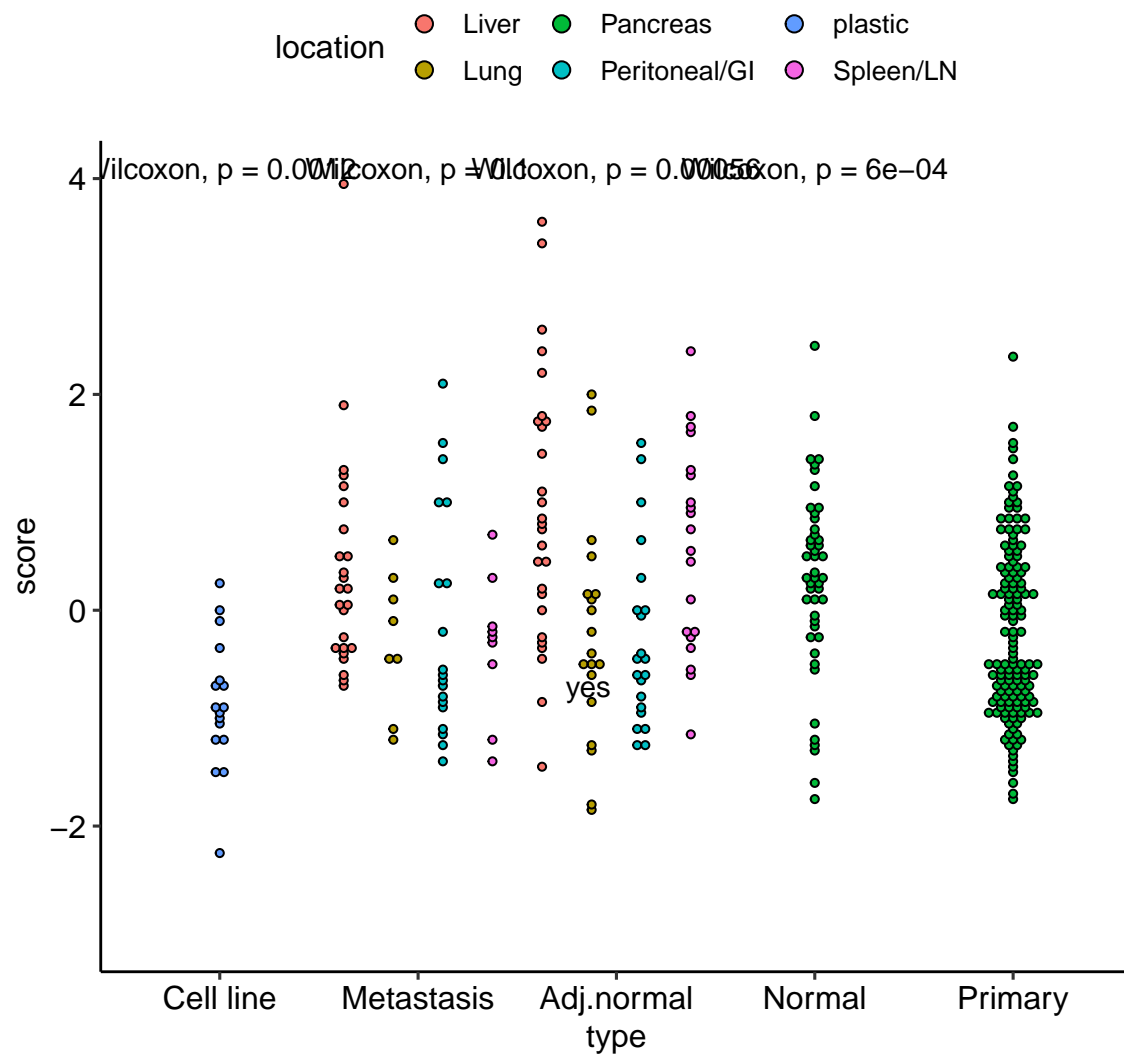
## Moffitt.F2.top250 score



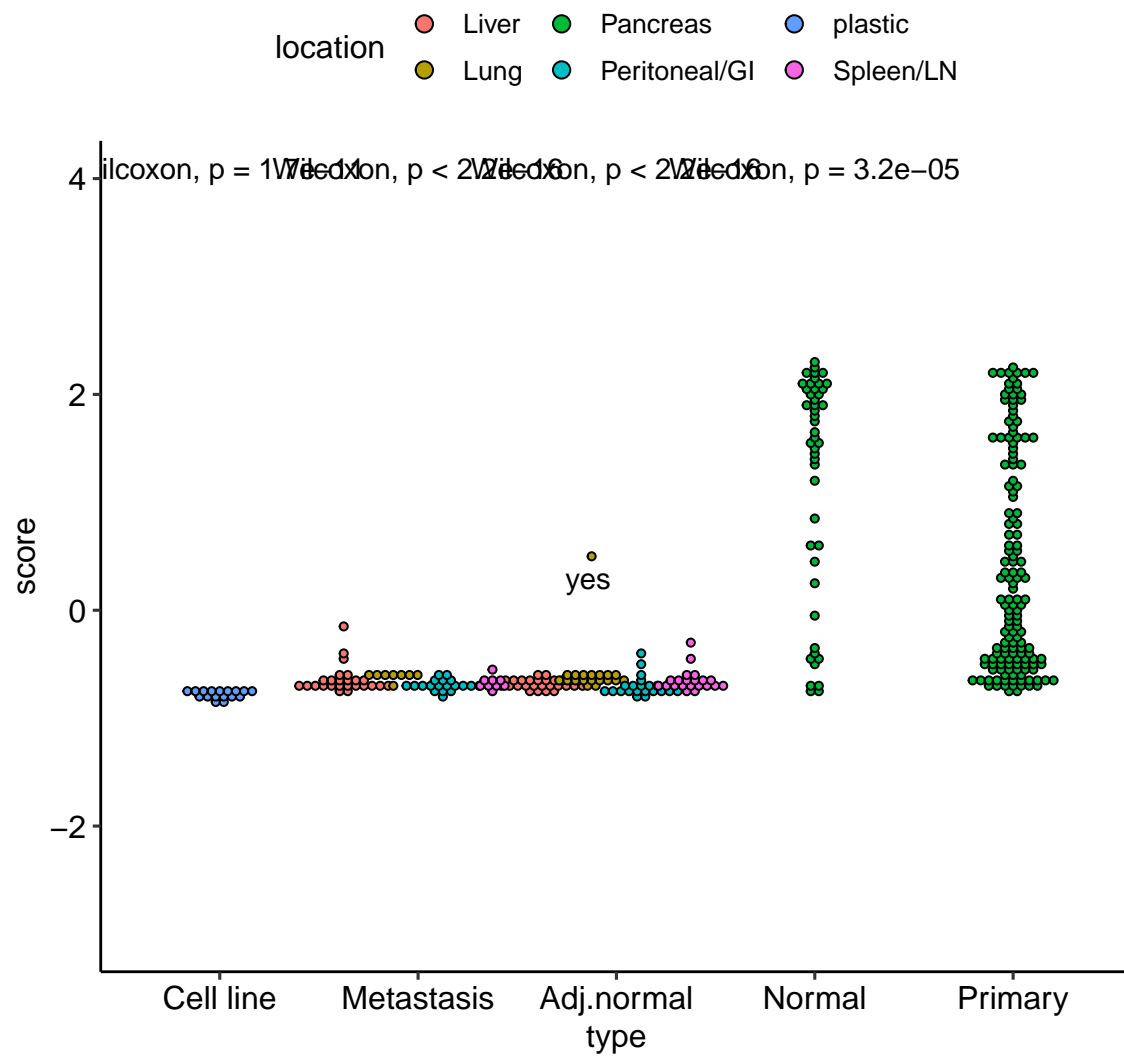
## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

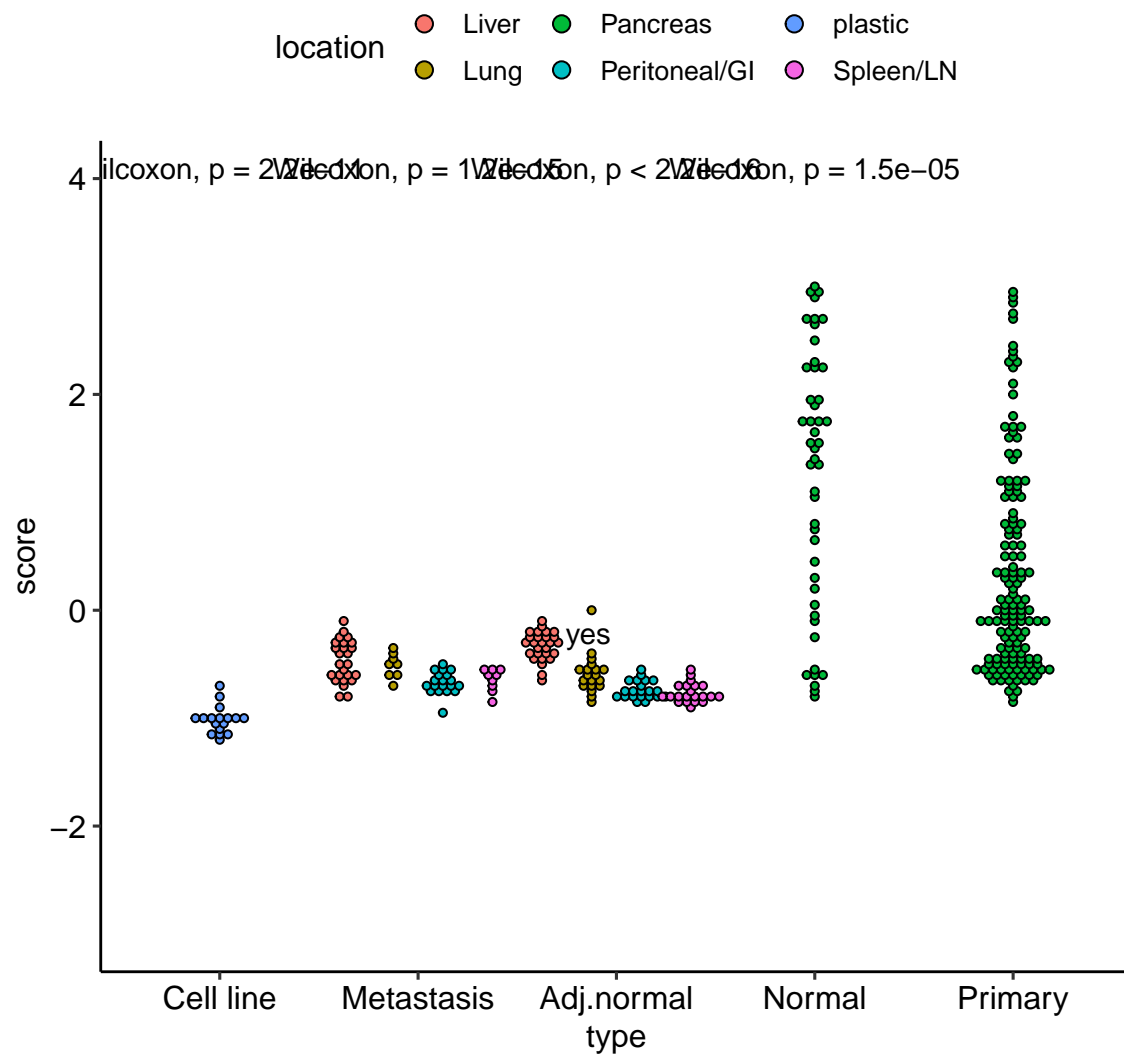
# Moffitt.F2.top100 score



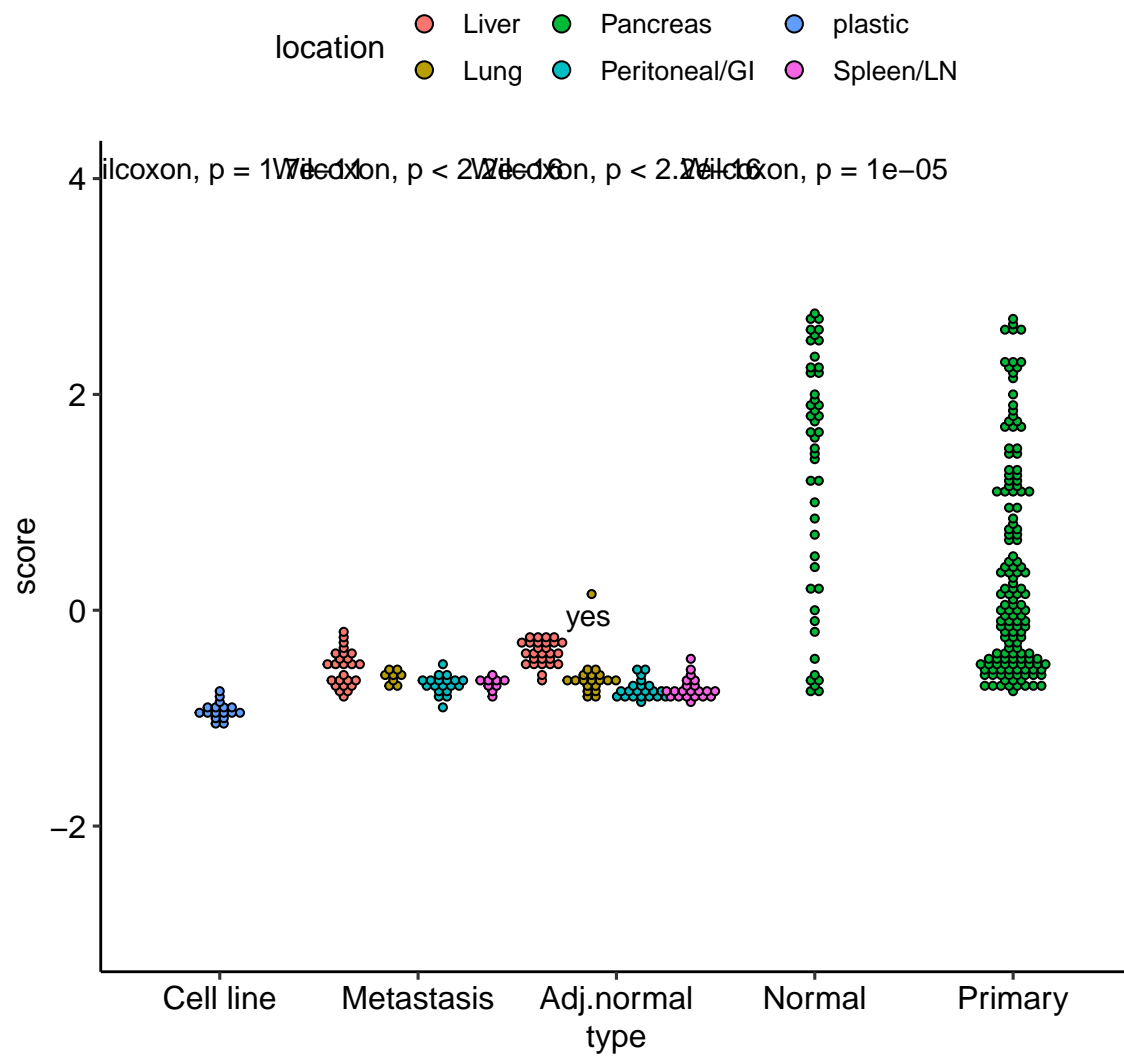
# Moffitt.F3\_Exocrine.top25 score



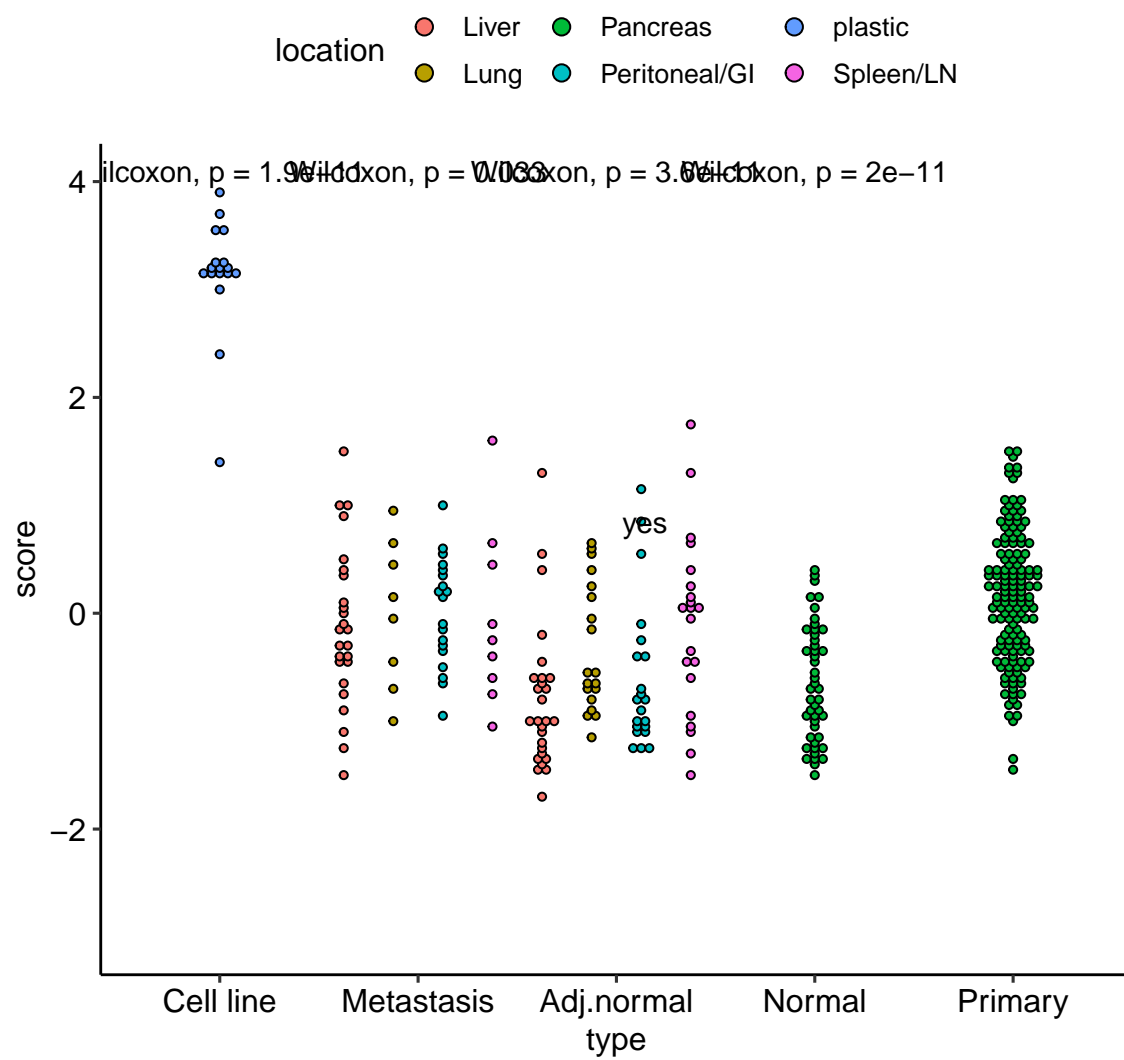
# Moffitt.F3\_Exocrine.top250 score



# Moffitt.F3\_Exocrine.top100 score

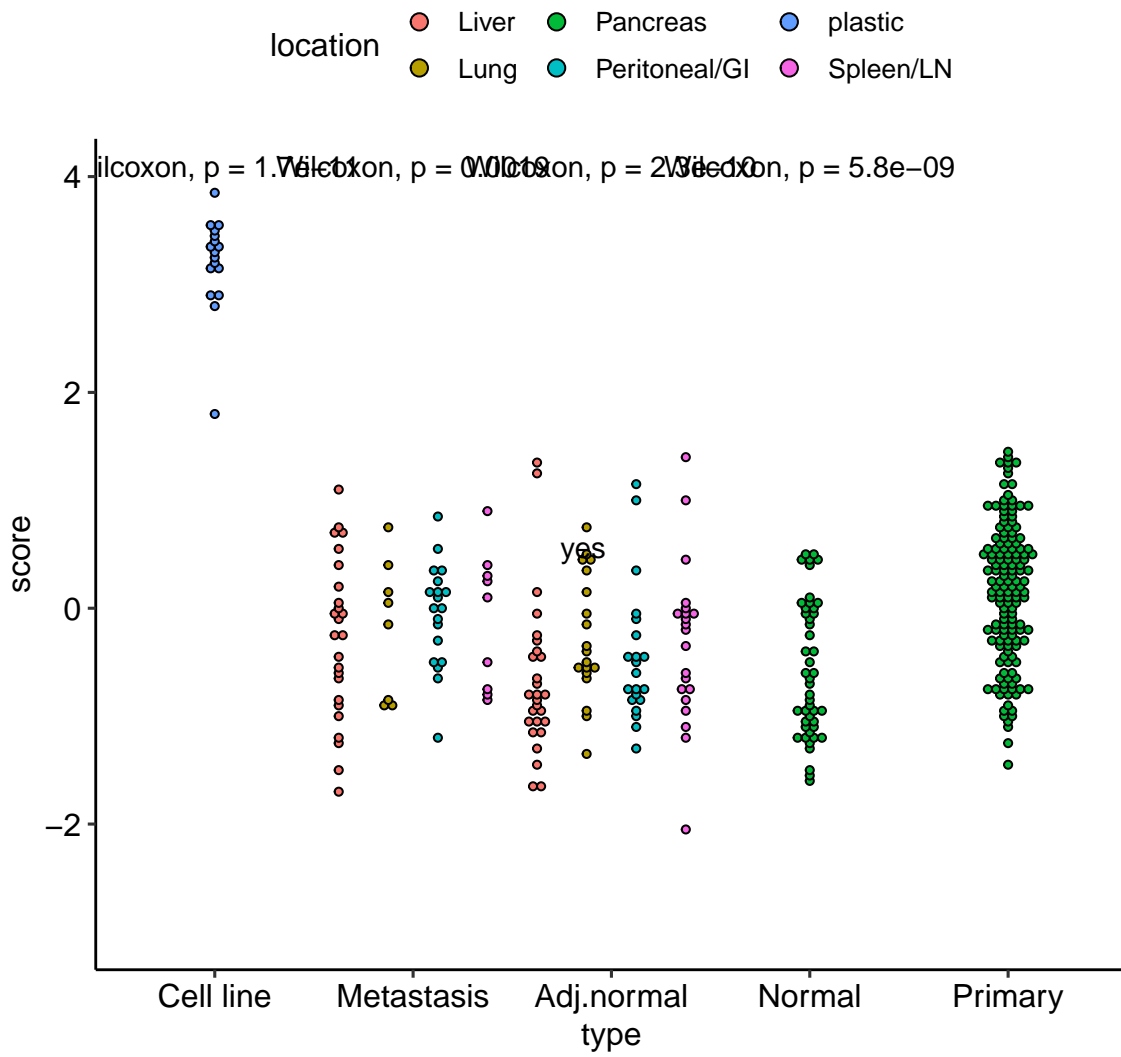


# Moffitt.F4\_CellCycle.top25 score

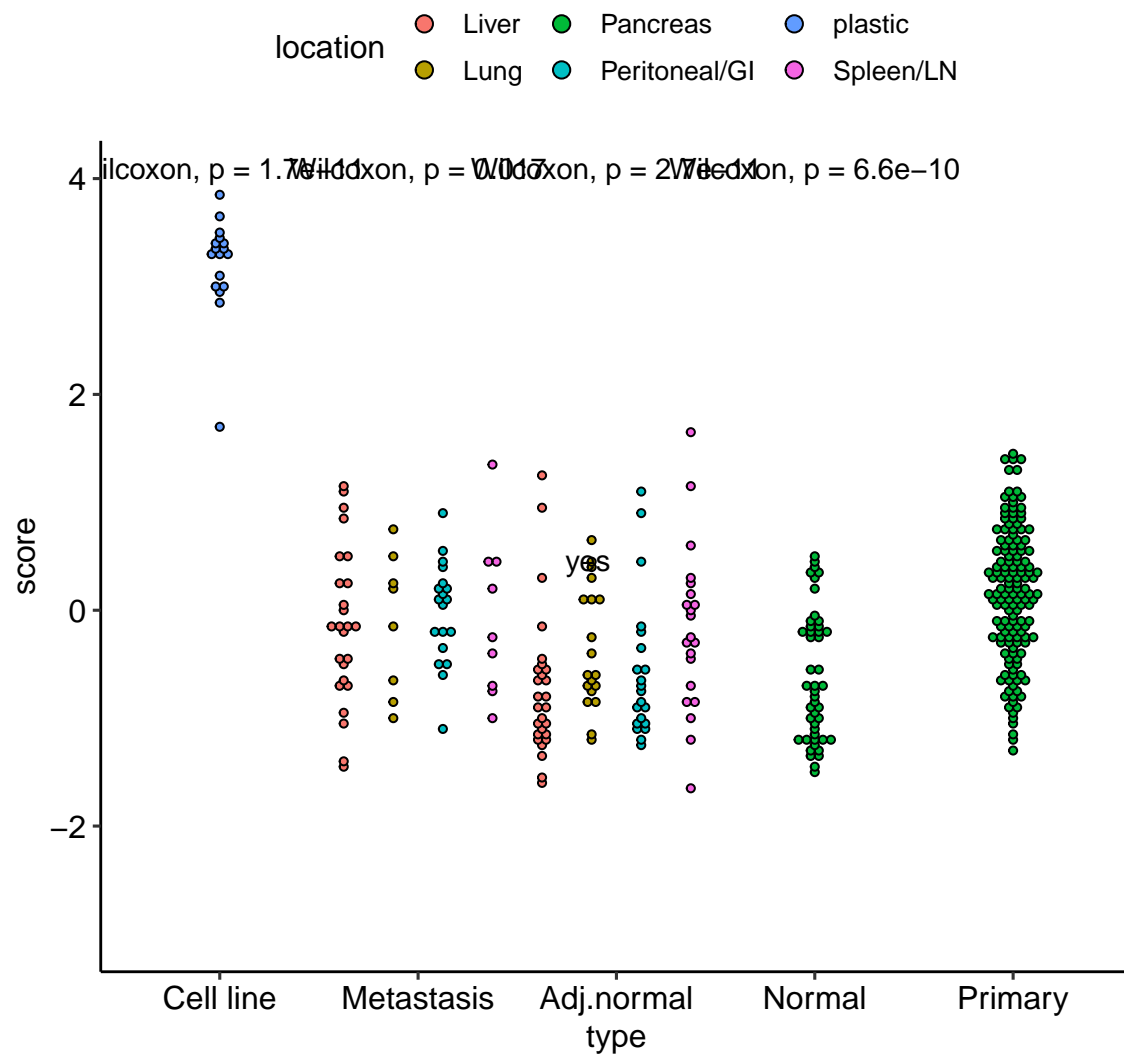




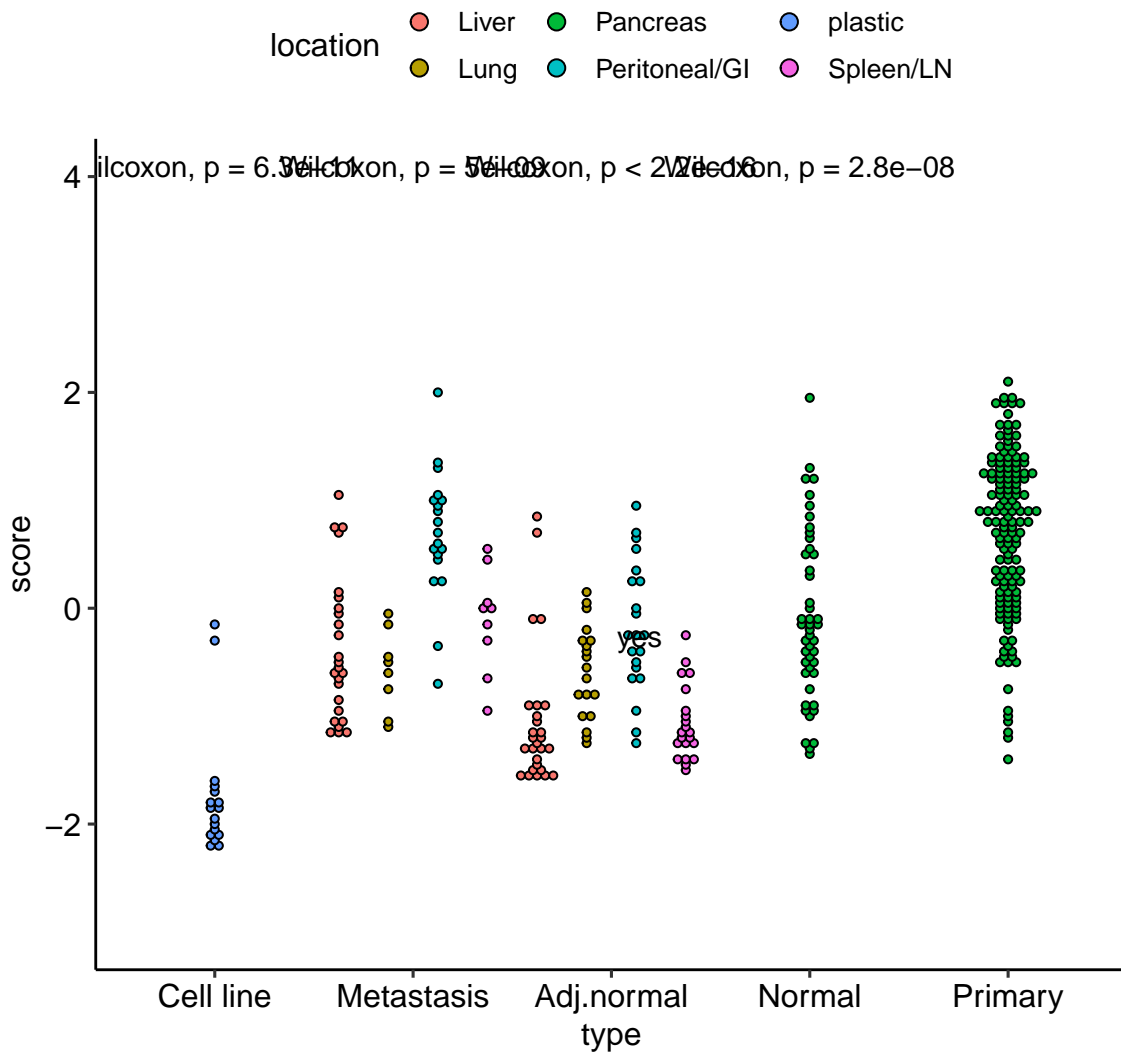
# Moffitt.F4\_CellCycle.top250 score



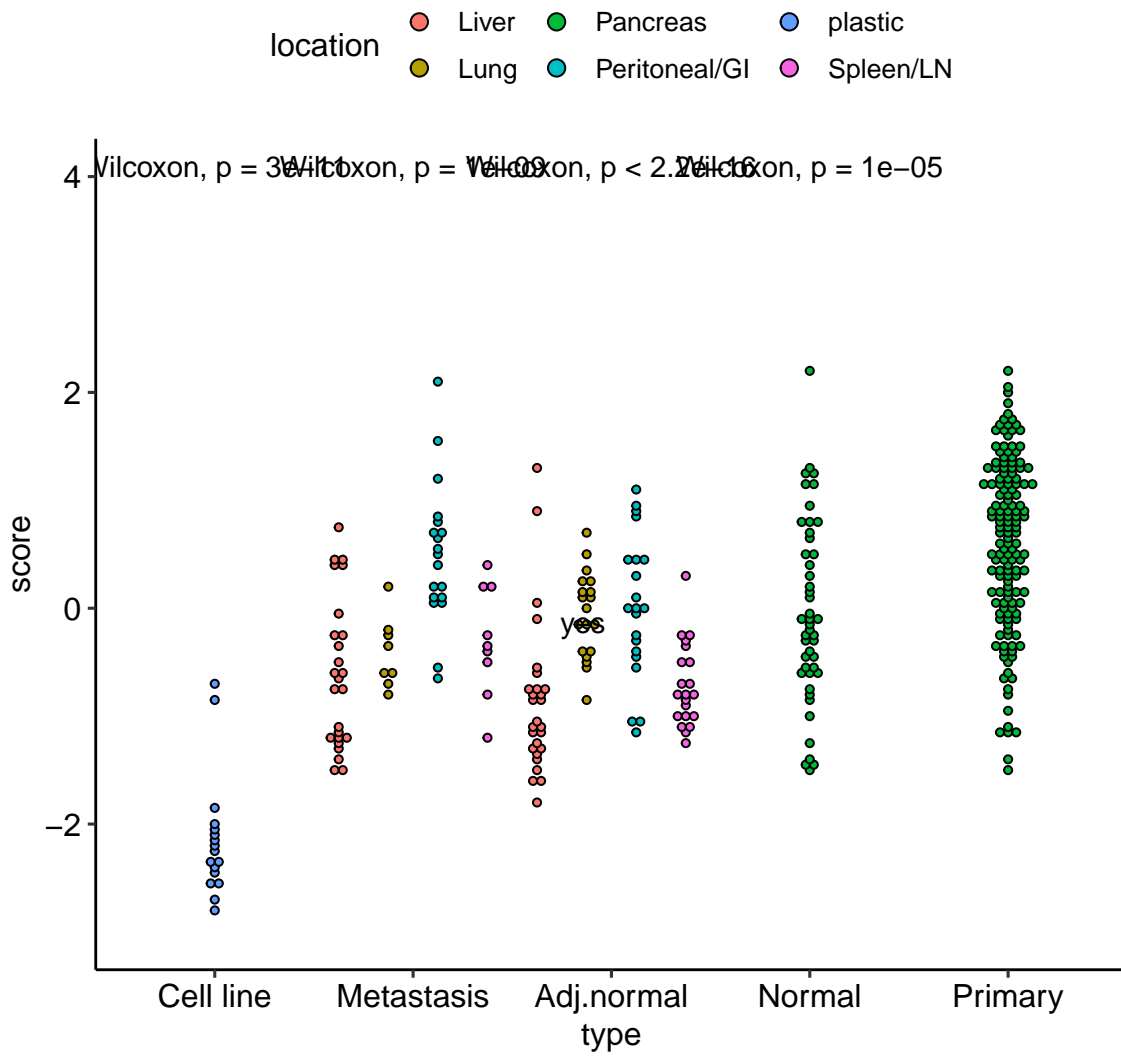
# Moffitt.F4\_CellCycle.top100 score



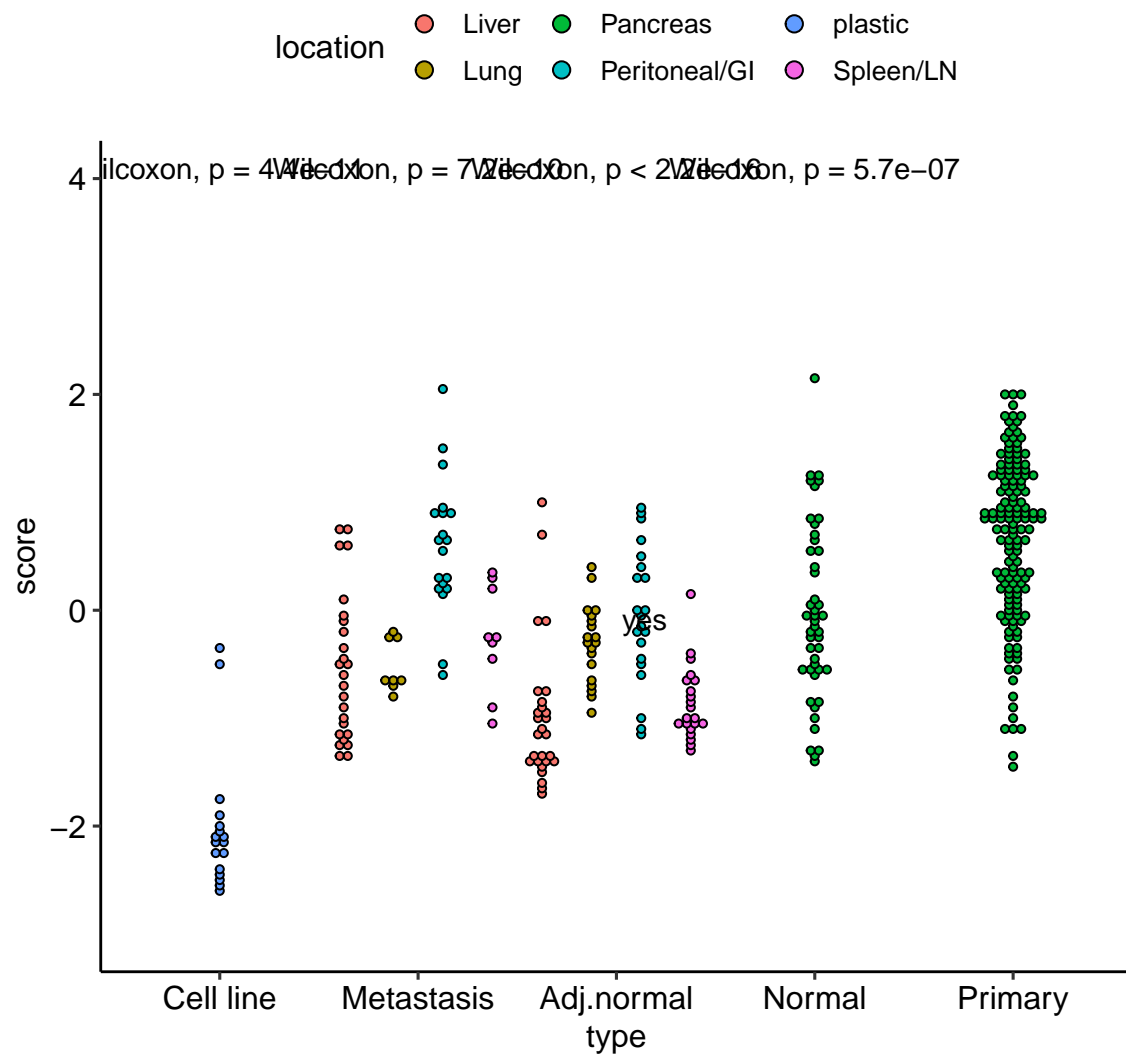
# Moffitt.F5\_ActivatedStroma.top25 score



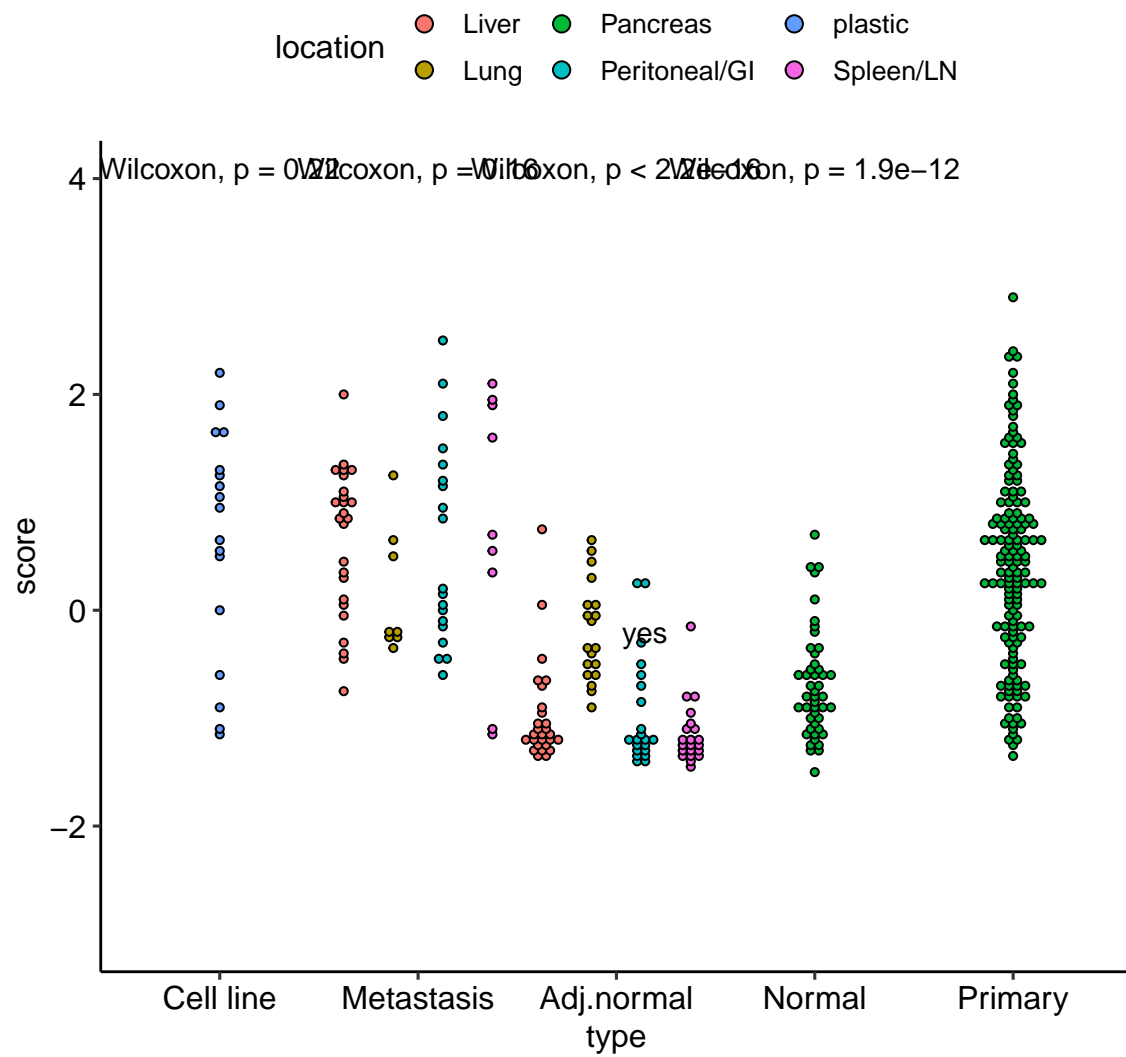
# Moffitt.F5\_ActivatedStroma.top250 score



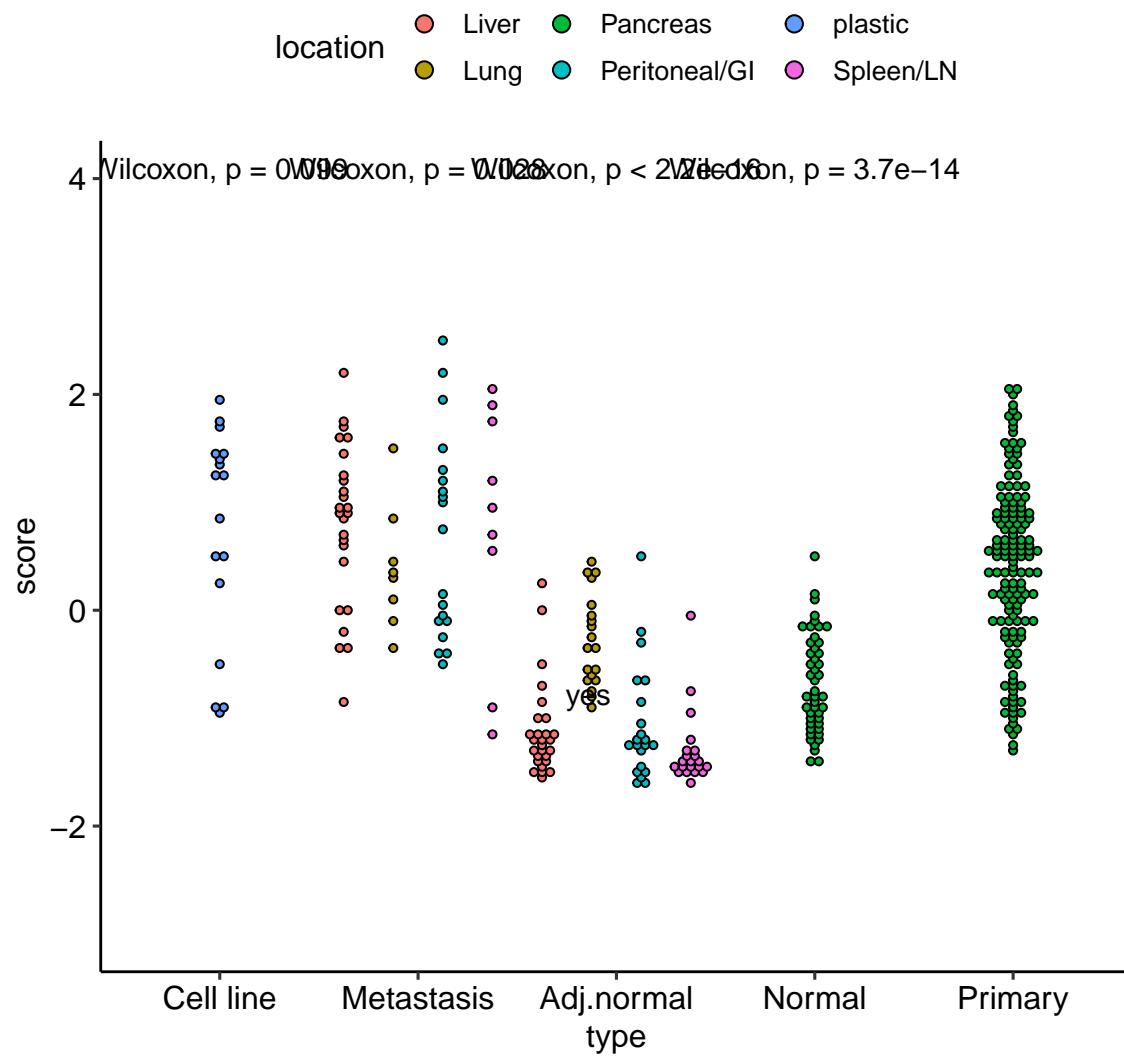
# Moffitt.F5\_ActivatedStroma.top100 score



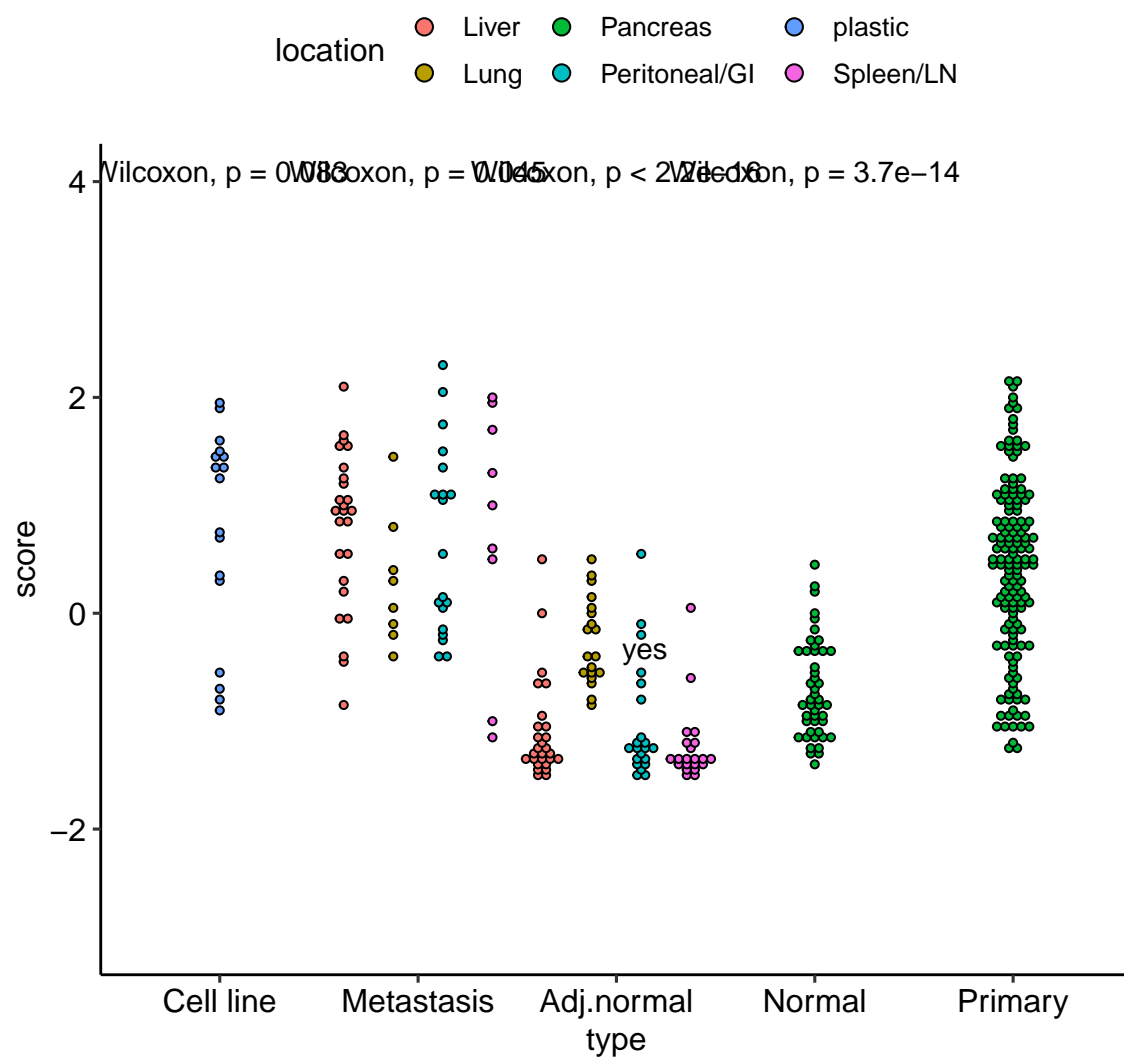
# Moffitt.F6\_BasalLike.top25 score



# Moffitt.F6\_BasalLike.top250 score

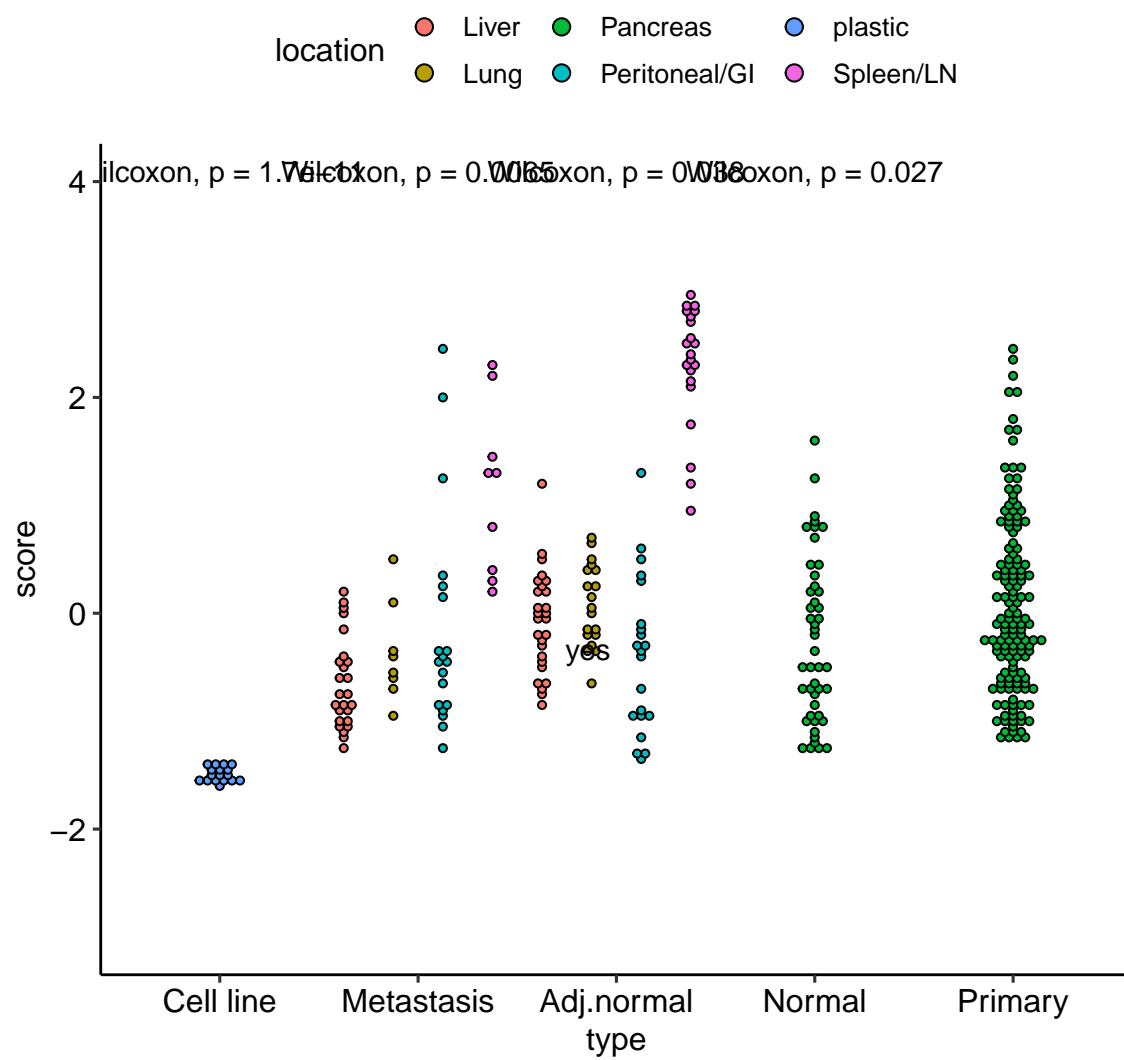


# Moffitt.F6\_BasalLike.top100 score

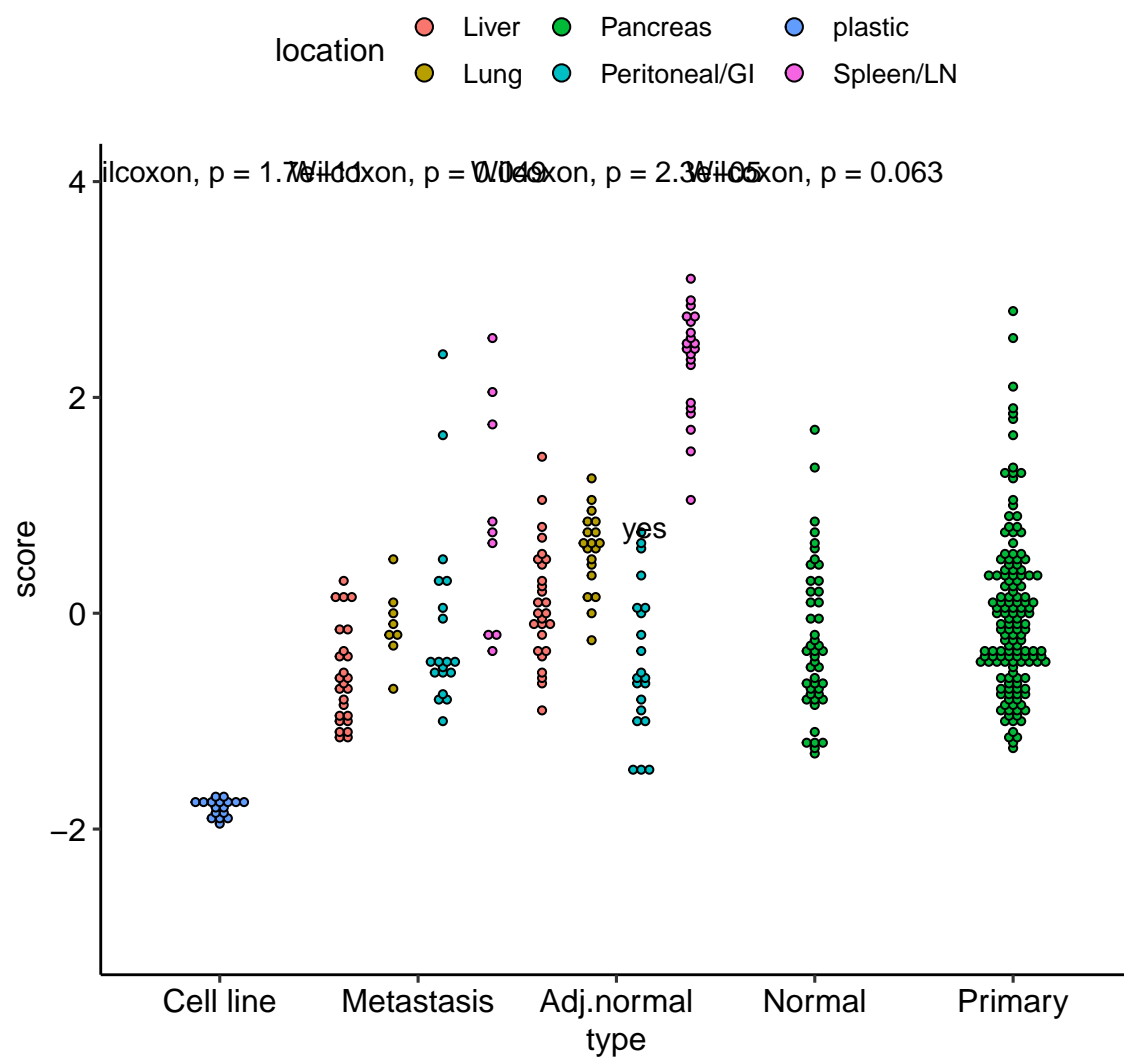




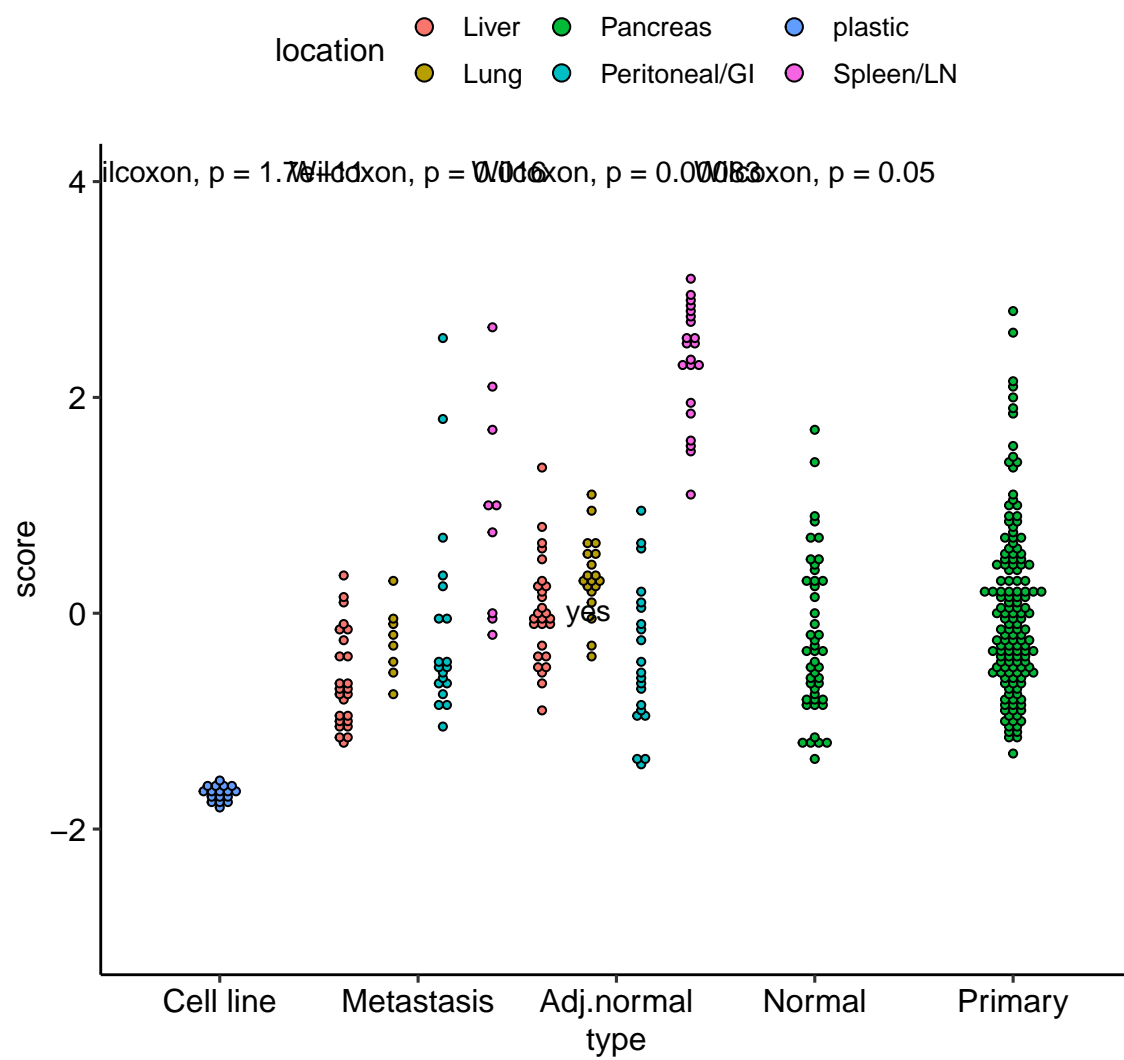
# Moffitt.F7\_Immune.top25 score



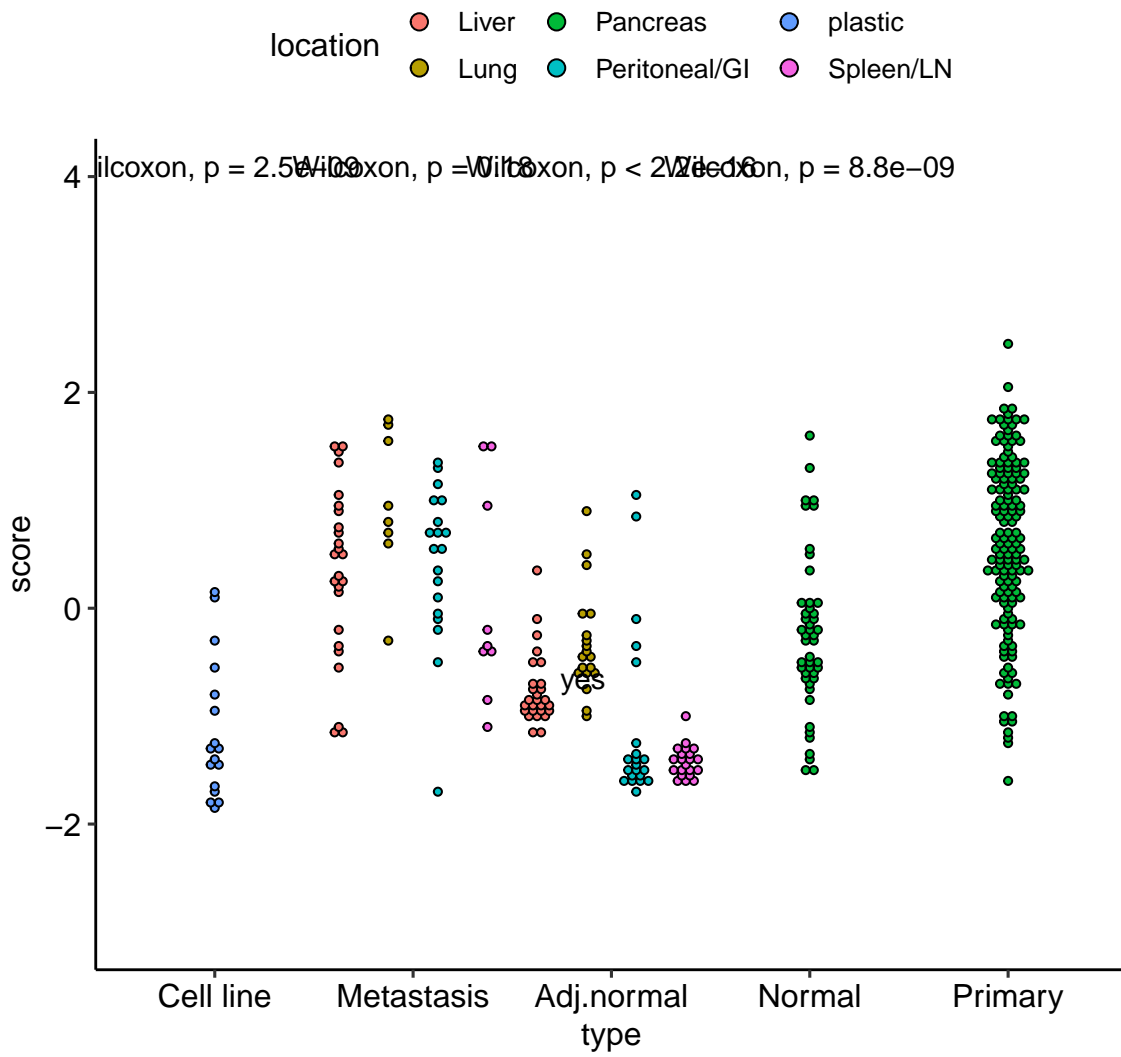
# Moffitt.F7\_Immune.top250 score



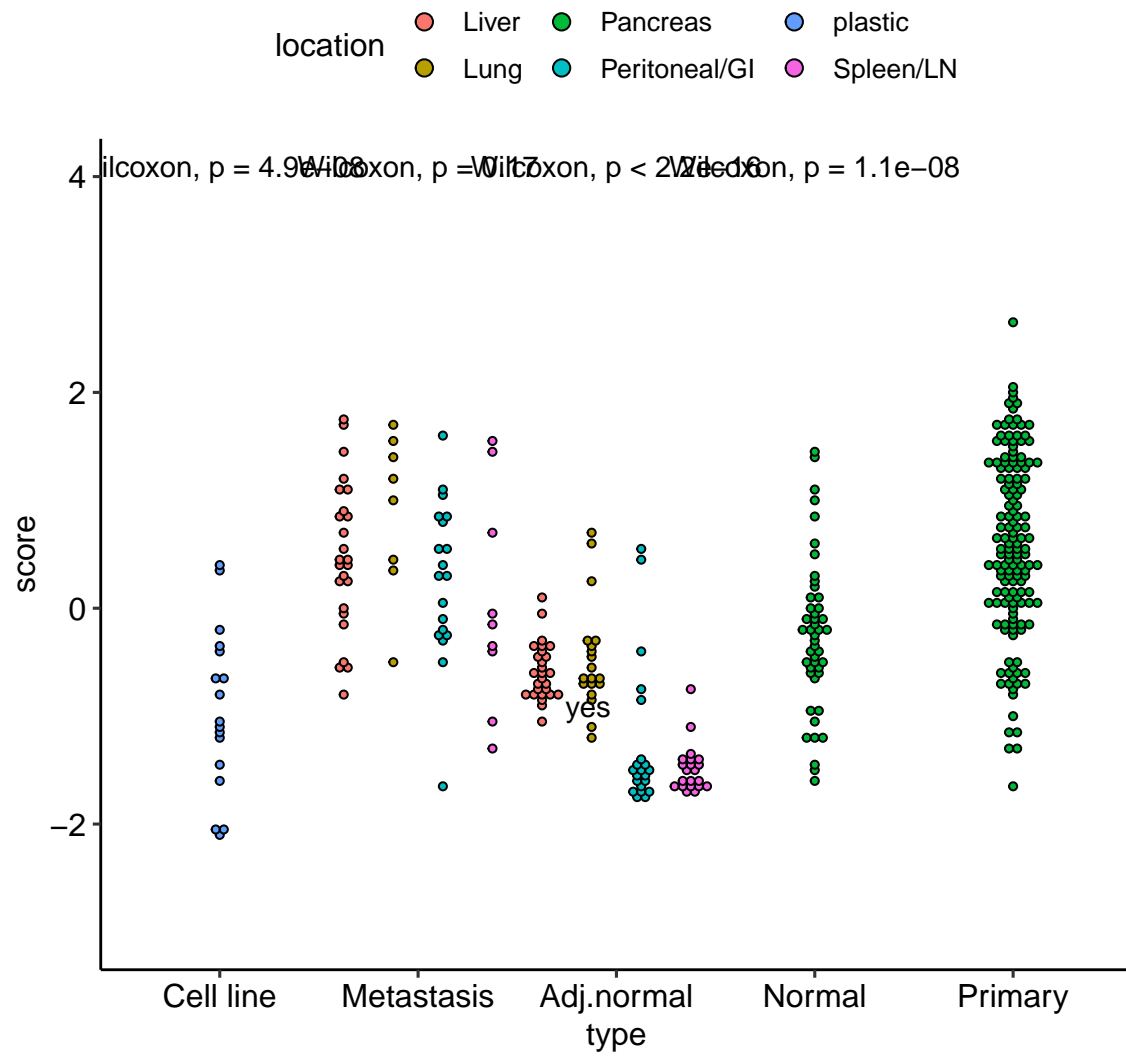
# Moffitt.F7\_Immune.top100 score



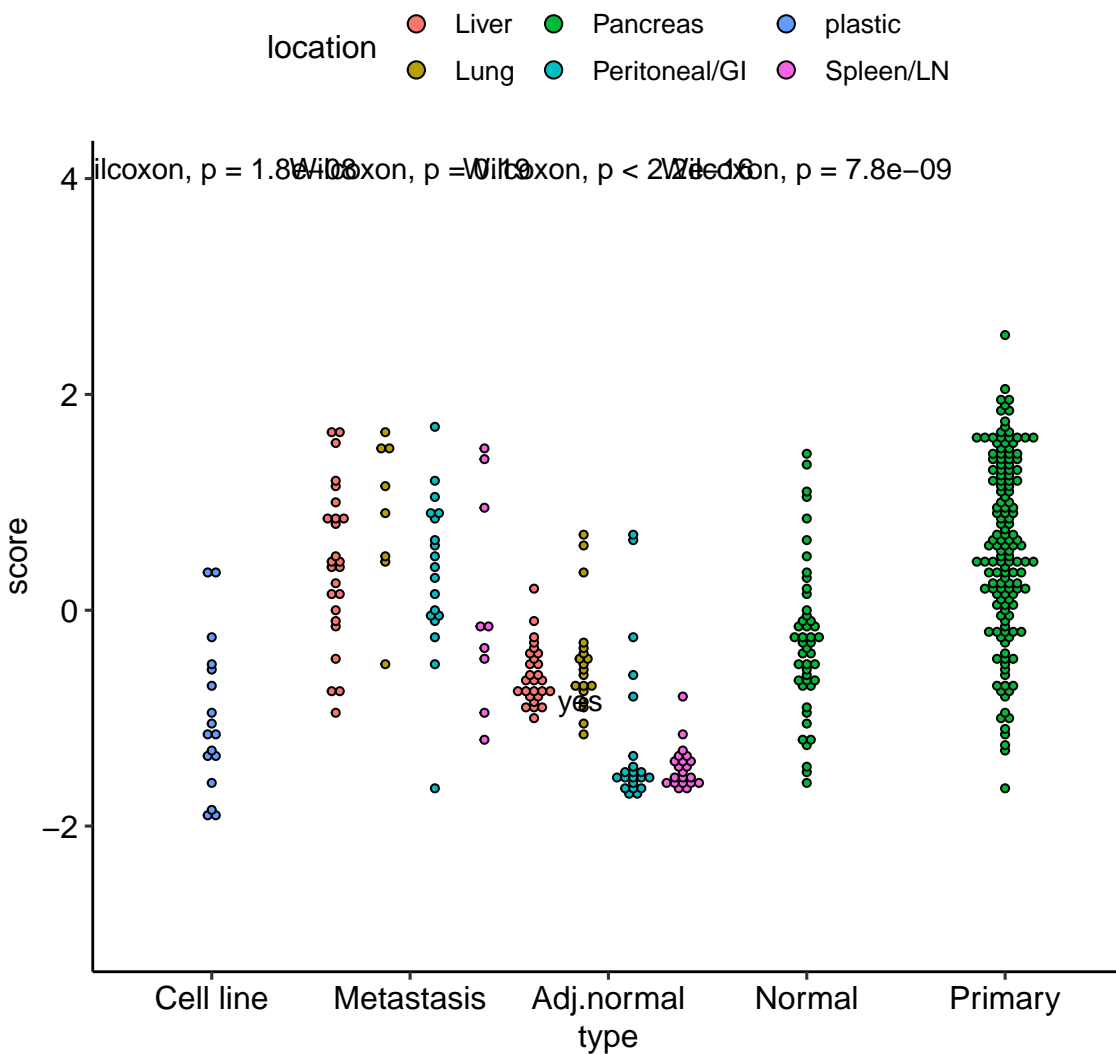
# Moffitt.F8\_Classical.top25 score



# Moffitt.F8\_Classical.top250 score



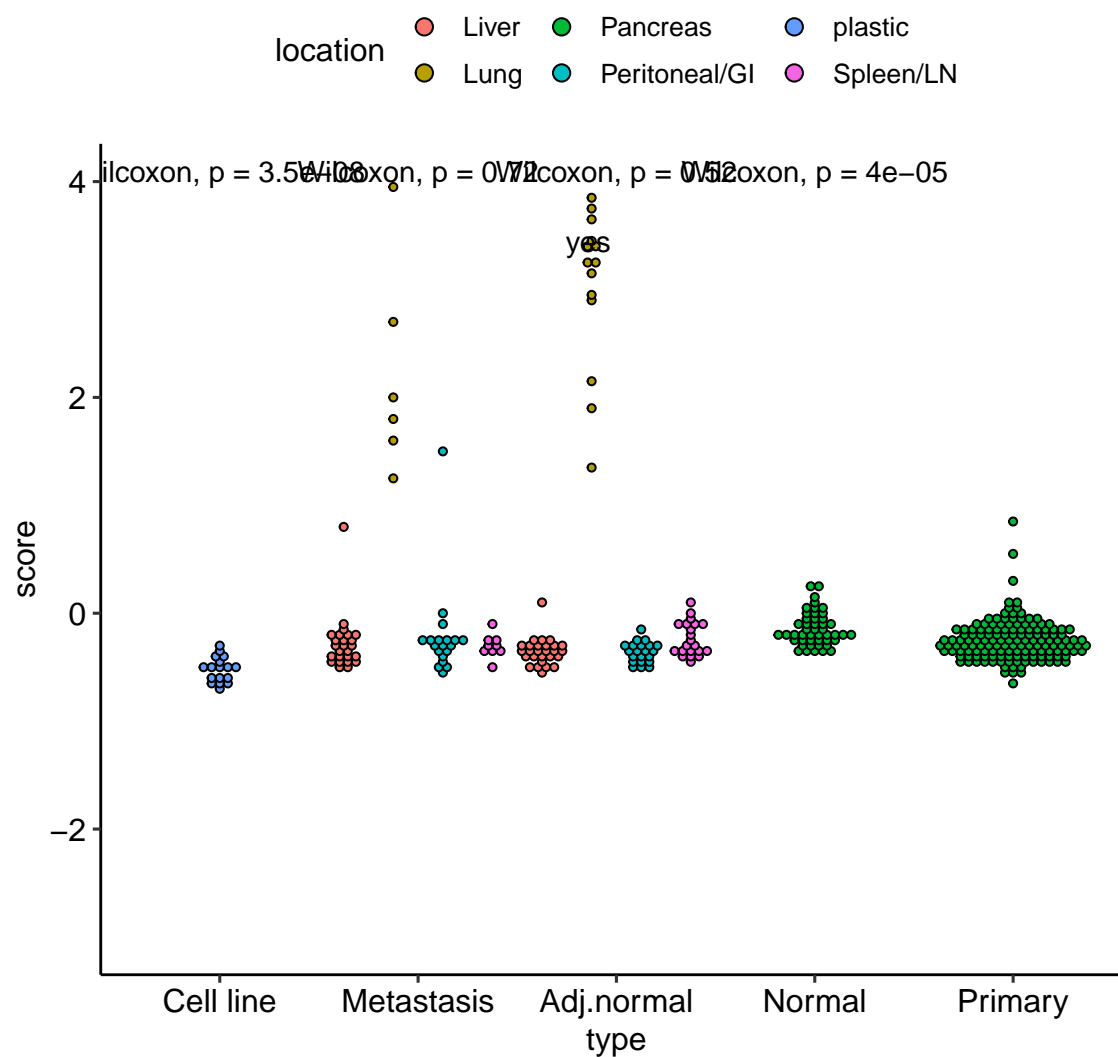
## Moffitt.F8\_Classical.top100 score



## Warning: Removed 7 rows containing non-finite values (stat\_bindot).

## Warning: Removed 7 rows containing non-finite values (stat\_compare\_means).

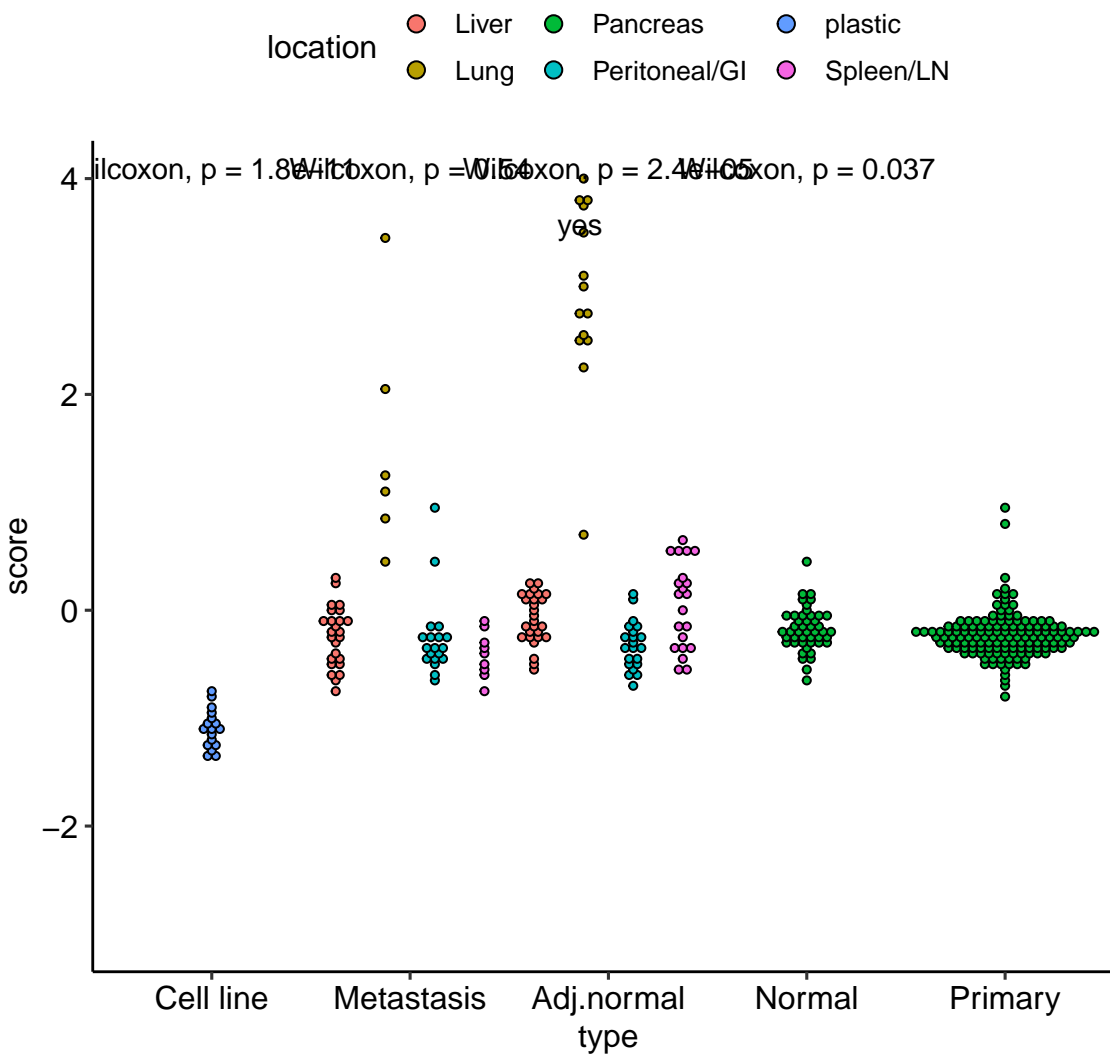
## Moffitt.F9\_Lung.top25 score



## Warning: Removed 7 rows containing non-finite values (stat\_bindot).

## Warning: Removed 7 rows containing non-finite values (stat\_compare\_means).

## Moffitt.F9\_Lung.top250 score

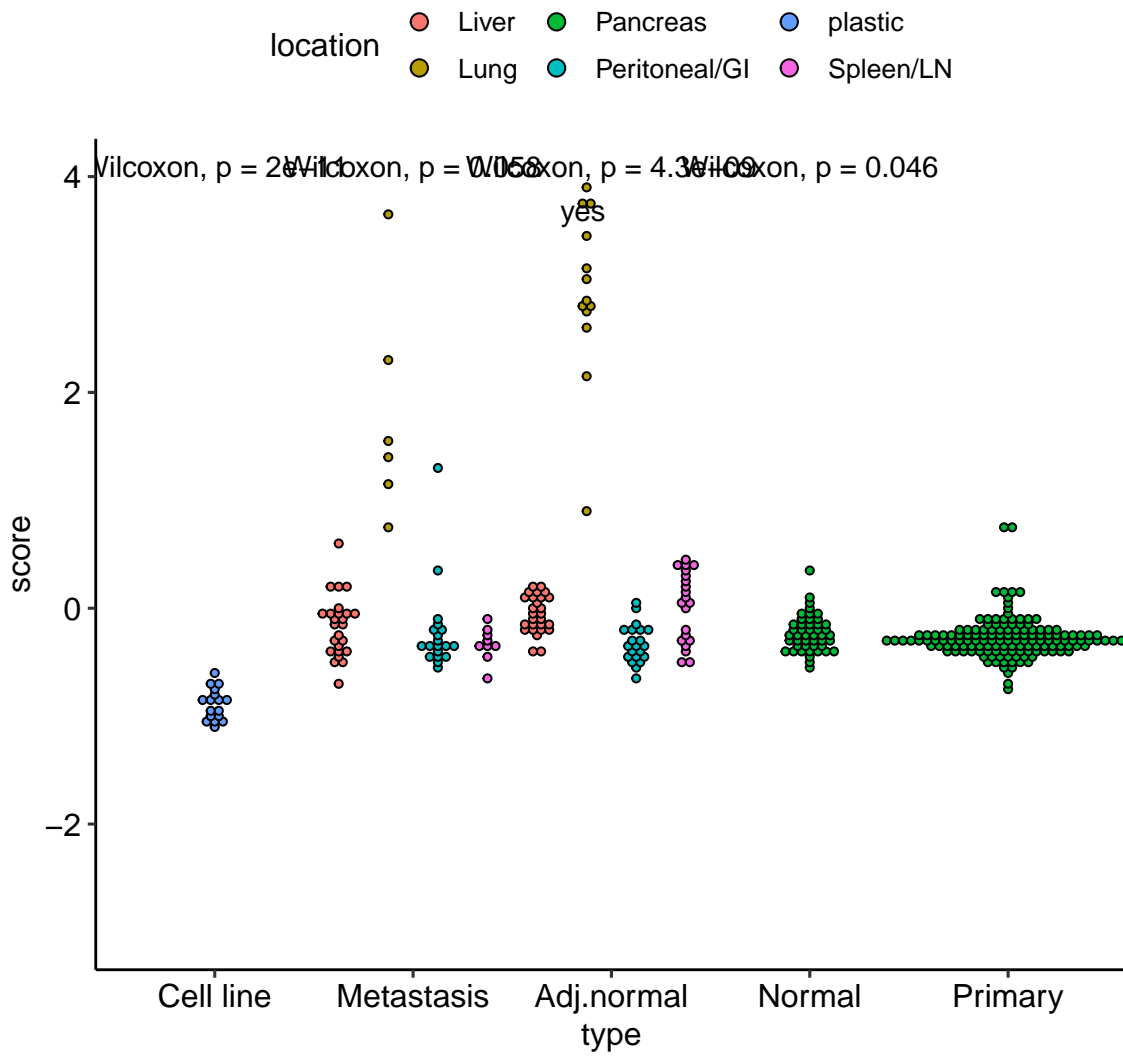


## Warning: Removed 8 rows containing non-finite values (stat\_bindot).

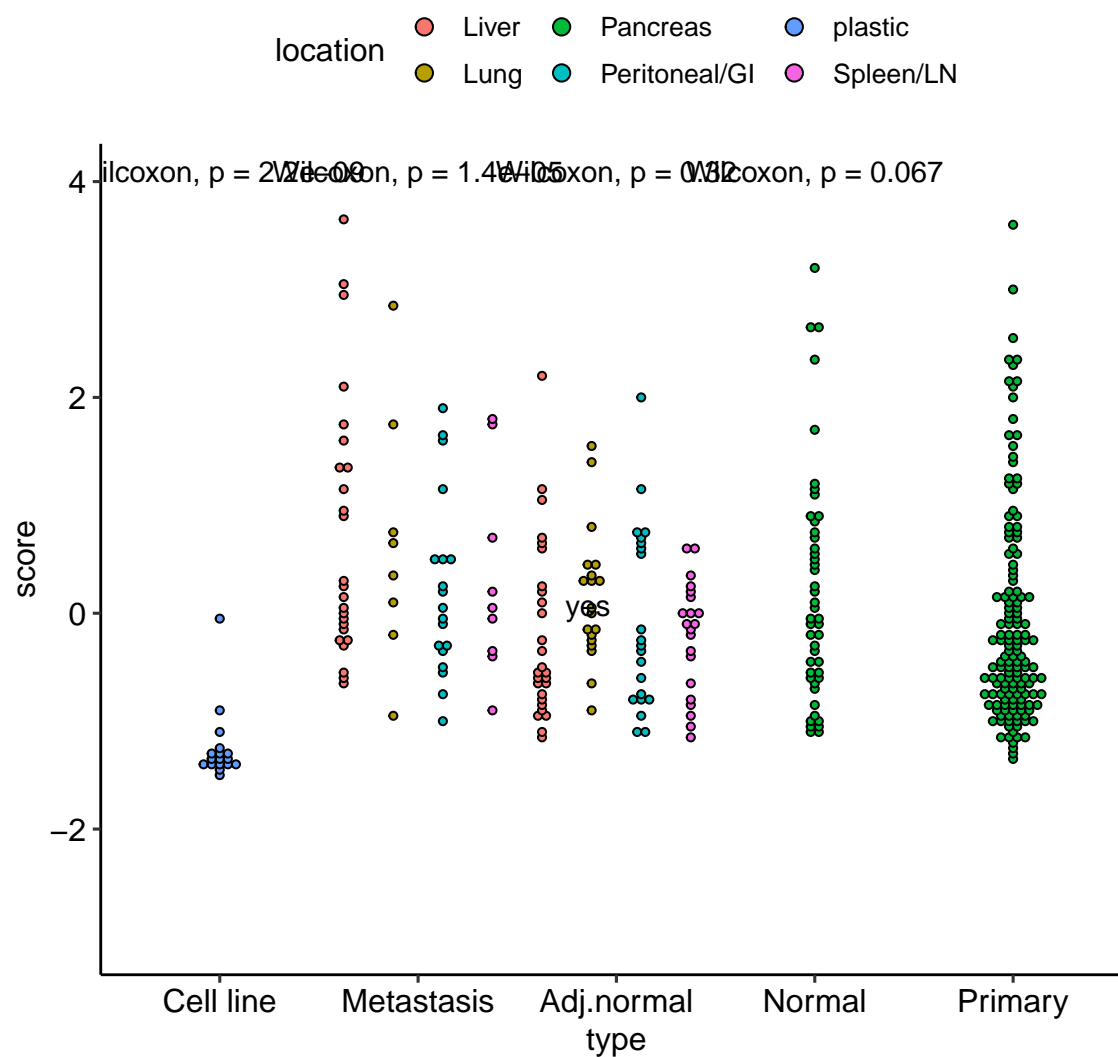
## Warning: Removed 8 rows containing non-finite values (stat\_compare\_means).



# Moffitt.F9\_Lung.top100 score



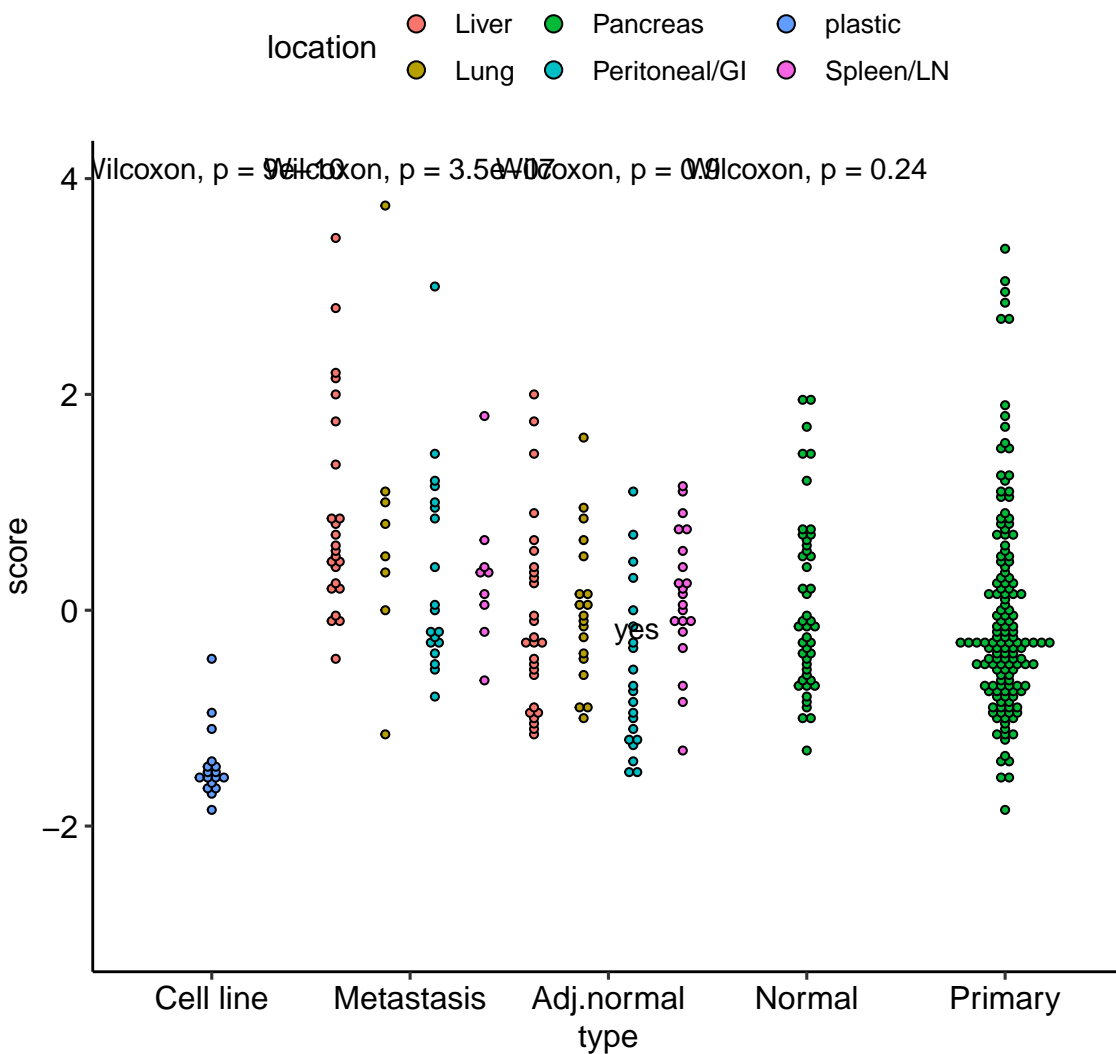
## Moffitt.F10.top25 score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

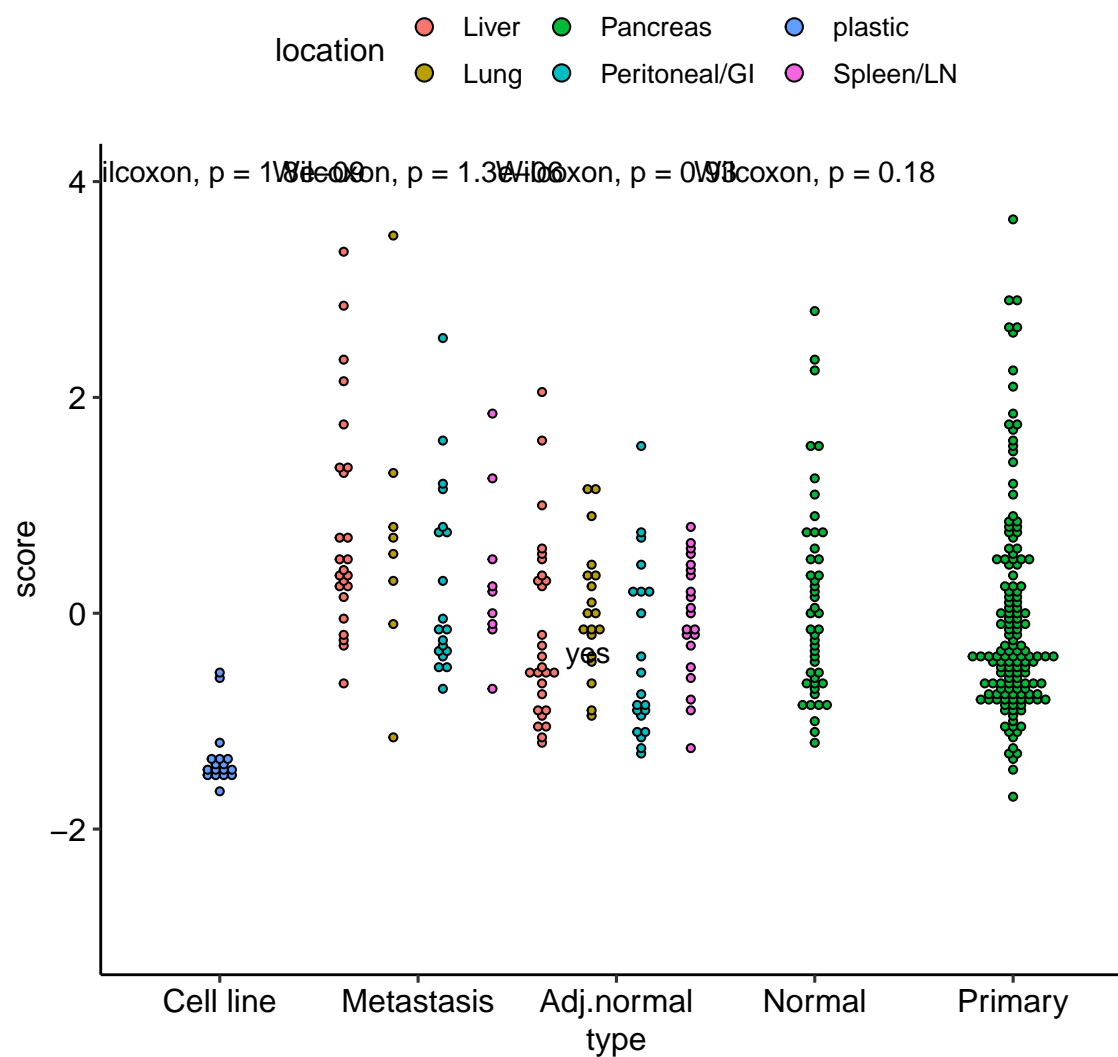
## Moffitt.F10.top250 score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

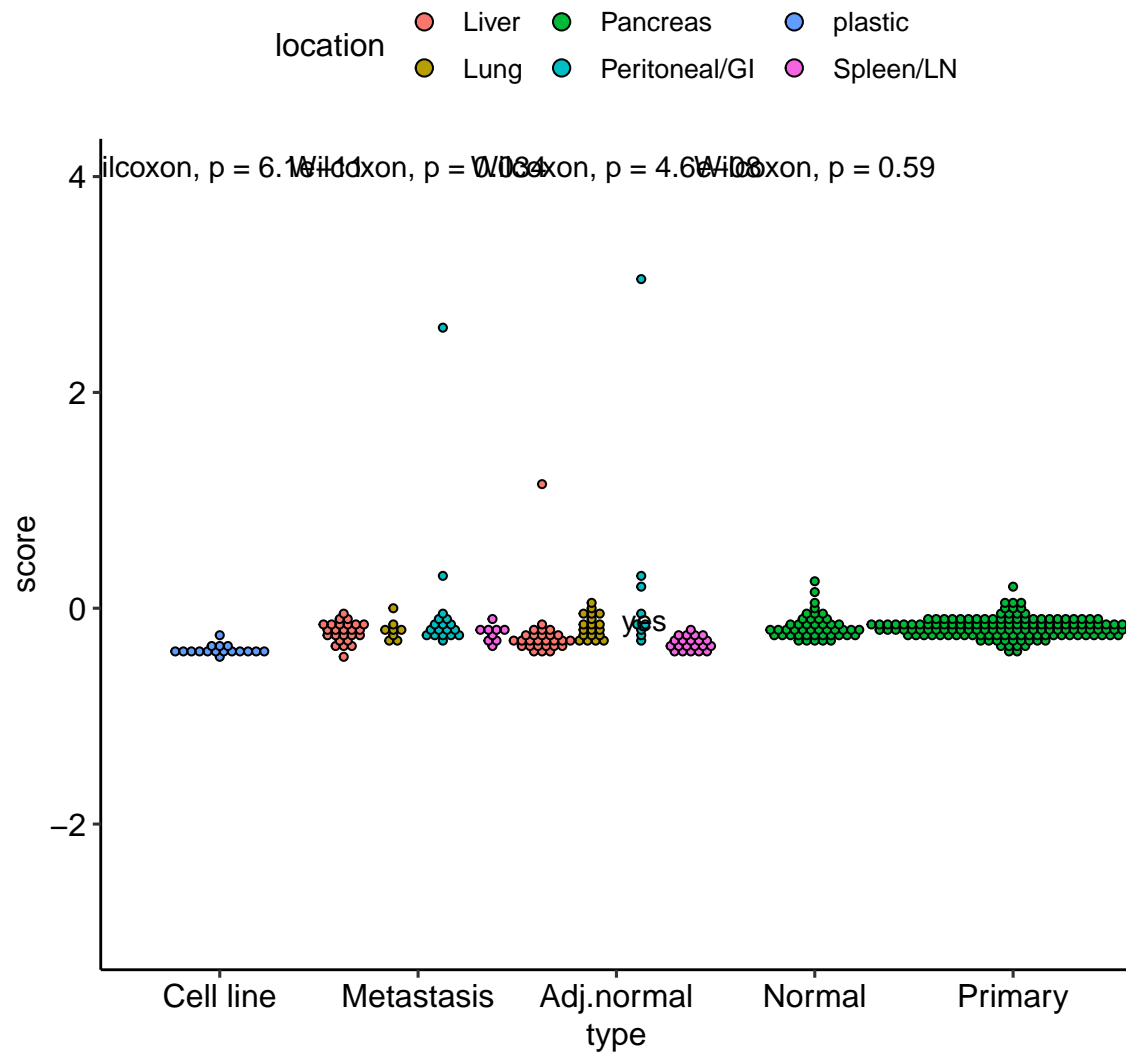
## Moffitt.F10.top100 score



## Warning: Removed 13 rows containing non-finite values (stat\_bindot).

## Warning: Removed 13 rows containing non-finite values (stat\_compare\_means).

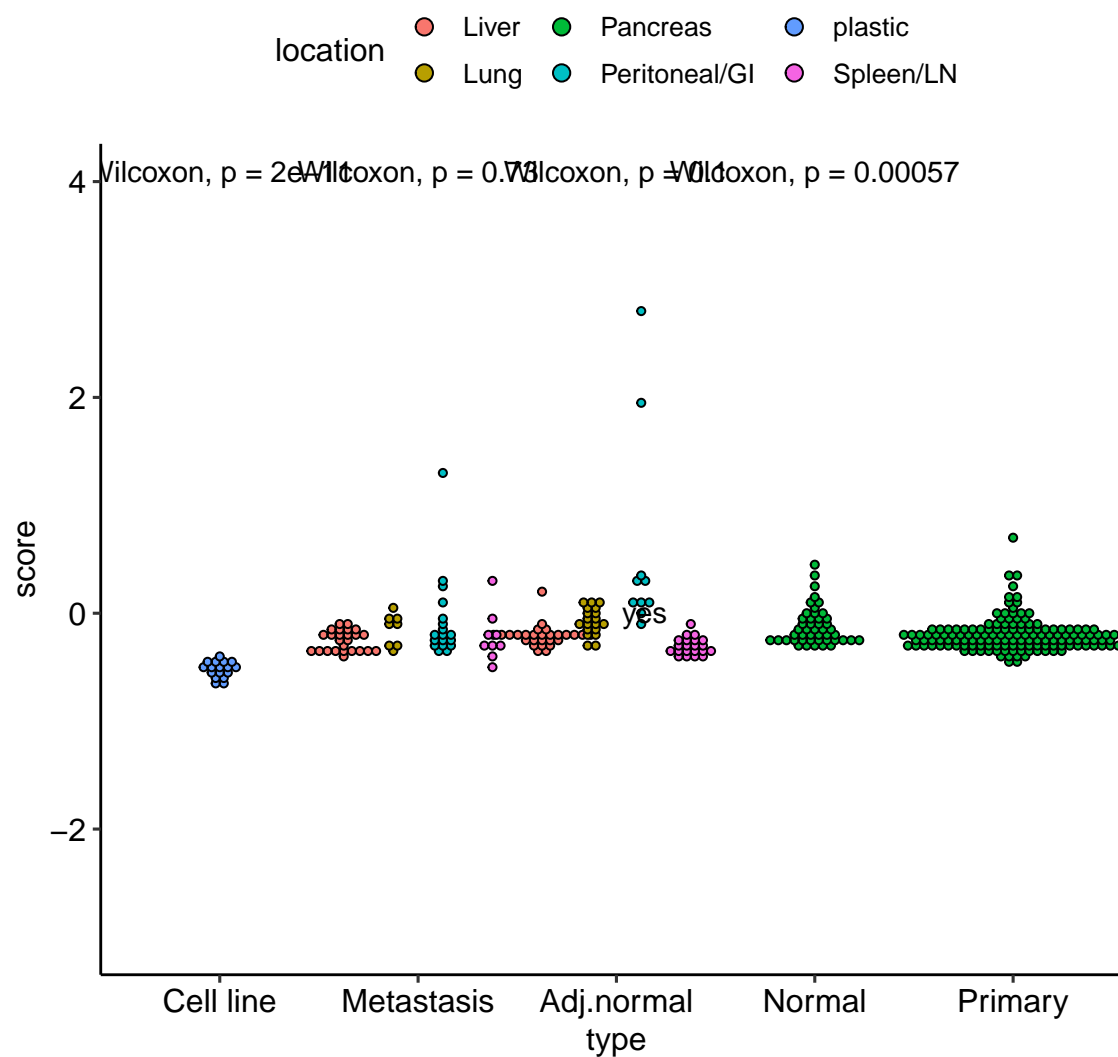
## Moffitt.F11\_Muscle.top25 score



## Warning: Removed 12 rows containing non-finite values (stat\_bindot).

## Warning: Removed 12 rows containing non-finite values (stat\_compare\_means).

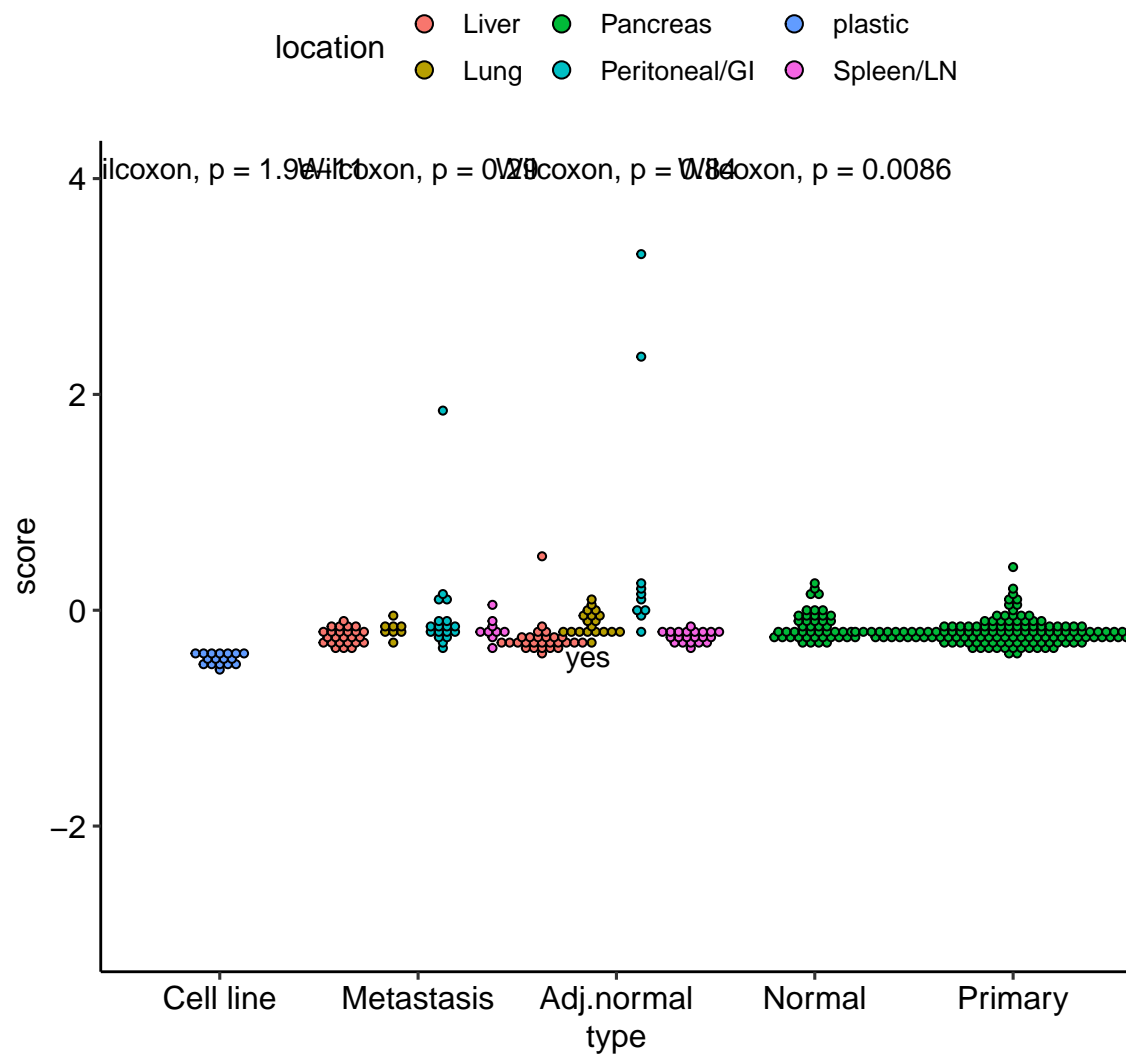
## Moffitt.F11\_Muscle.top250 score



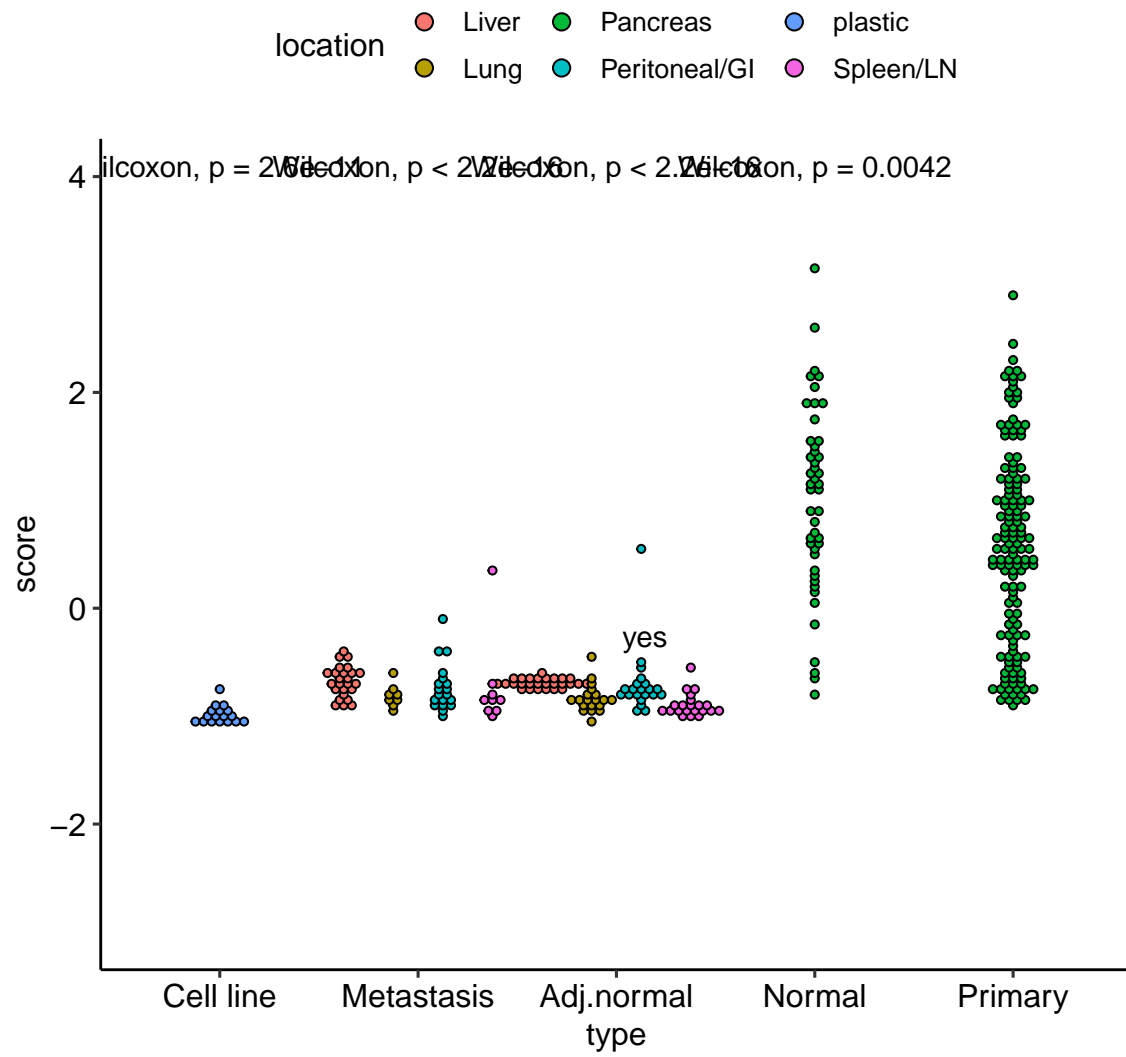
## Warning: Removed 12 rows containing non-finite values (stat\_bindot).

## Warning: Removed 12 rows containing non-finite values (stat\_compare\_means).

# Moffitt.F11\_Muscle.top100 score

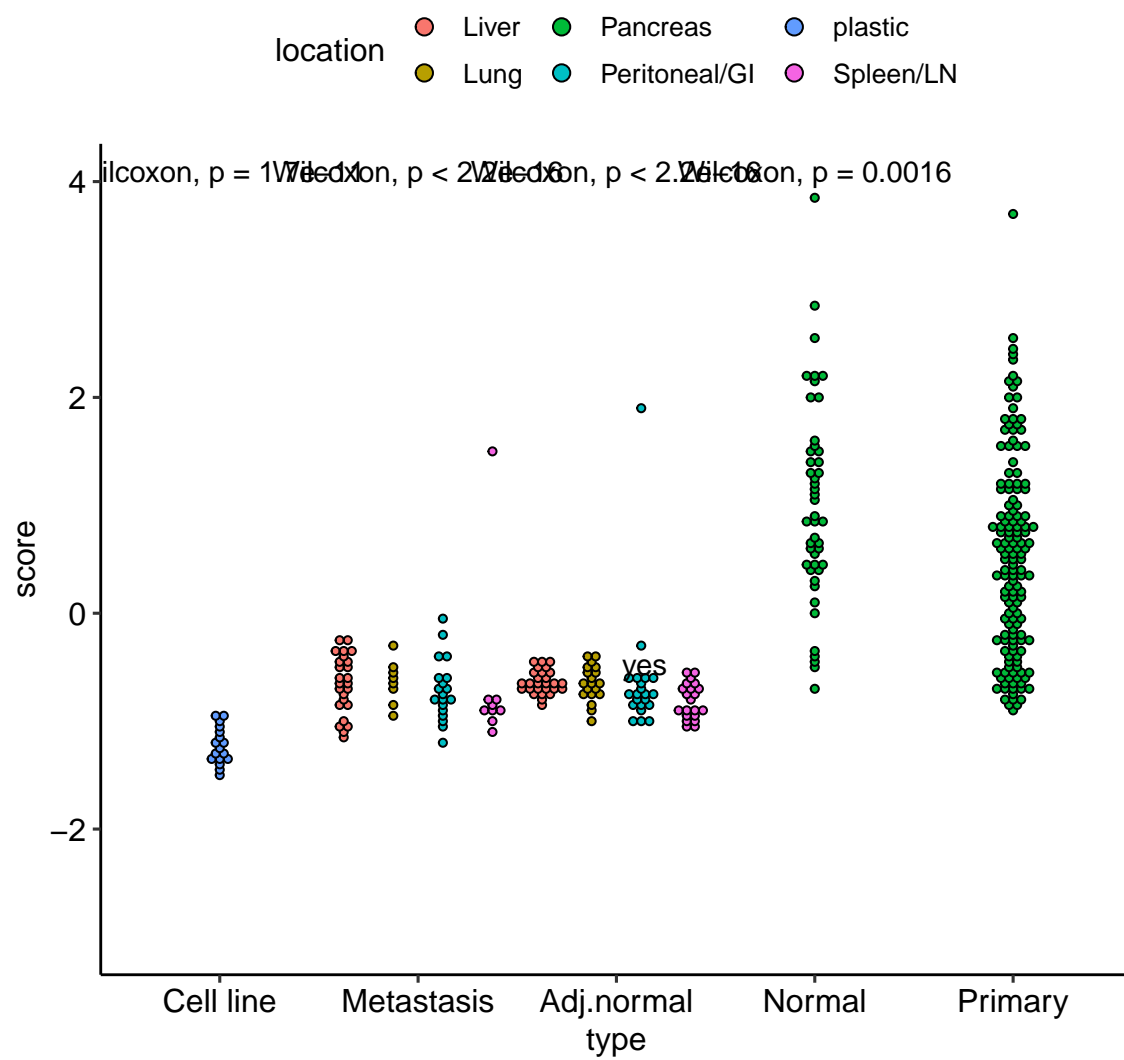


# Moffitt.F12\_Endocrine.top25 score

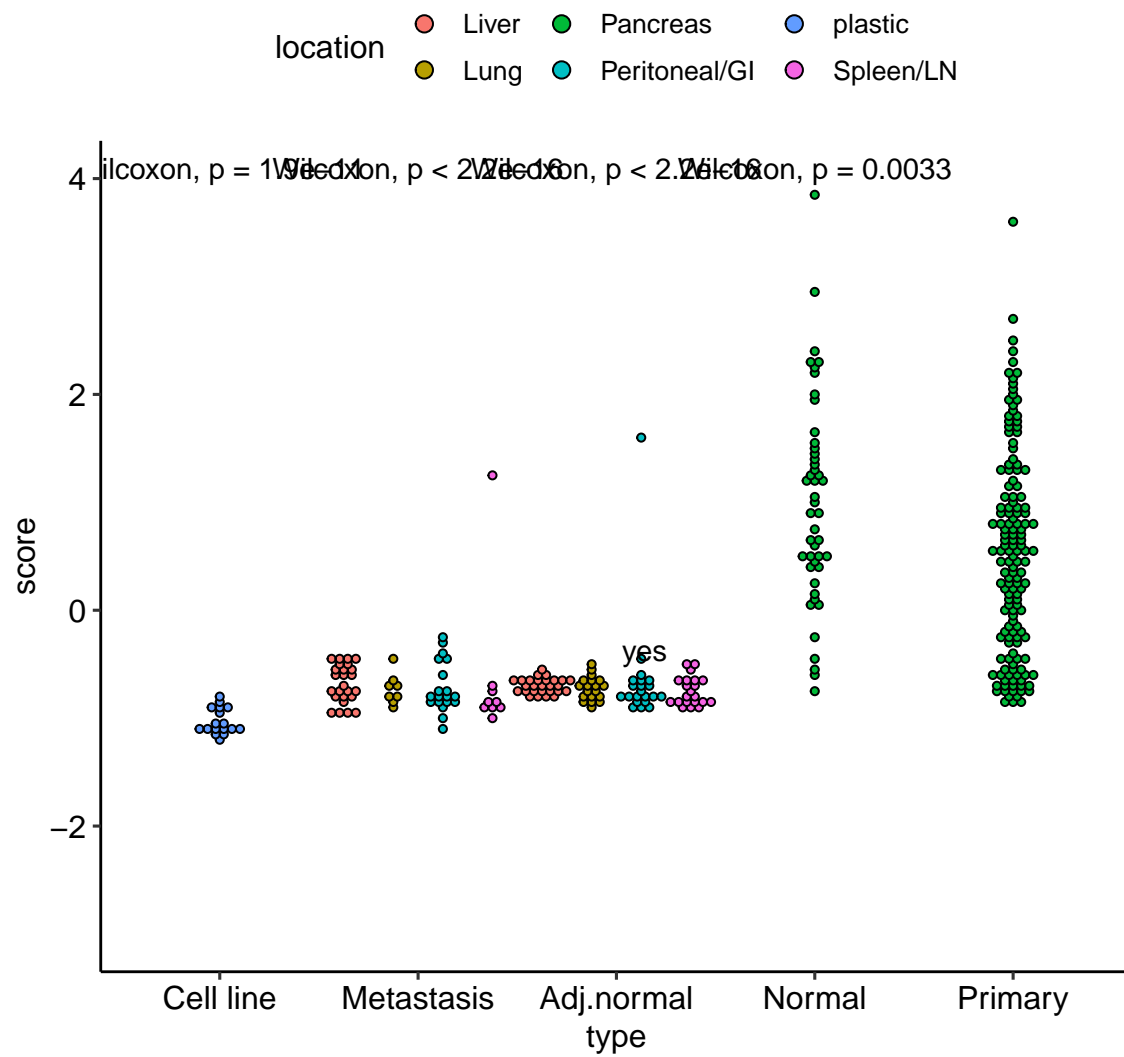




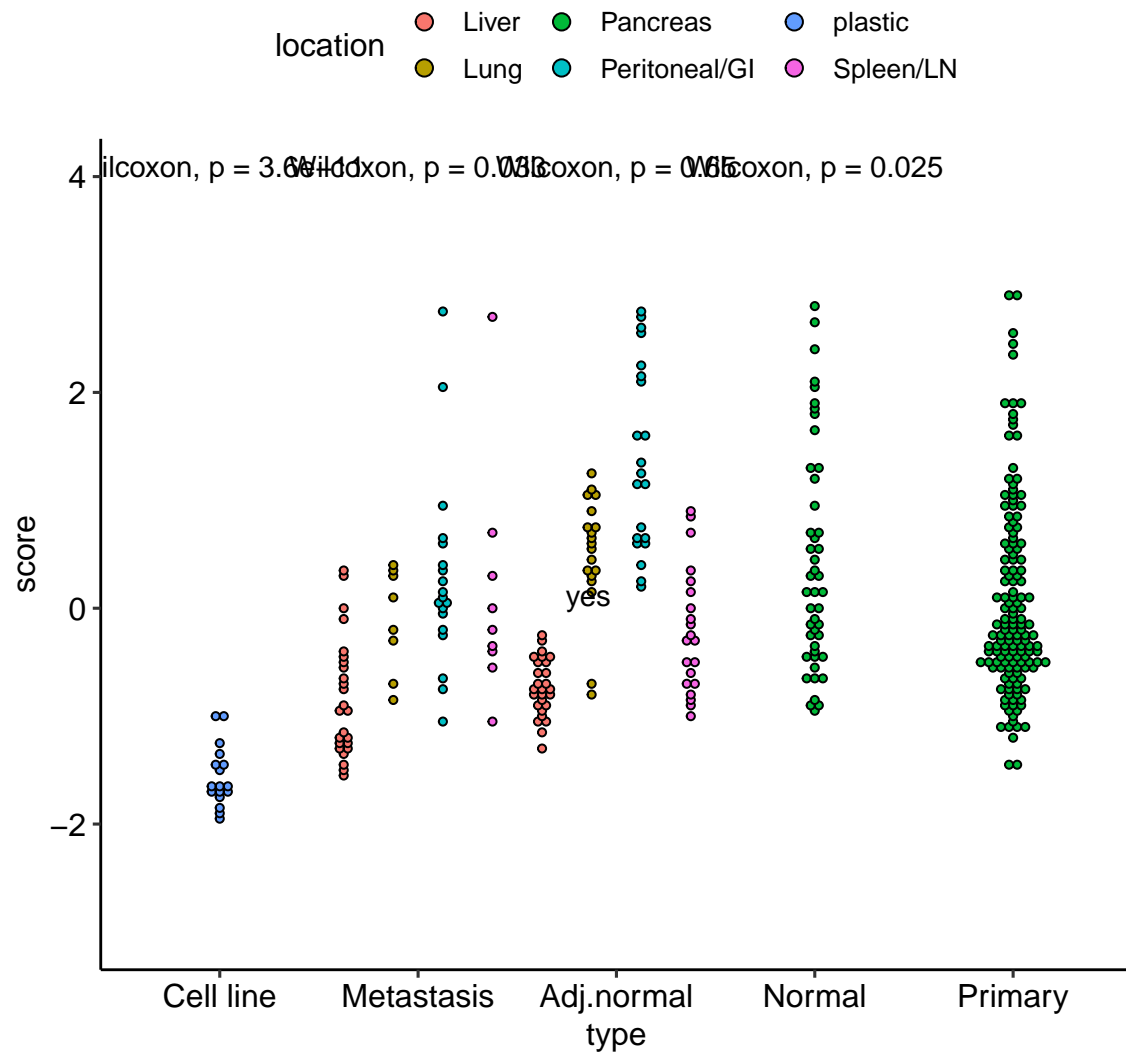
# Moffitt.F12\_Endocrine.top250 score



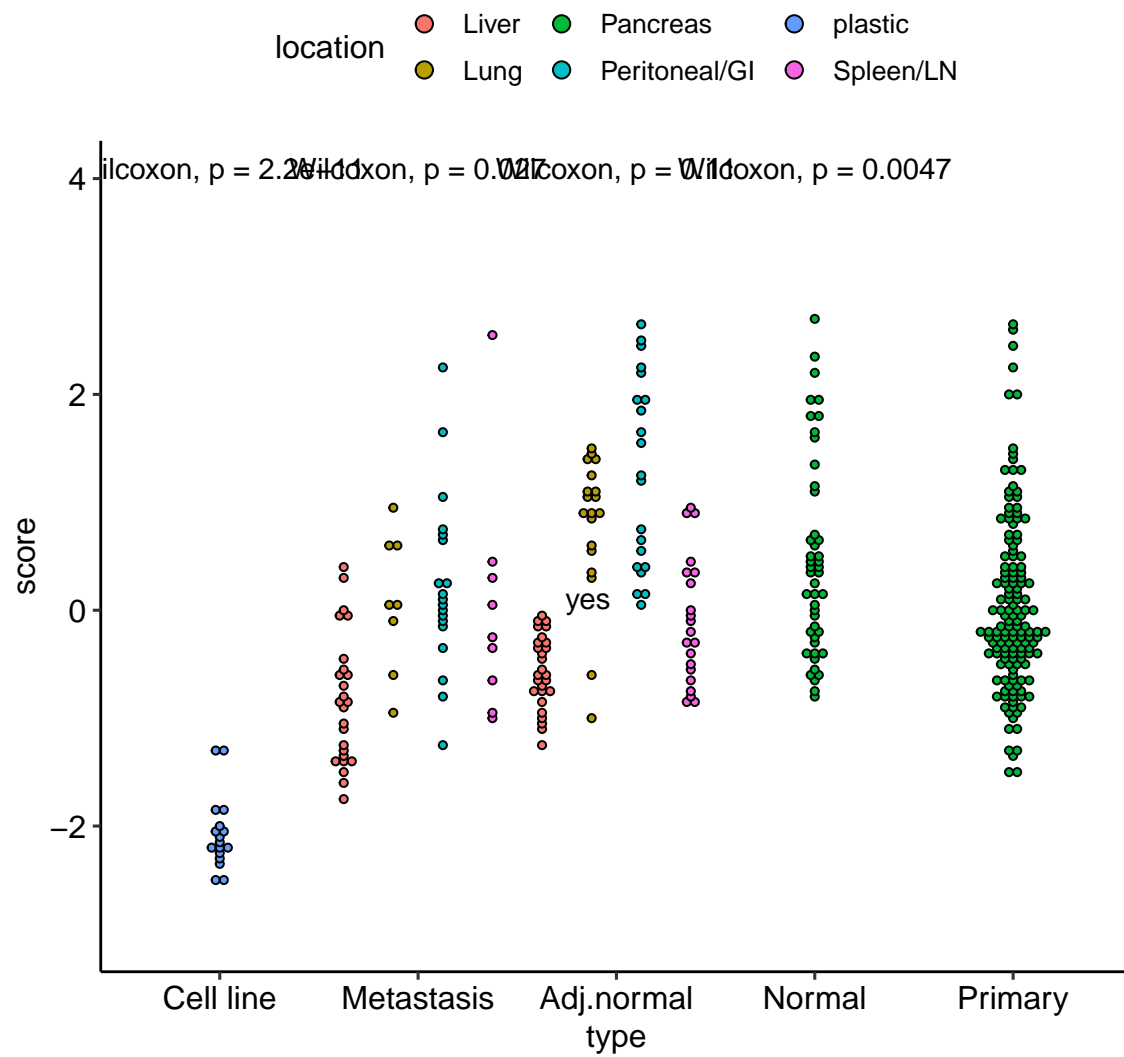
# Moffitt.F12\_Endocrine.top100 score



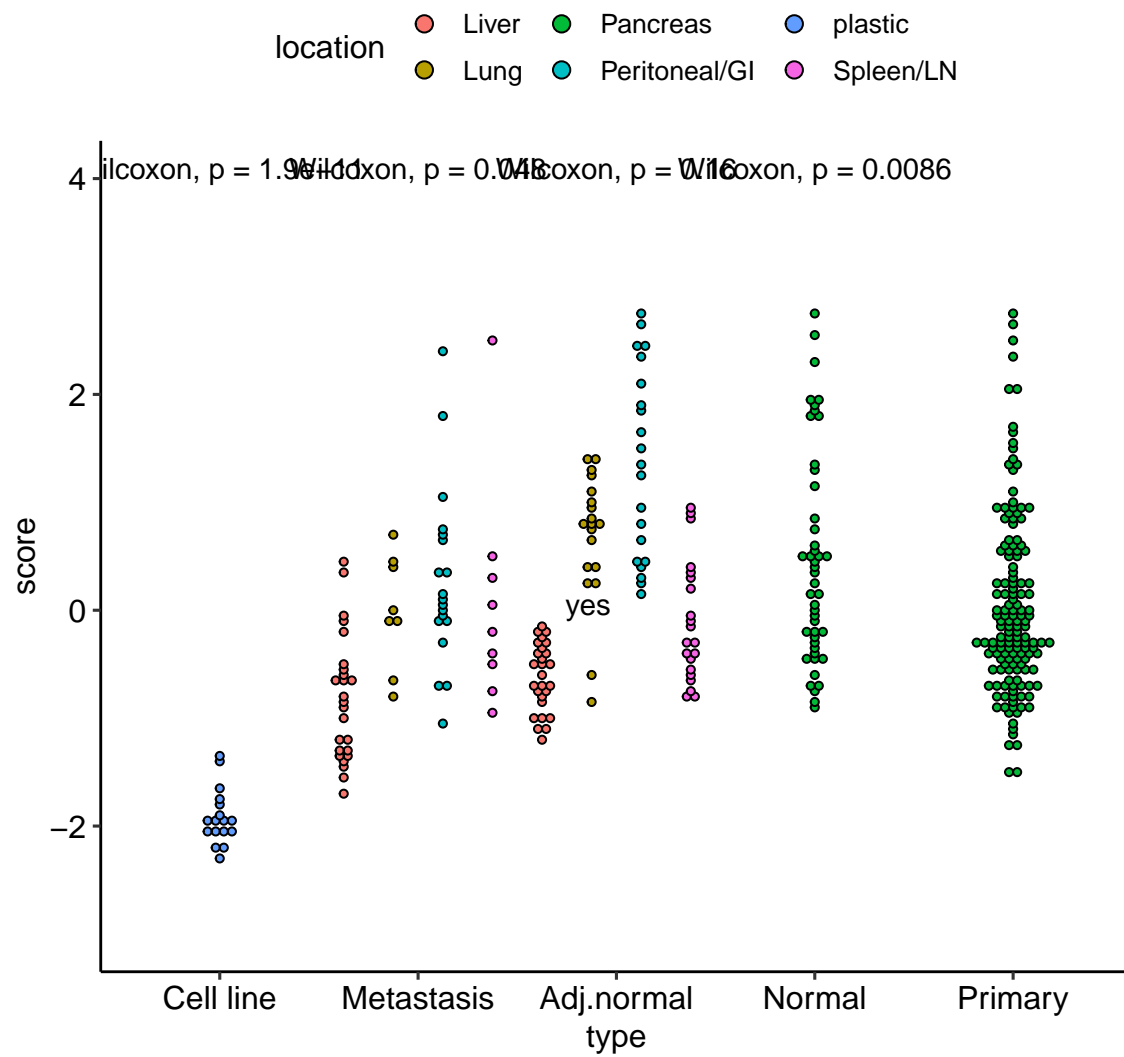
# Moffitt.F13\_NormalStroma.top25 score



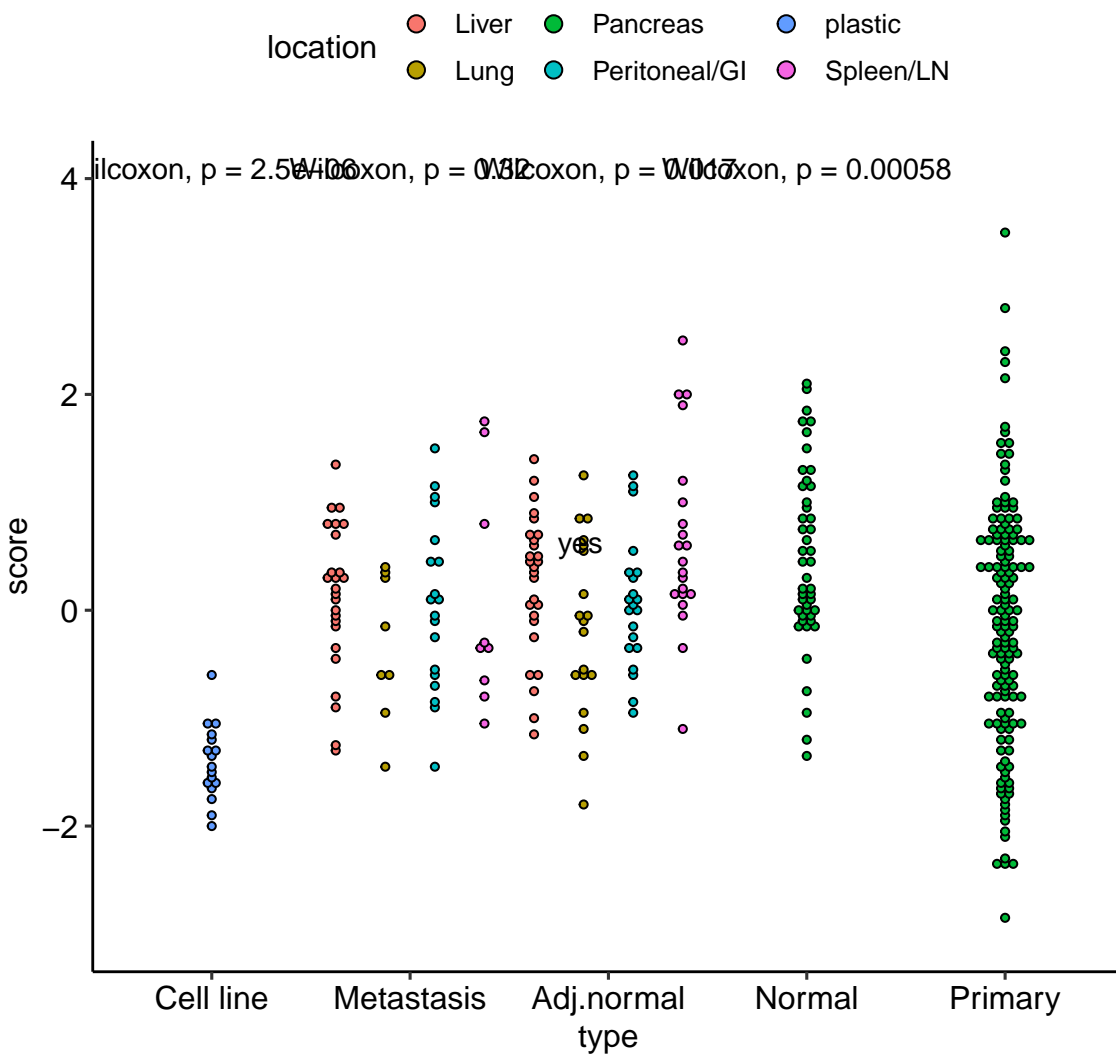
# Moffitt.F13\_NormalStroma.top250 score



# Moffitt.F13\_NormalStroma.top100 score



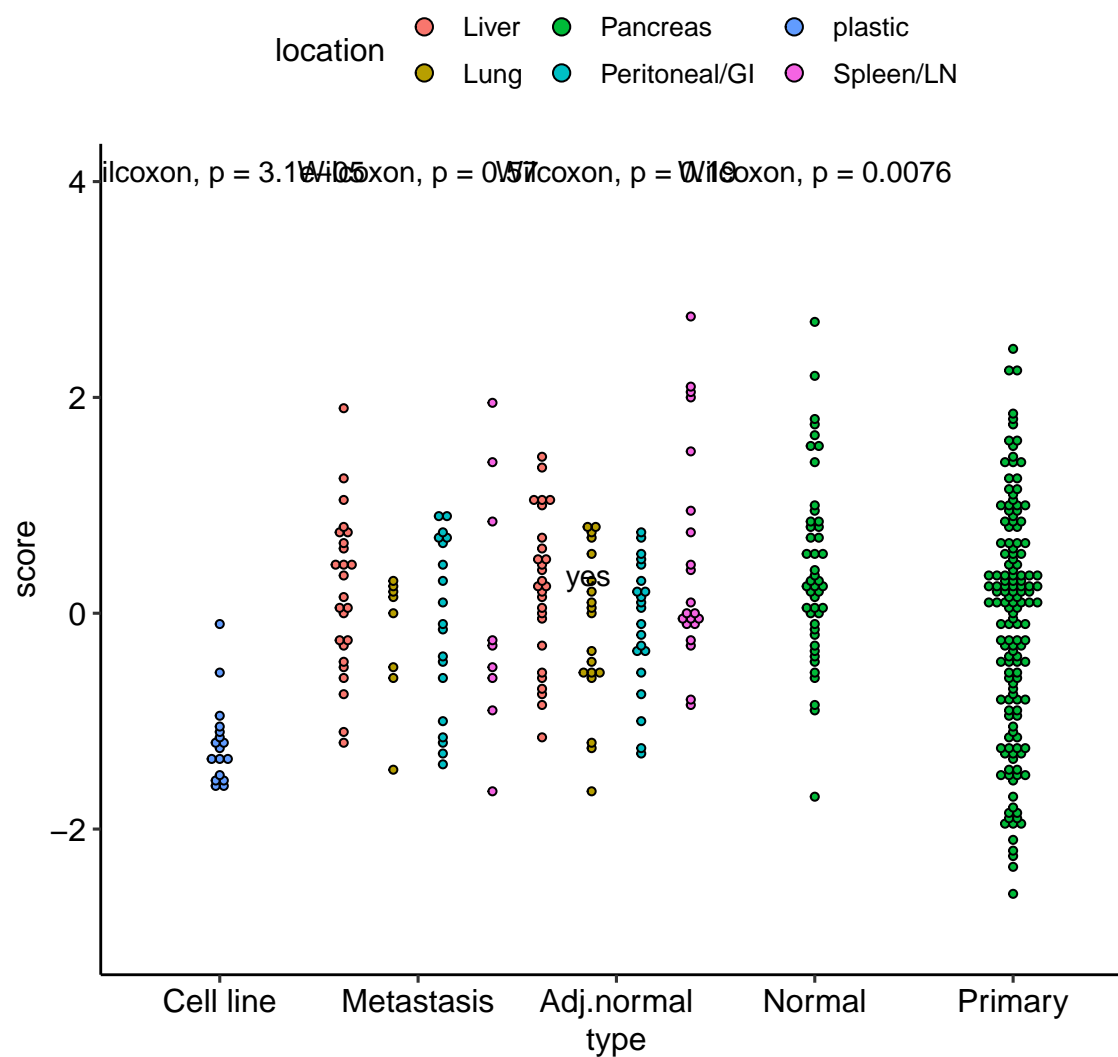
## Moffitt.F14.top25 score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

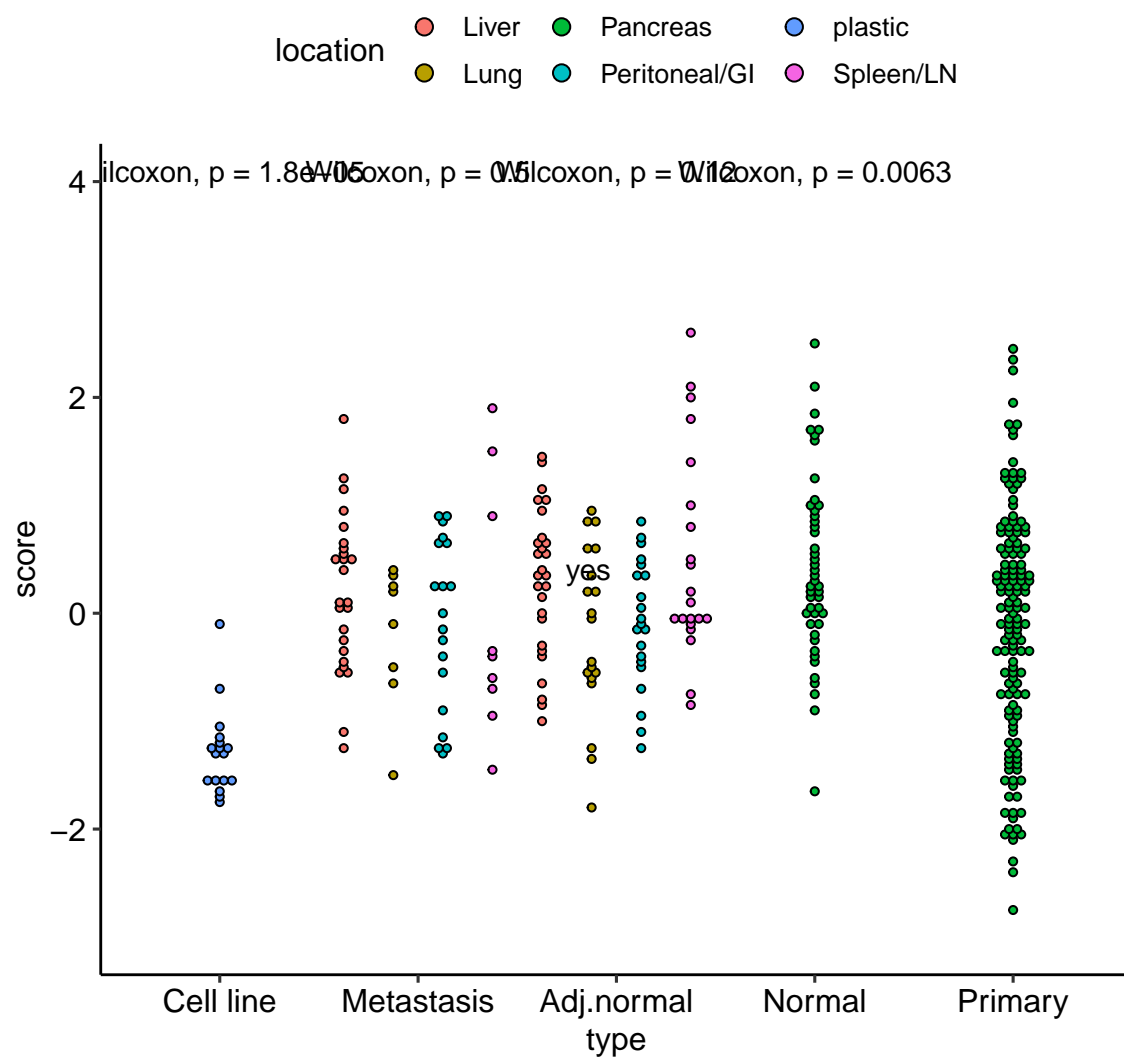
## Moffitt.F14.top250 score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

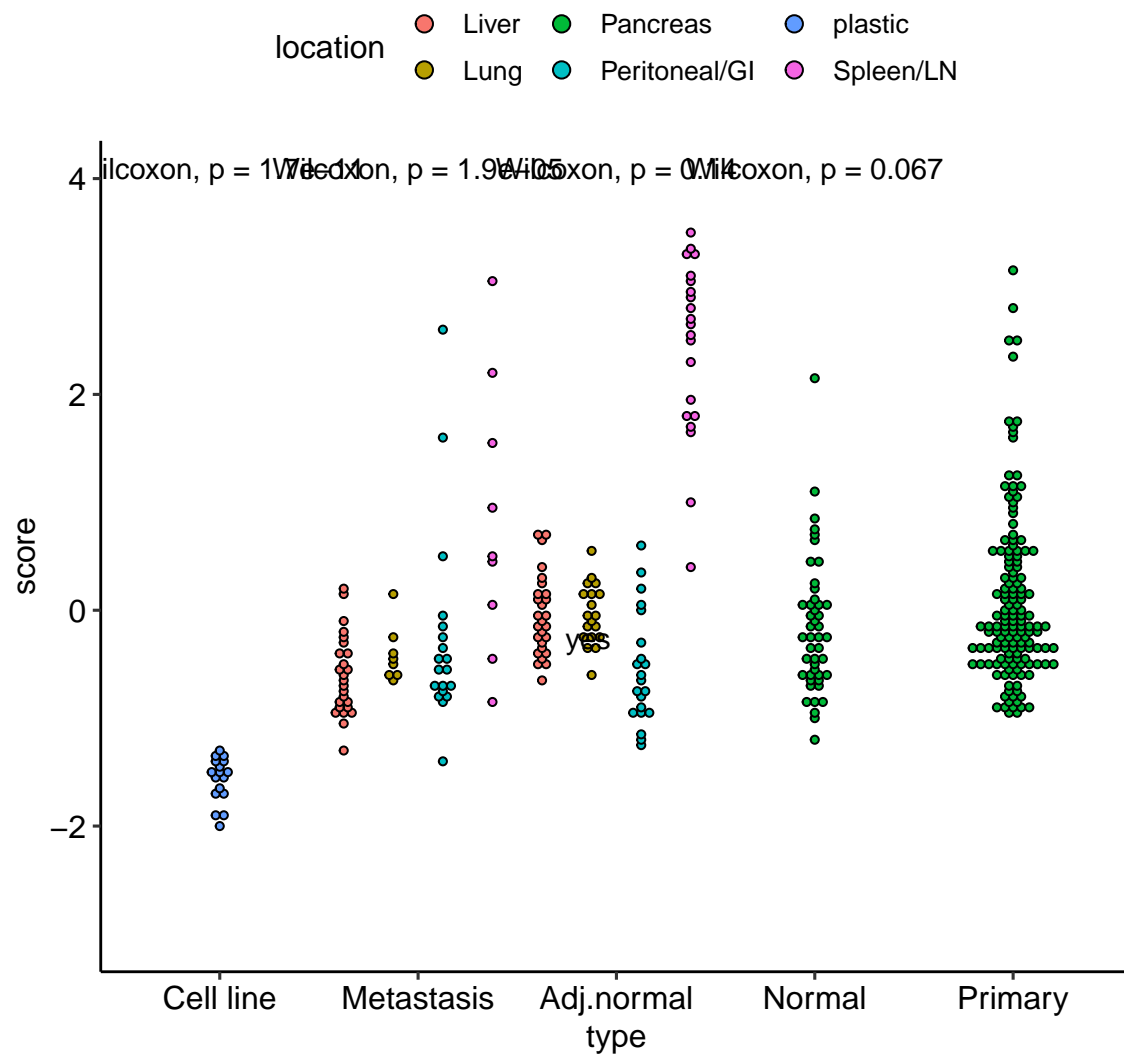
## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

# Moffitt.F14.top100 score

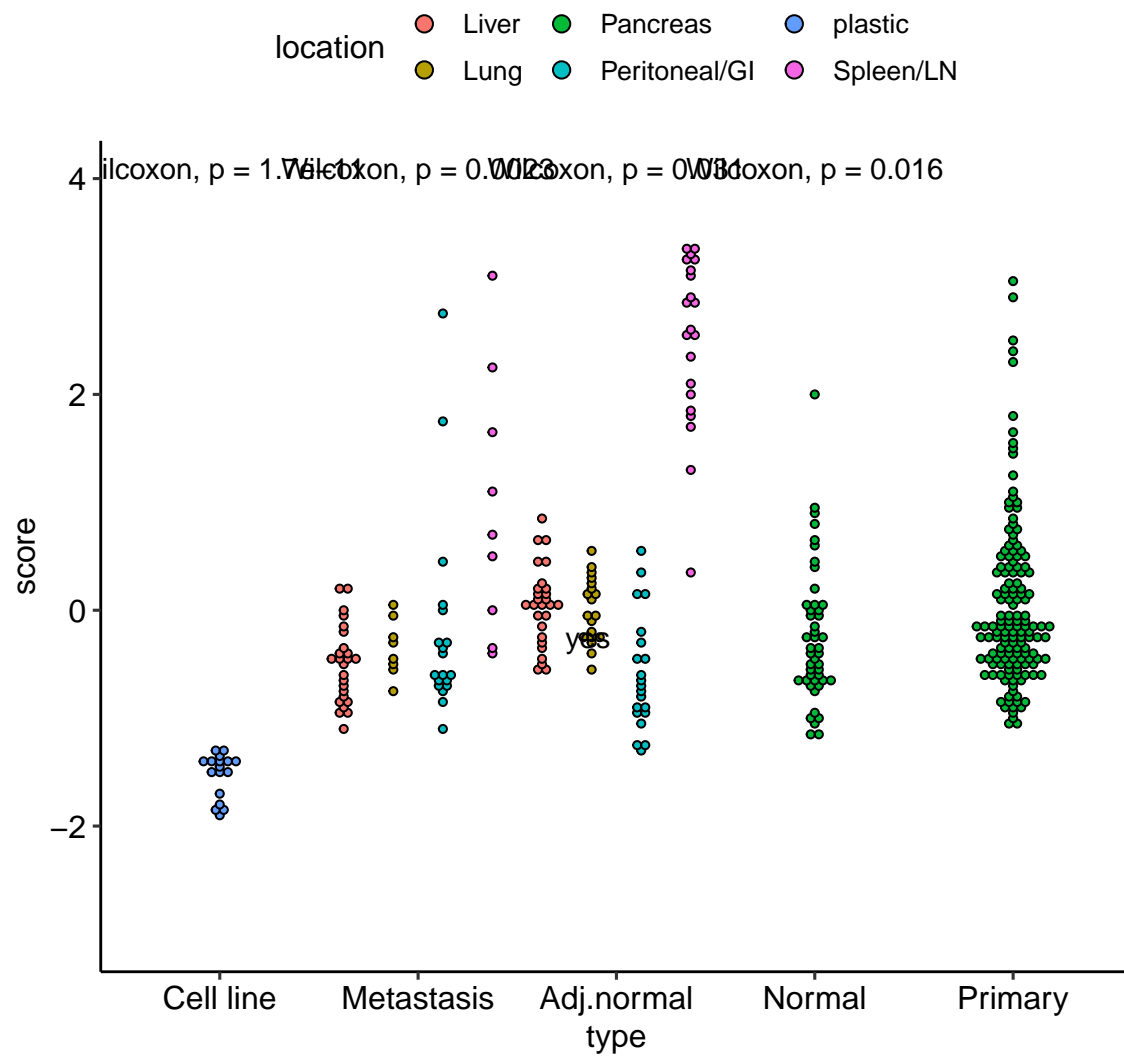




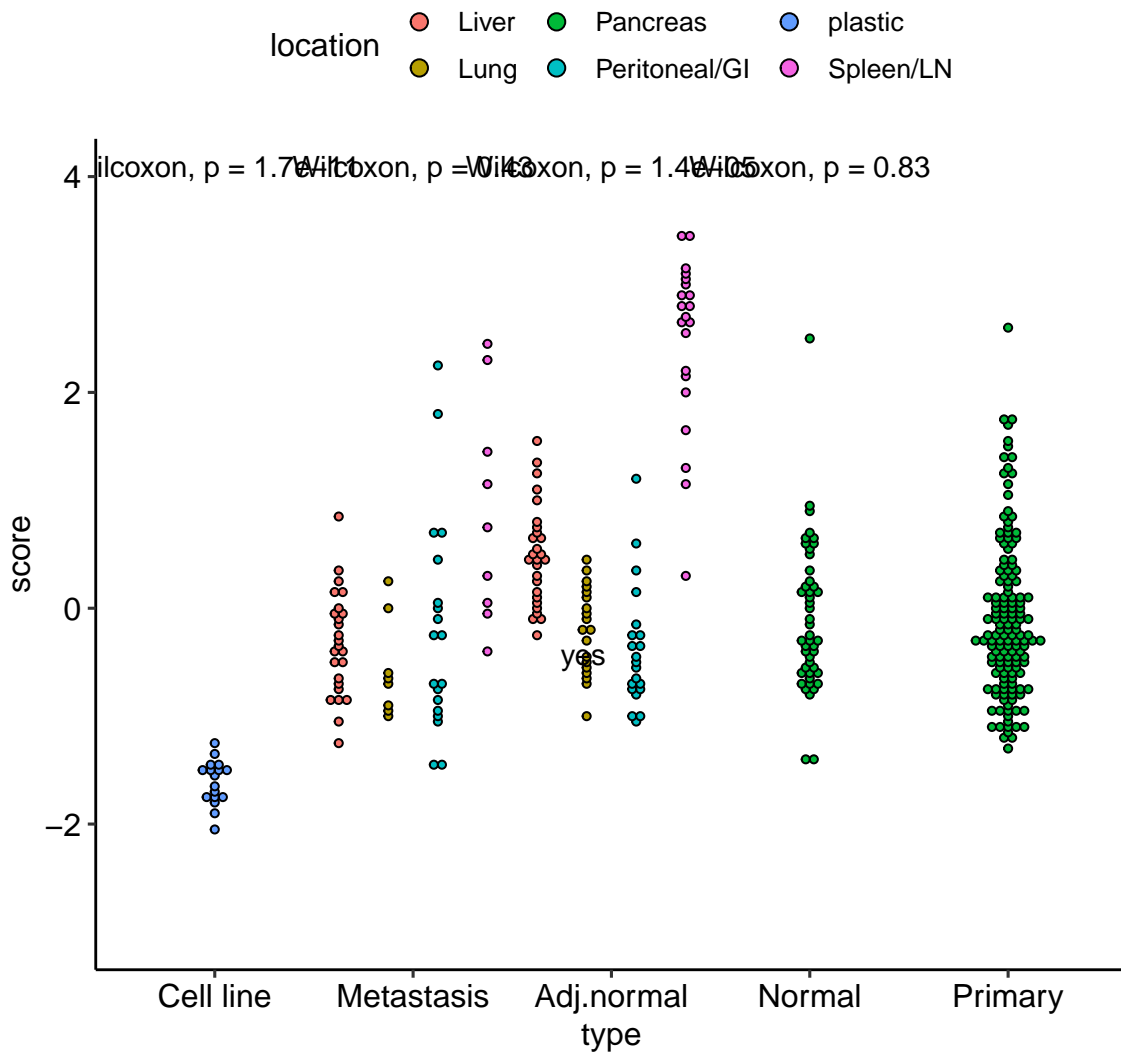
# CIBERSORT.B.cells.naive score



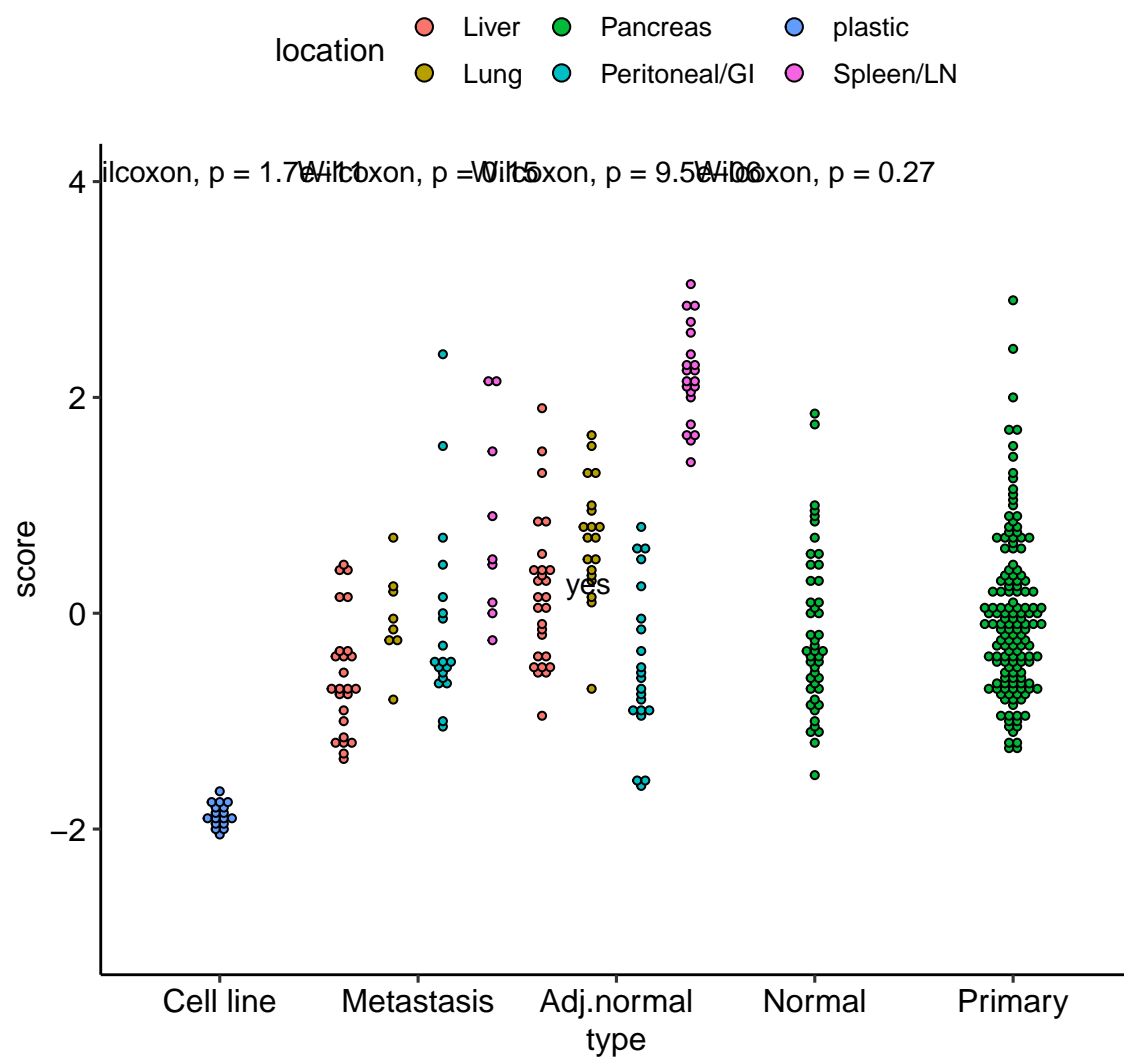
# CIBERSORT.B.cells.memory score



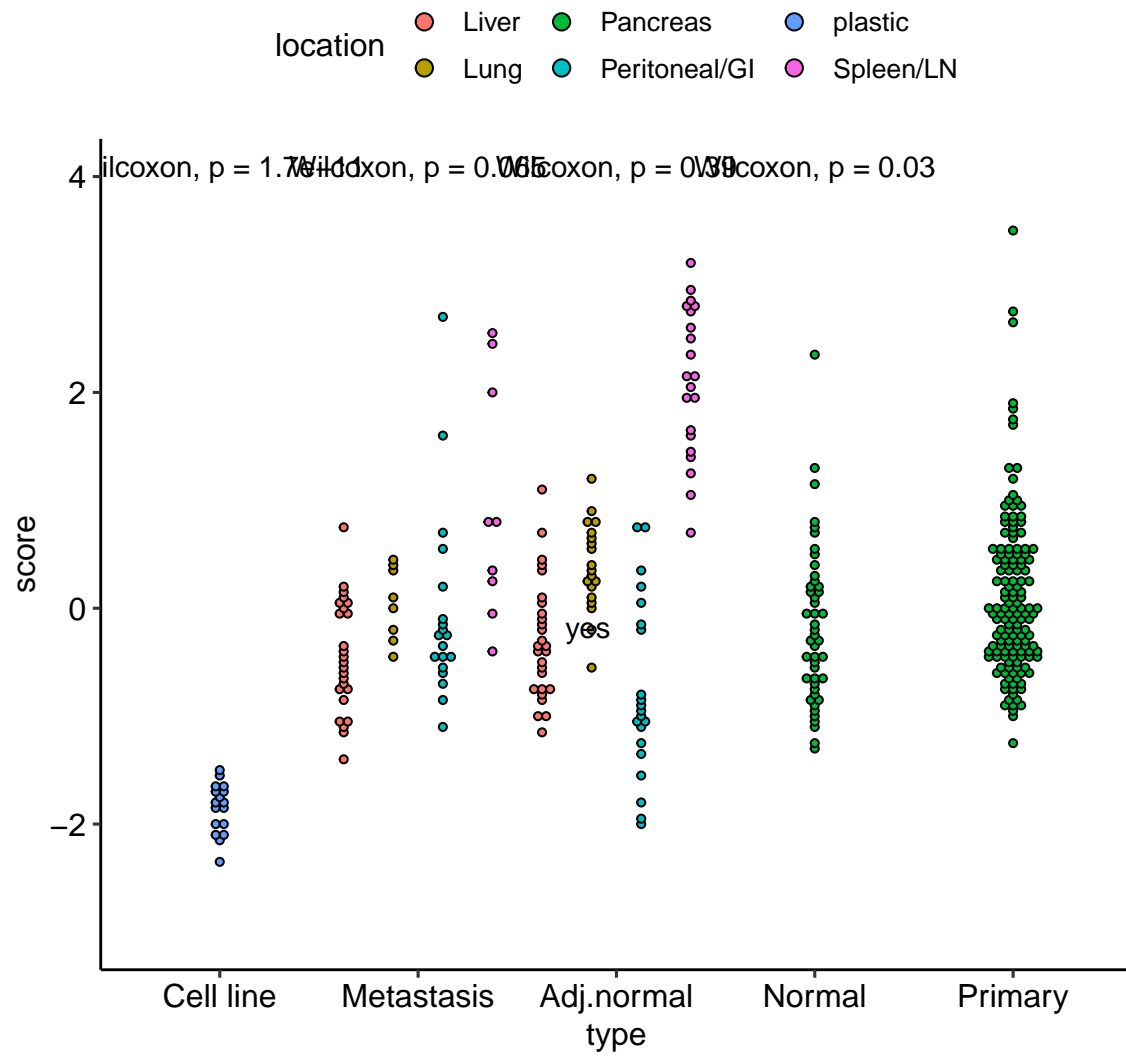
# CIBERSORT.Plasma.cells score



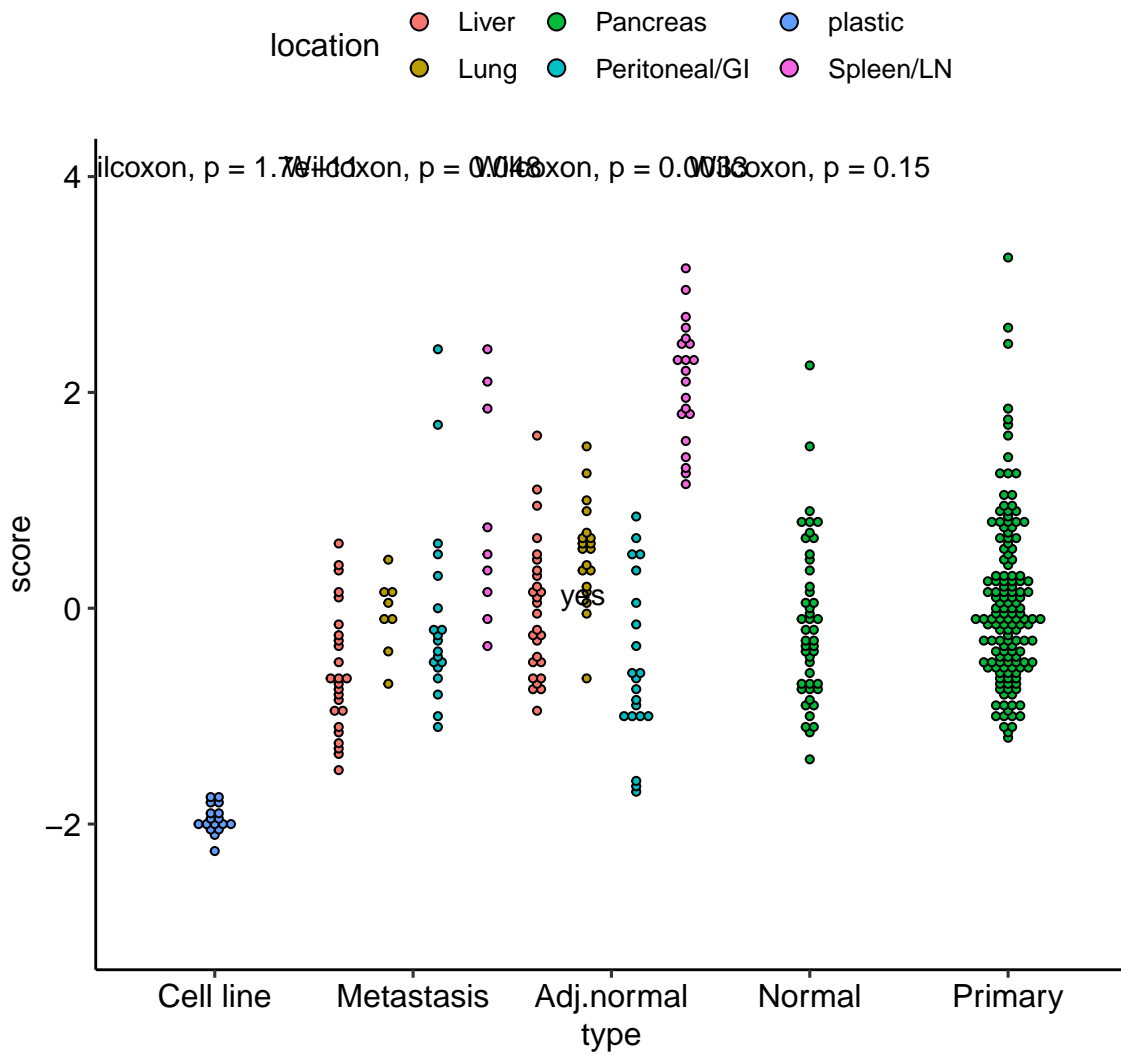
# CIBERSORT.T.cells.CD8 score



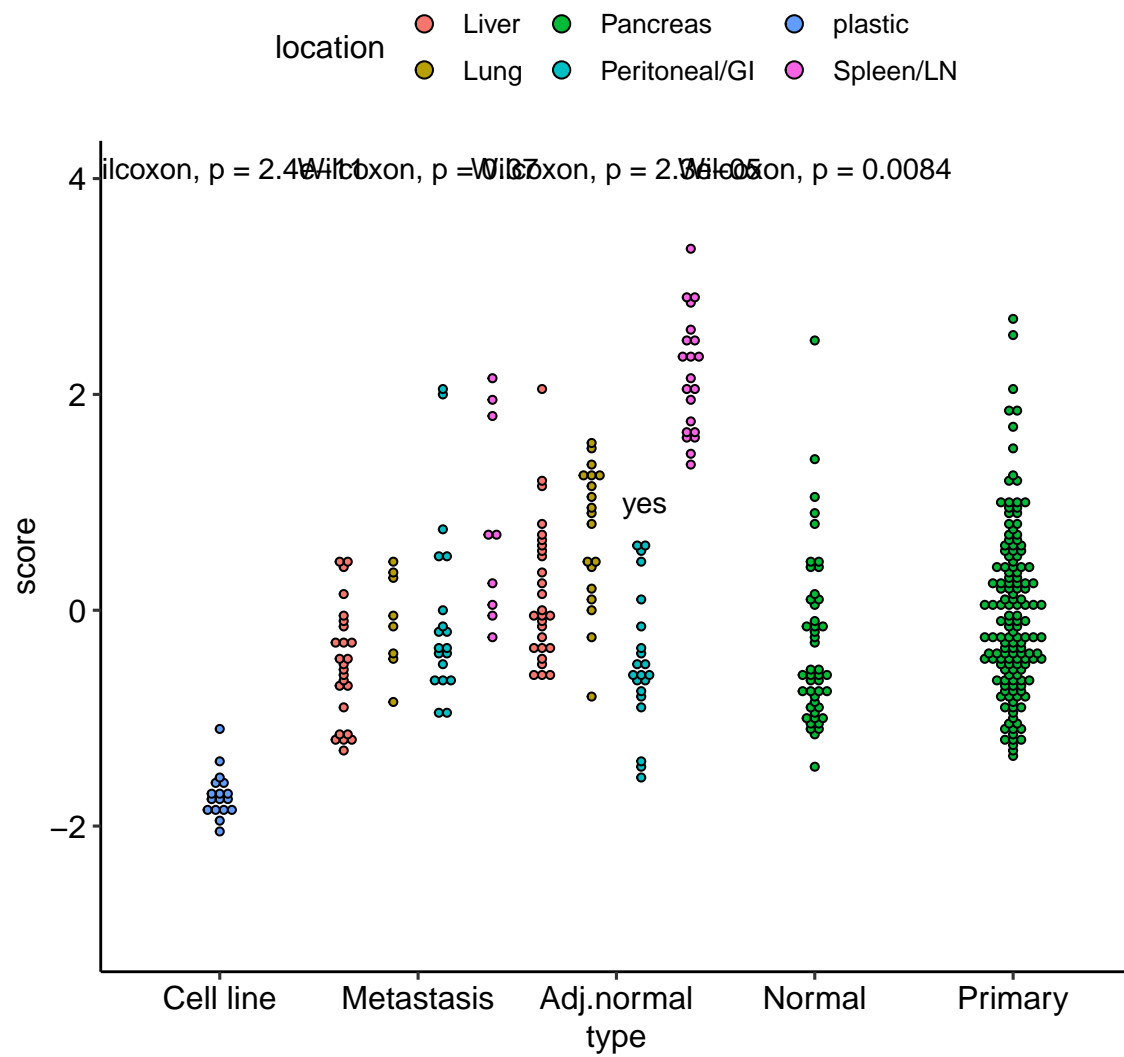
# CIBERSORT.T.cells.CD4.naive score



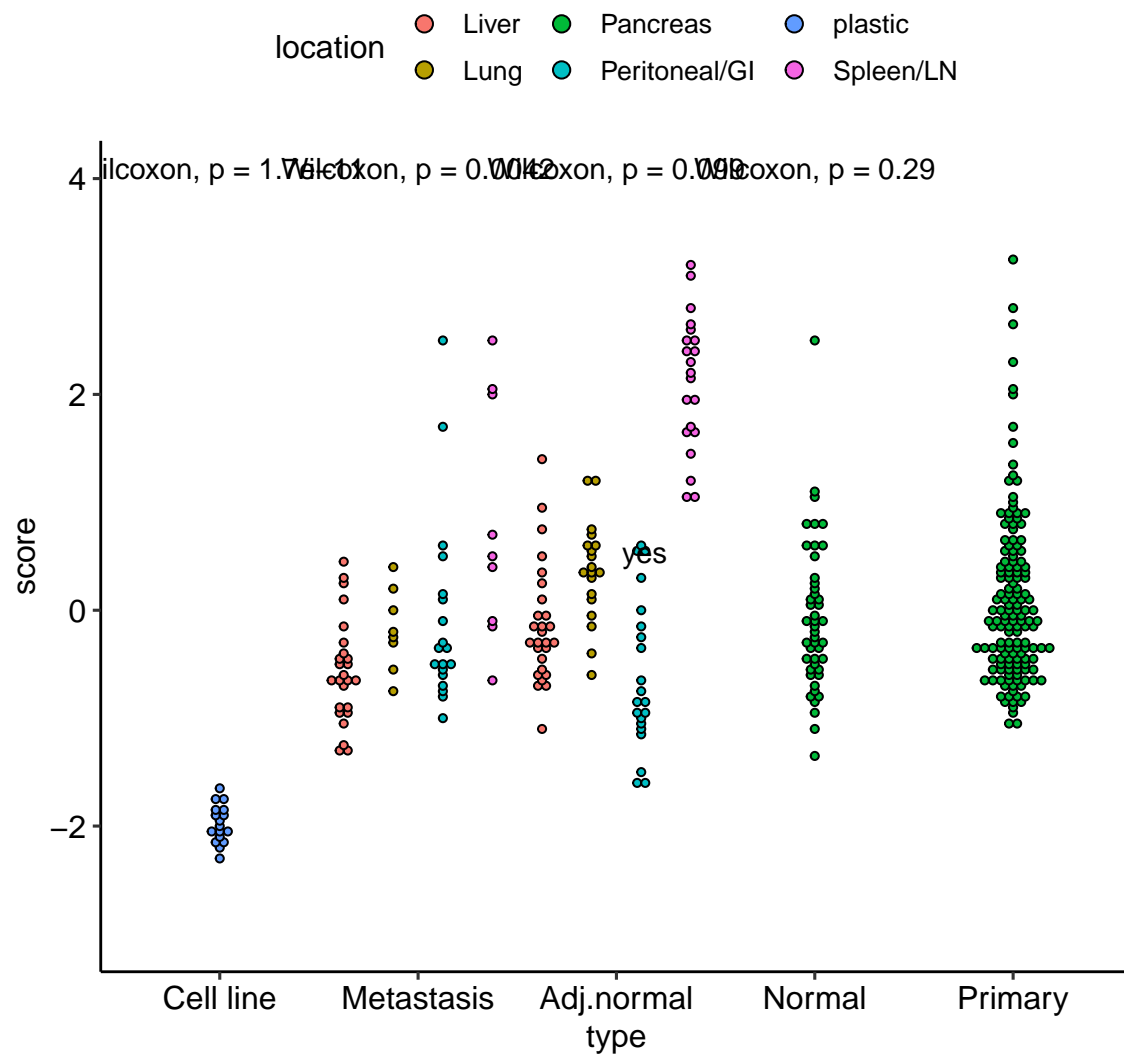
# CIBERSORT.T.cells.CD4.memory.resting score



# CIBERSORT.T.cells.CD4.memory.activated score

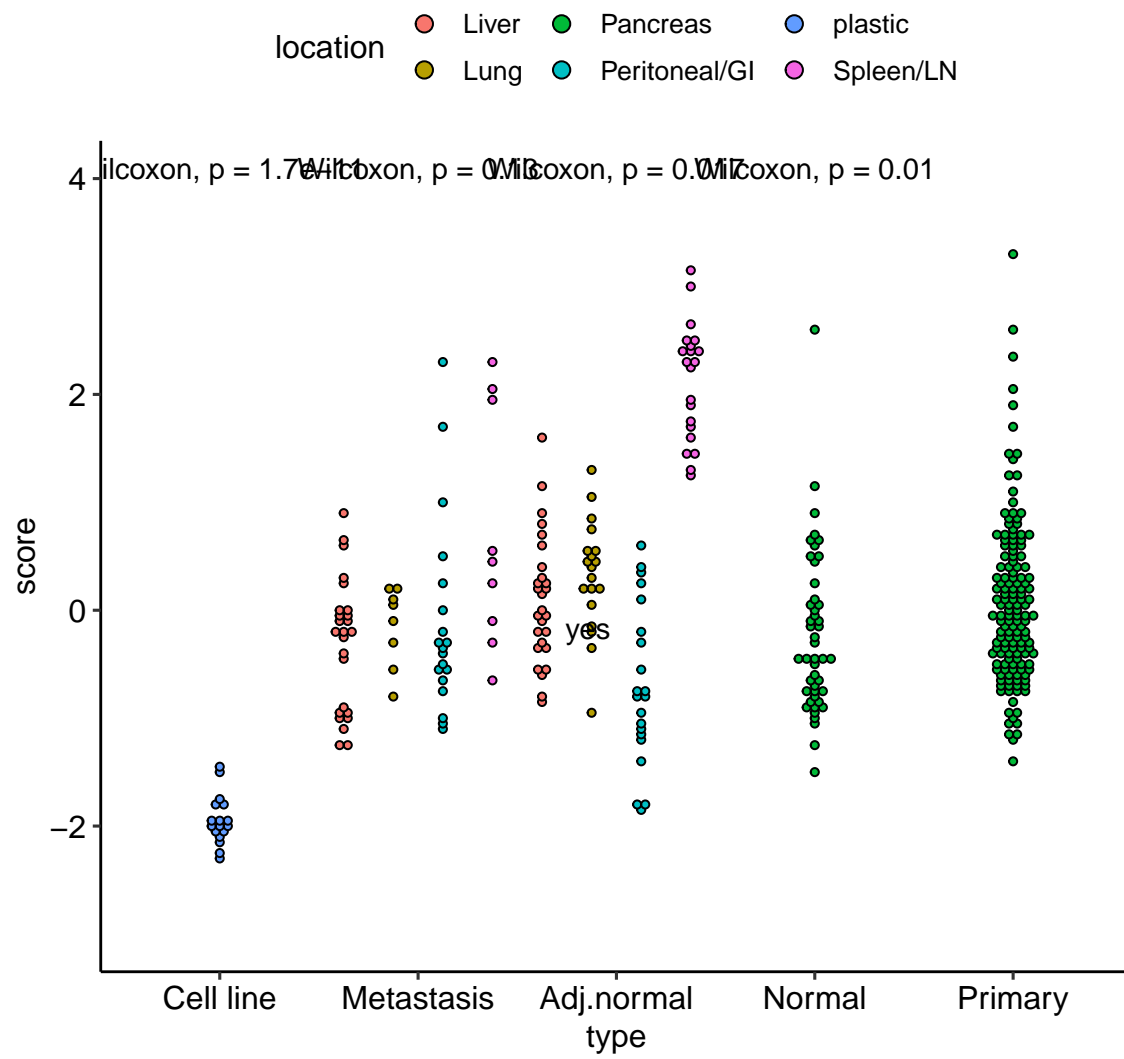


# CIBERSORT.T.cells.follicular.helper score

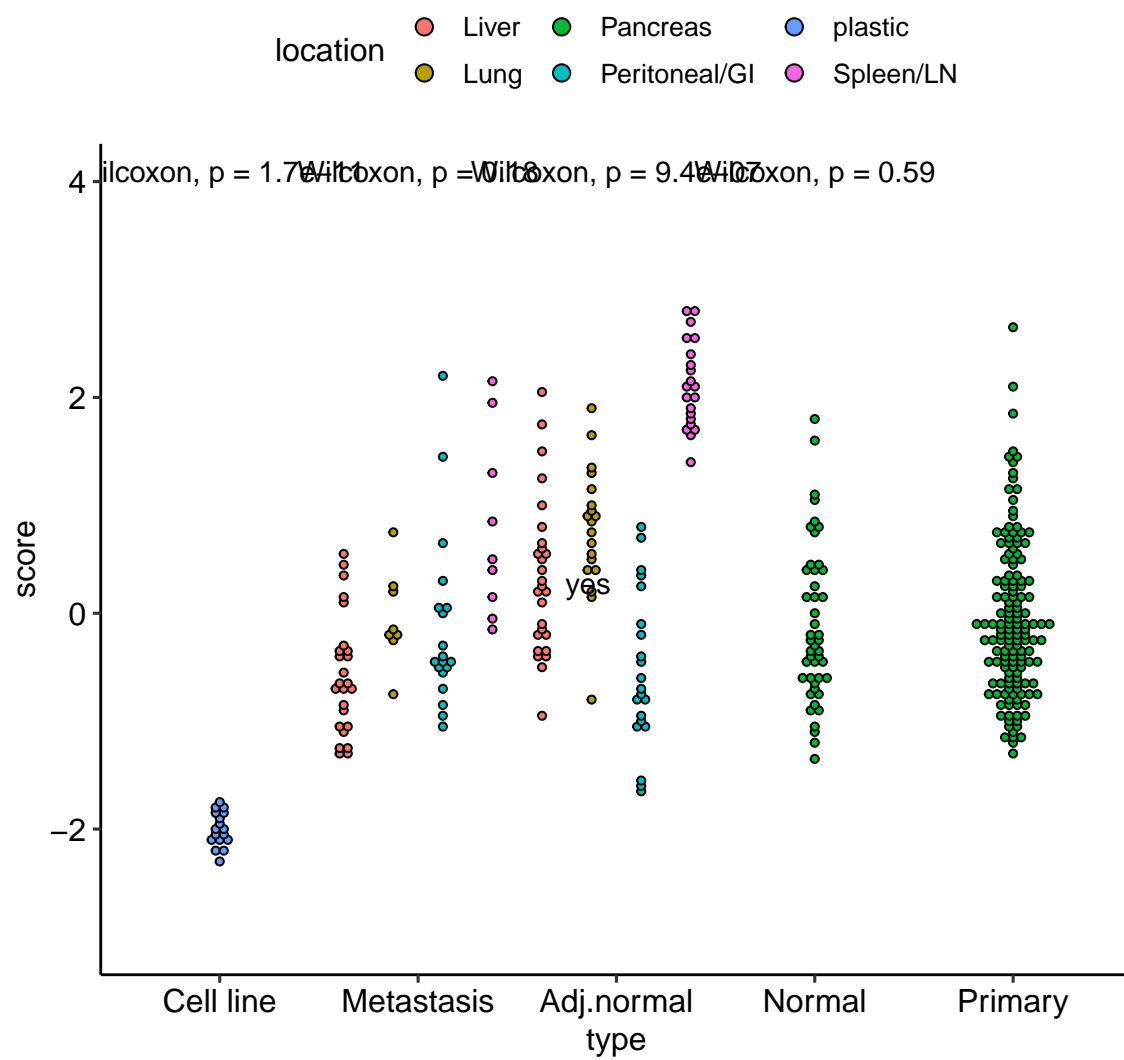




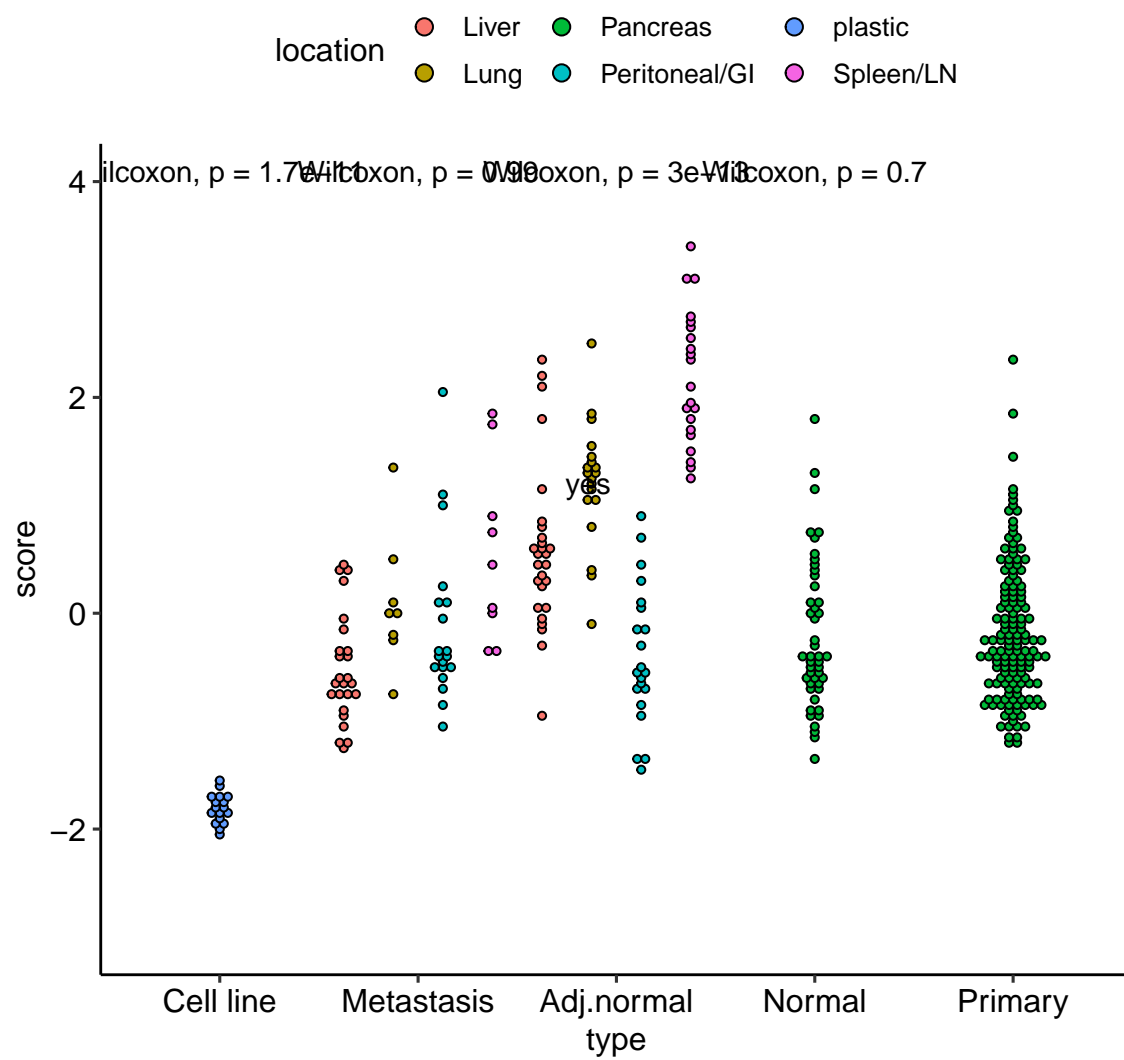
# CIBERSORT.T.cells.regulatory..Tregs. score



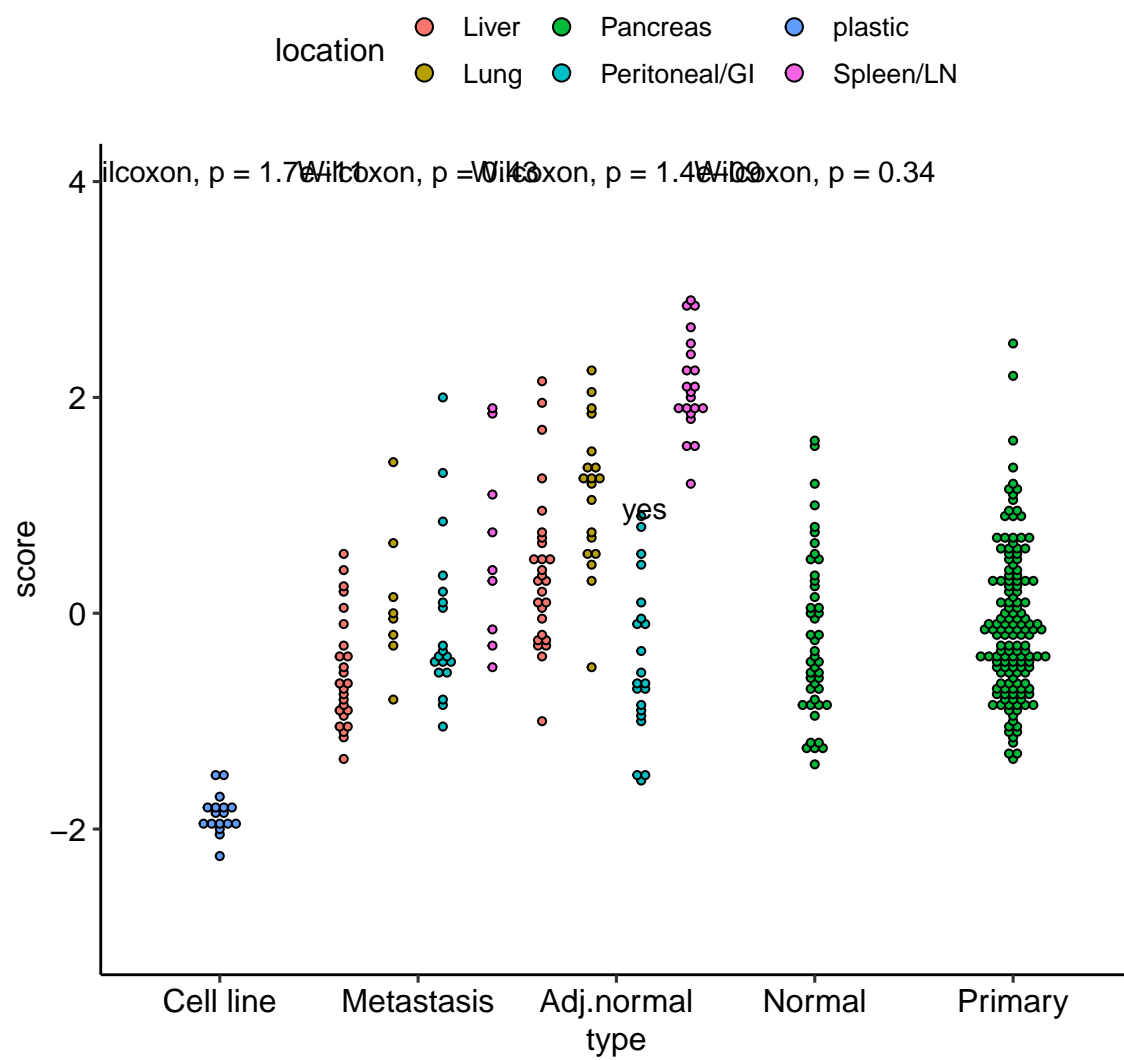
# CIBERSORT.T.cells.gamma.delta score



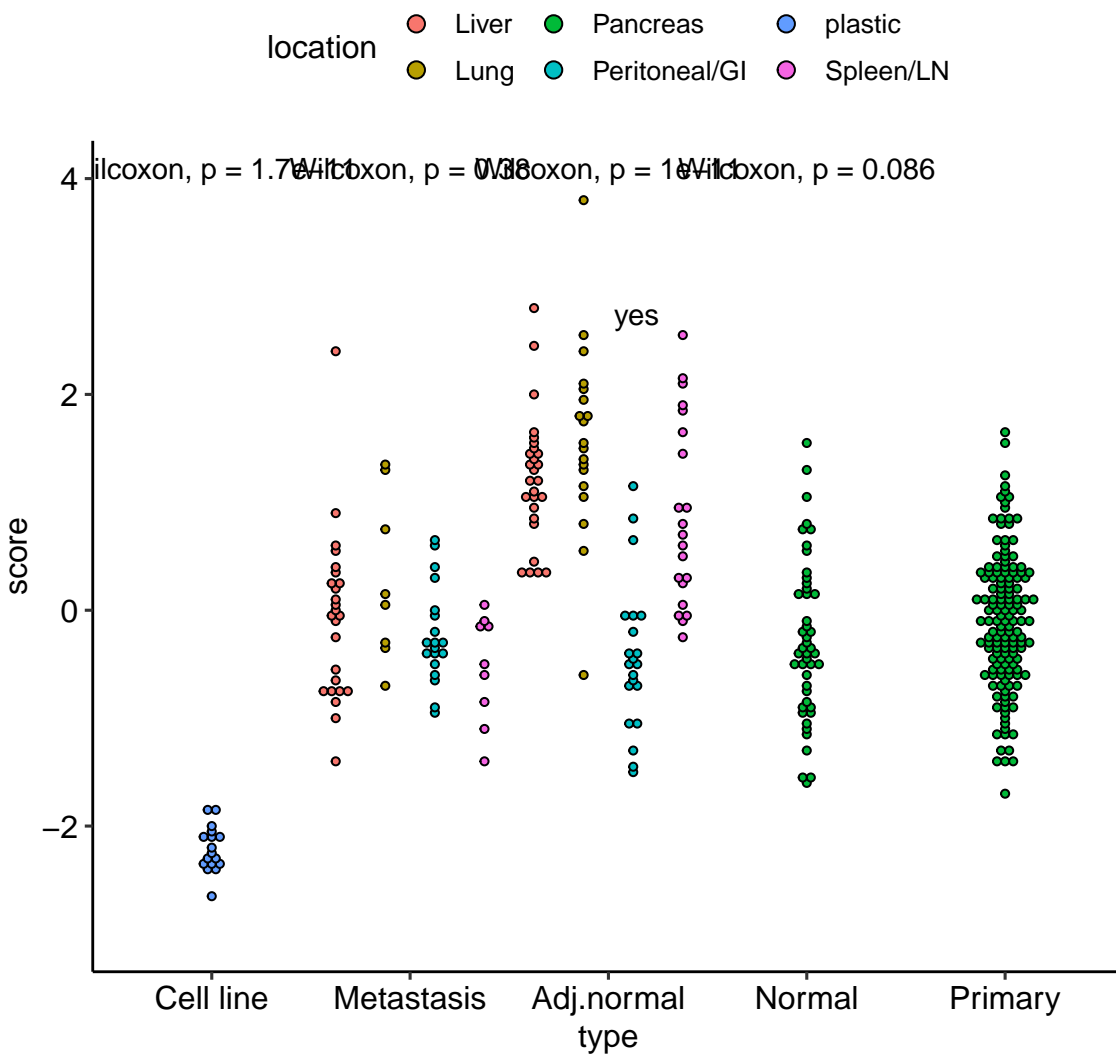
# CIBERSORT.NK.cells.resting score



# CIBERSORT.NK.cells.activated score



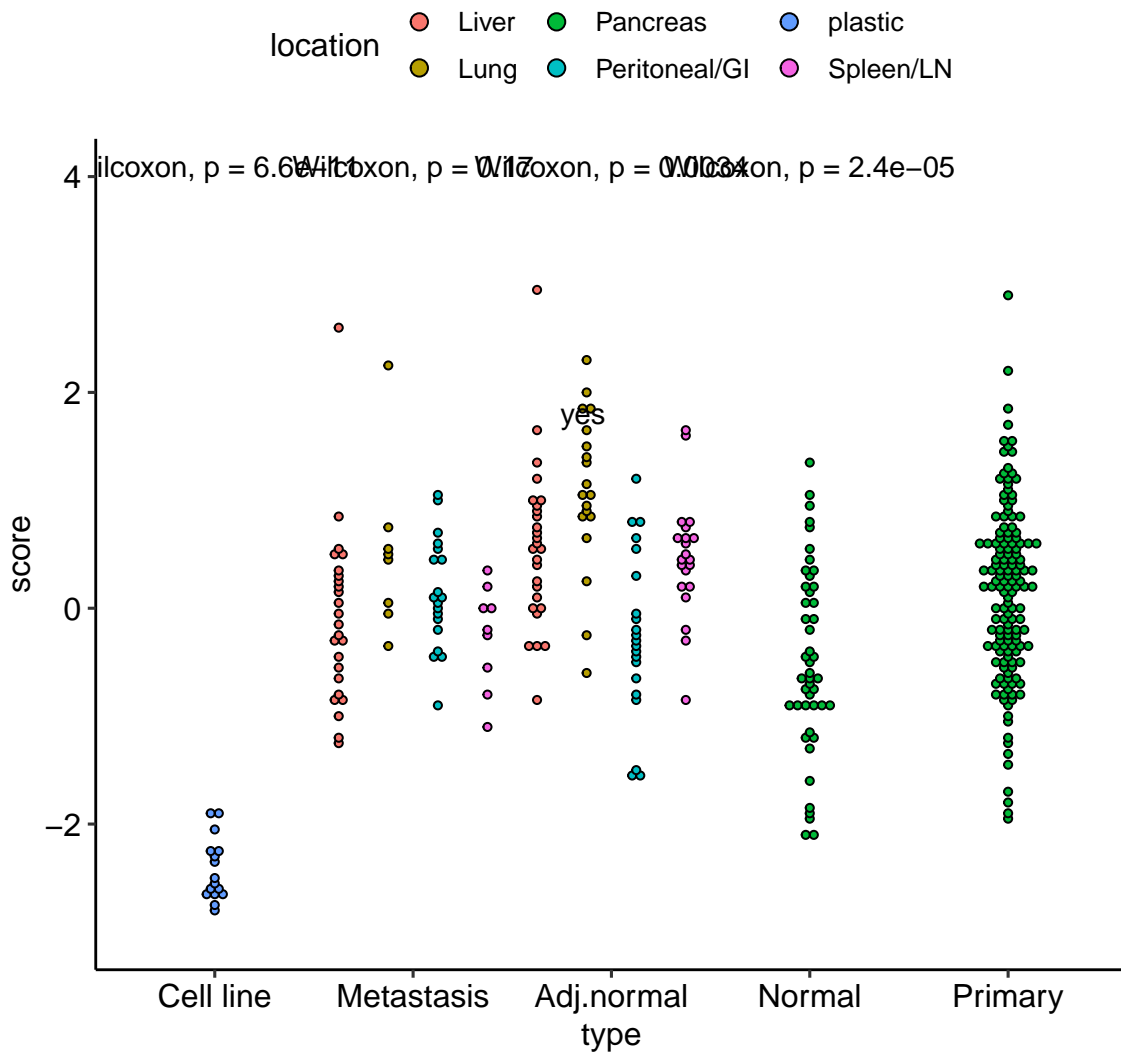
## CIBERSORT.Monocytes score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

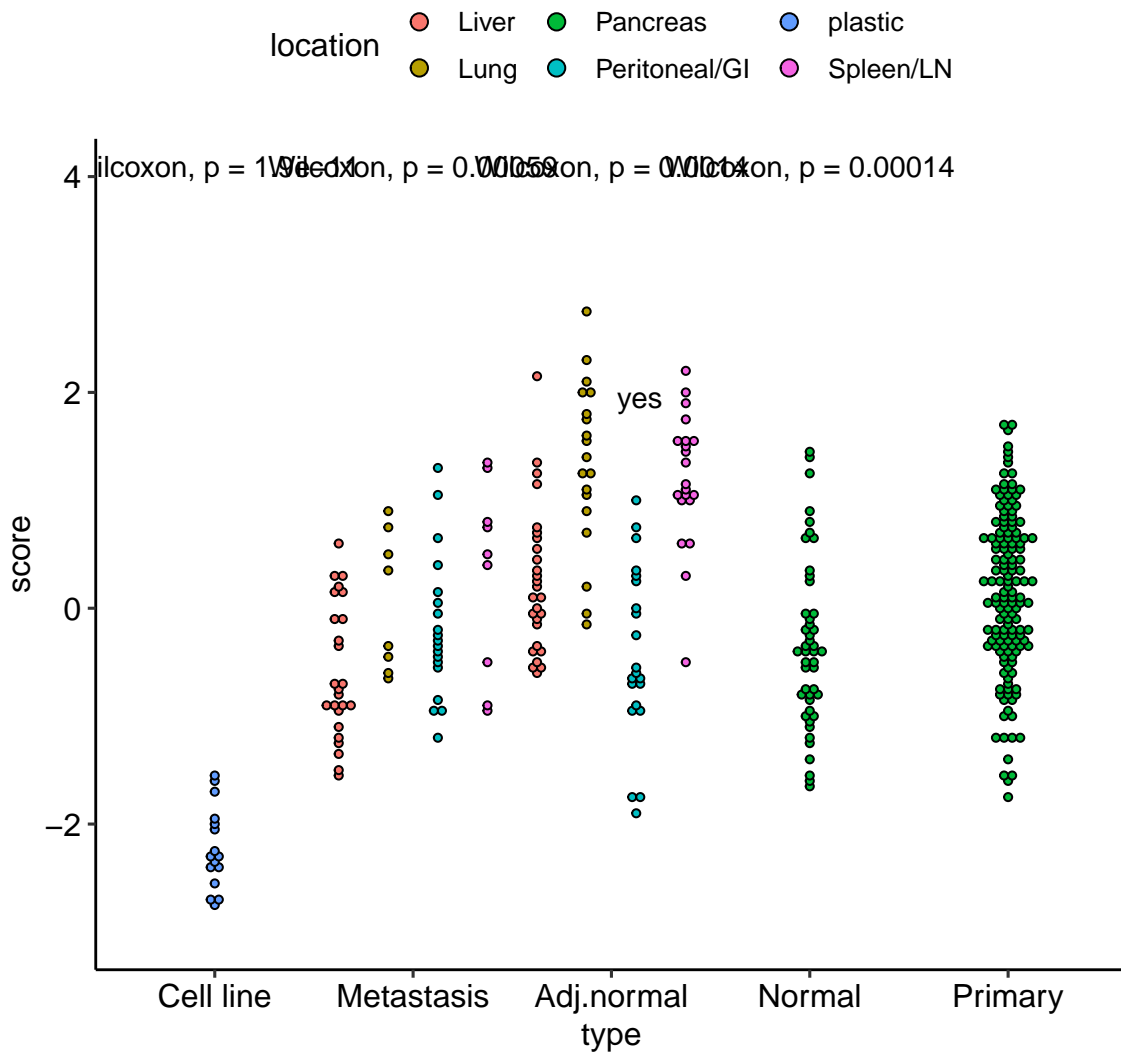
## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

# CIBERSORT.Macrophages.M0 score

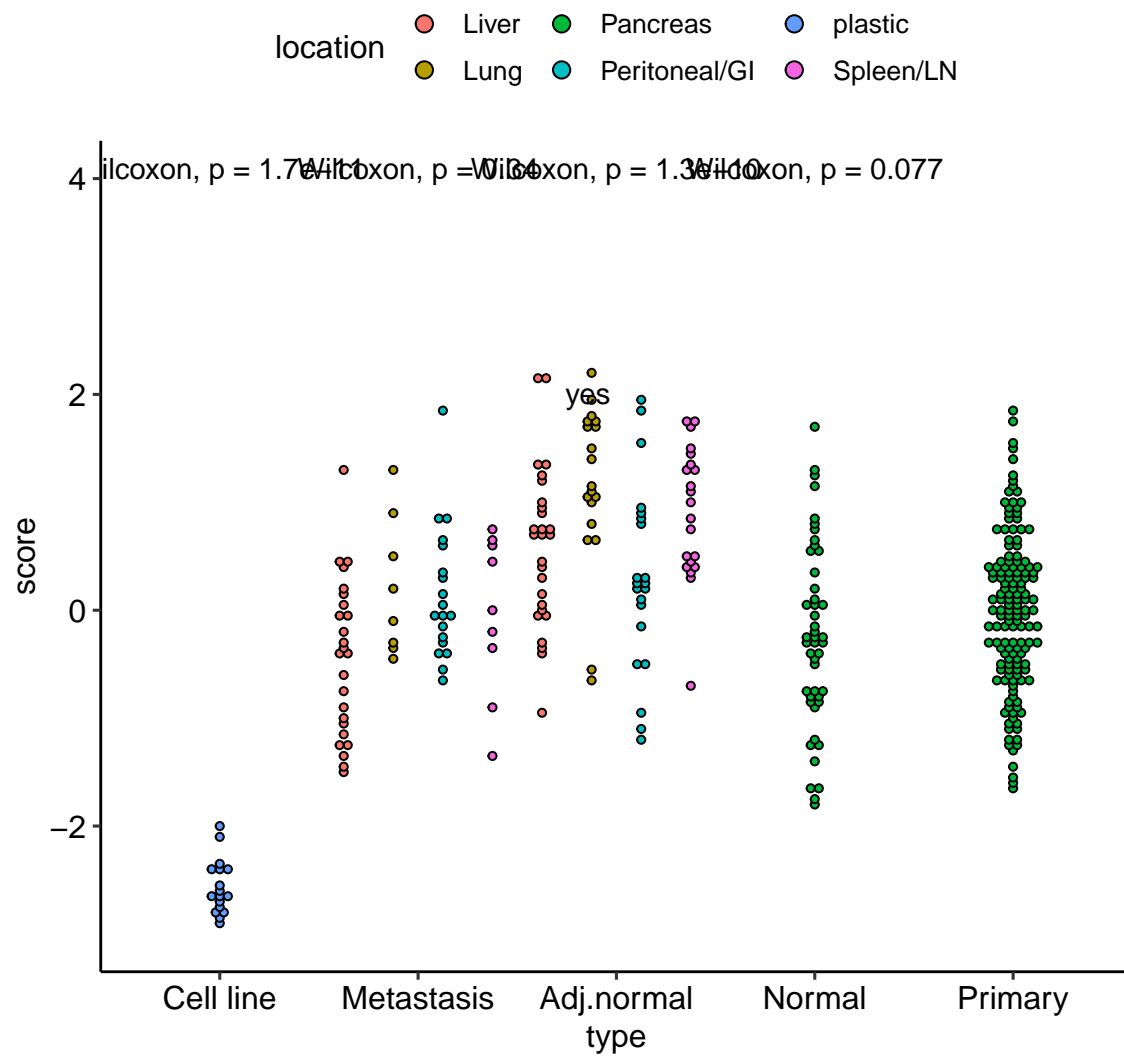


## Warning: Removed 1 rows containing missing values (geom\_dotplot).

# CIBERSORT.Macrophages.M1 score

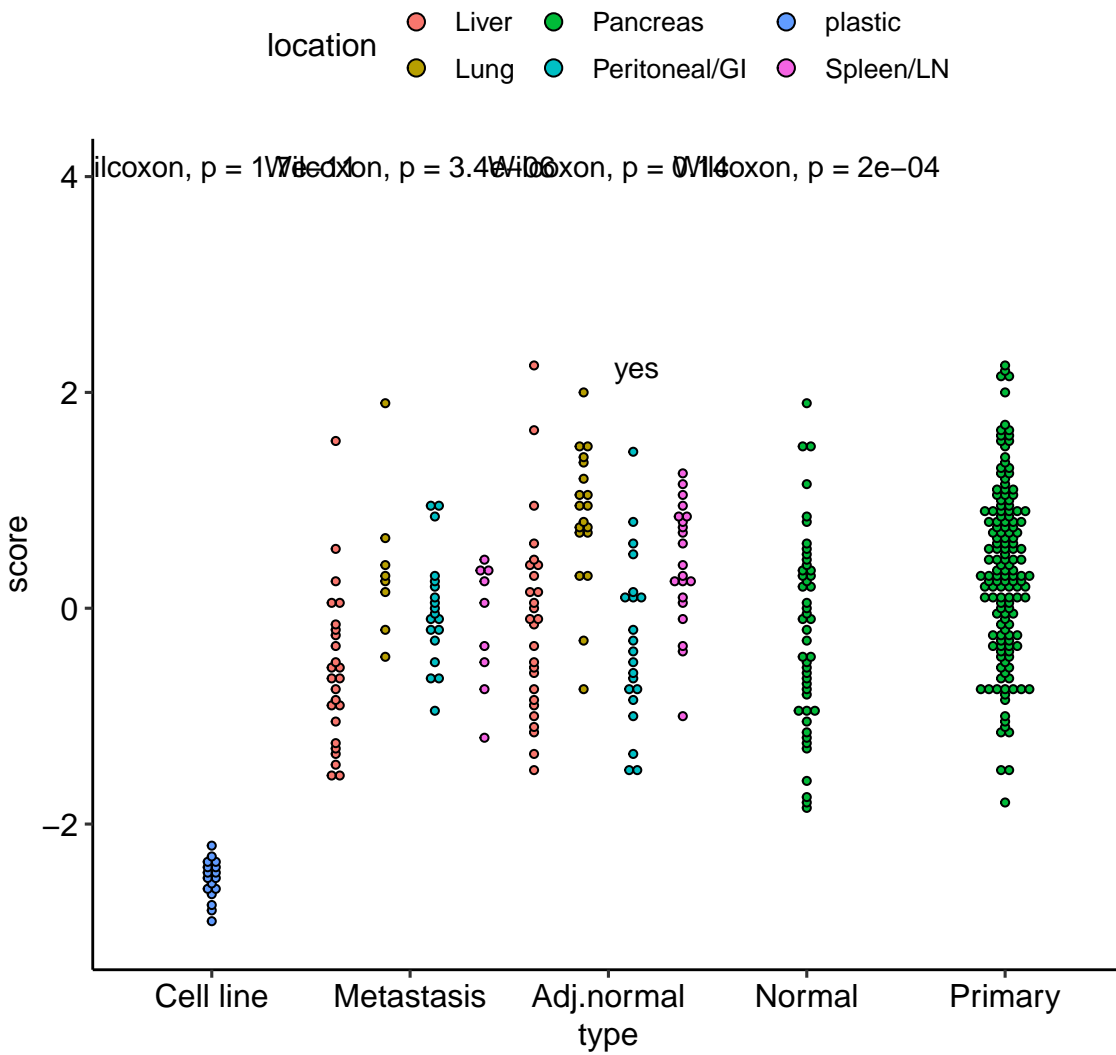


# CIBERSORT.Macrophages.M2 score





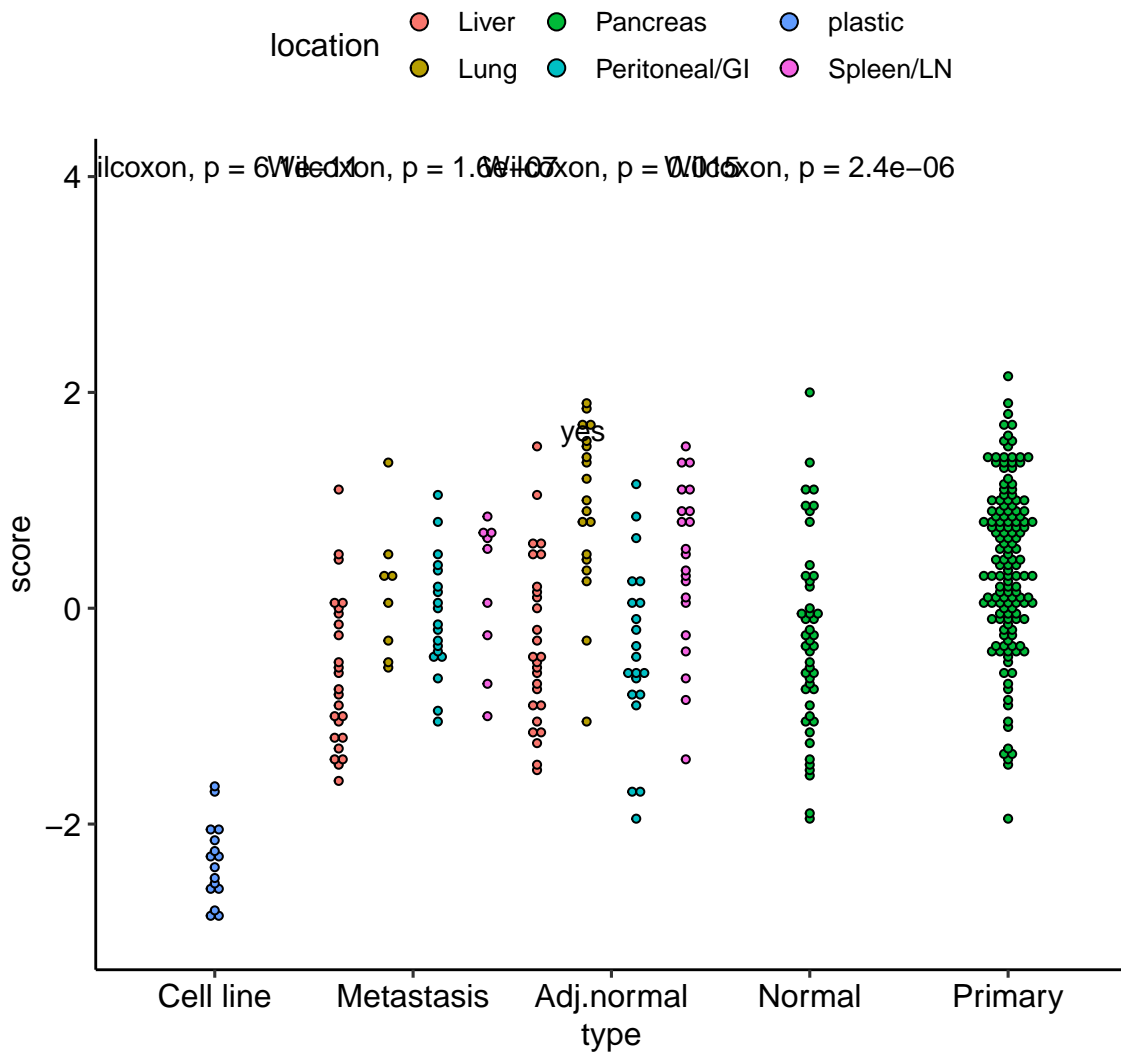
## CIBERSORT.Dendritic.cells.resting score



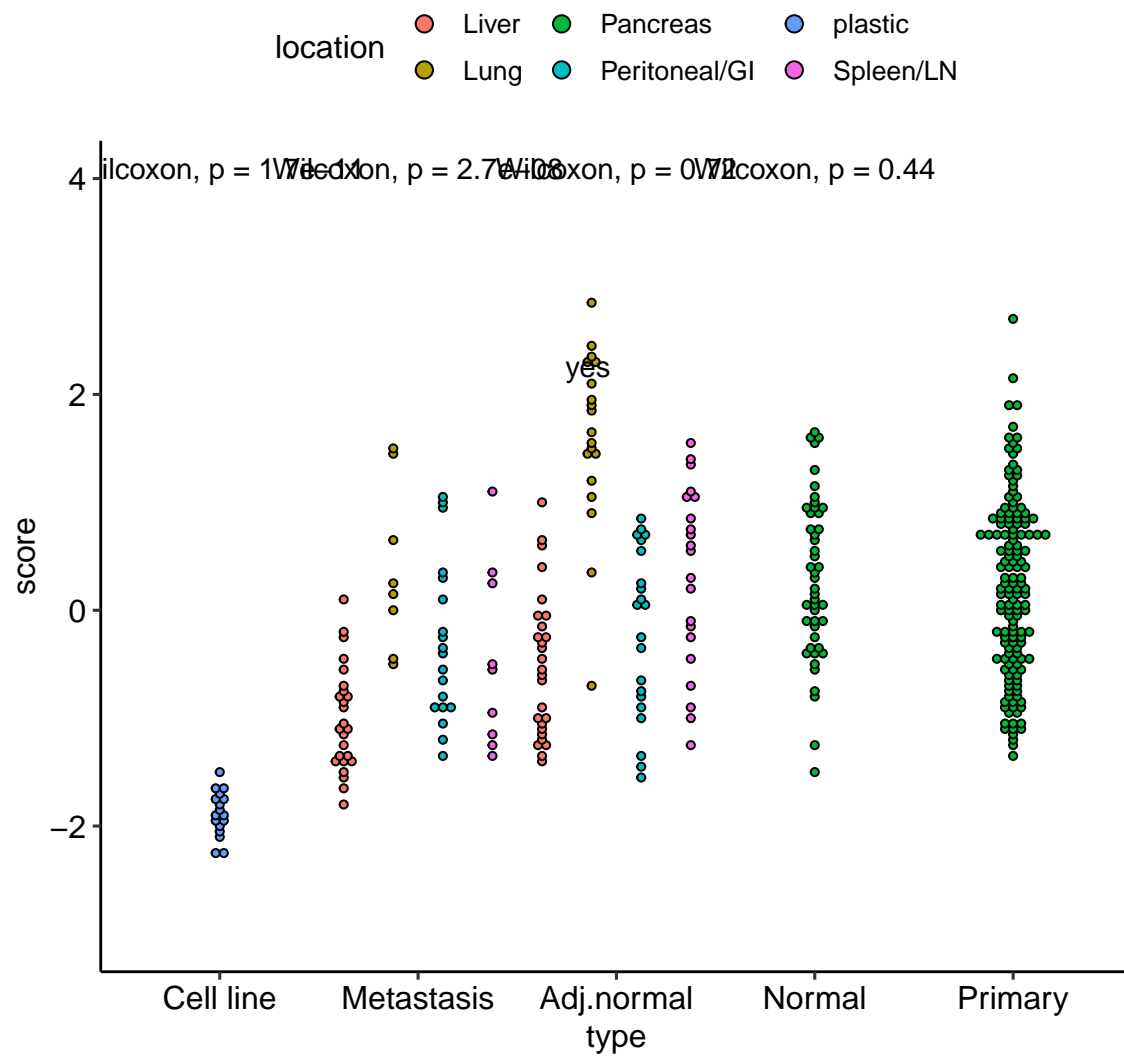
## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

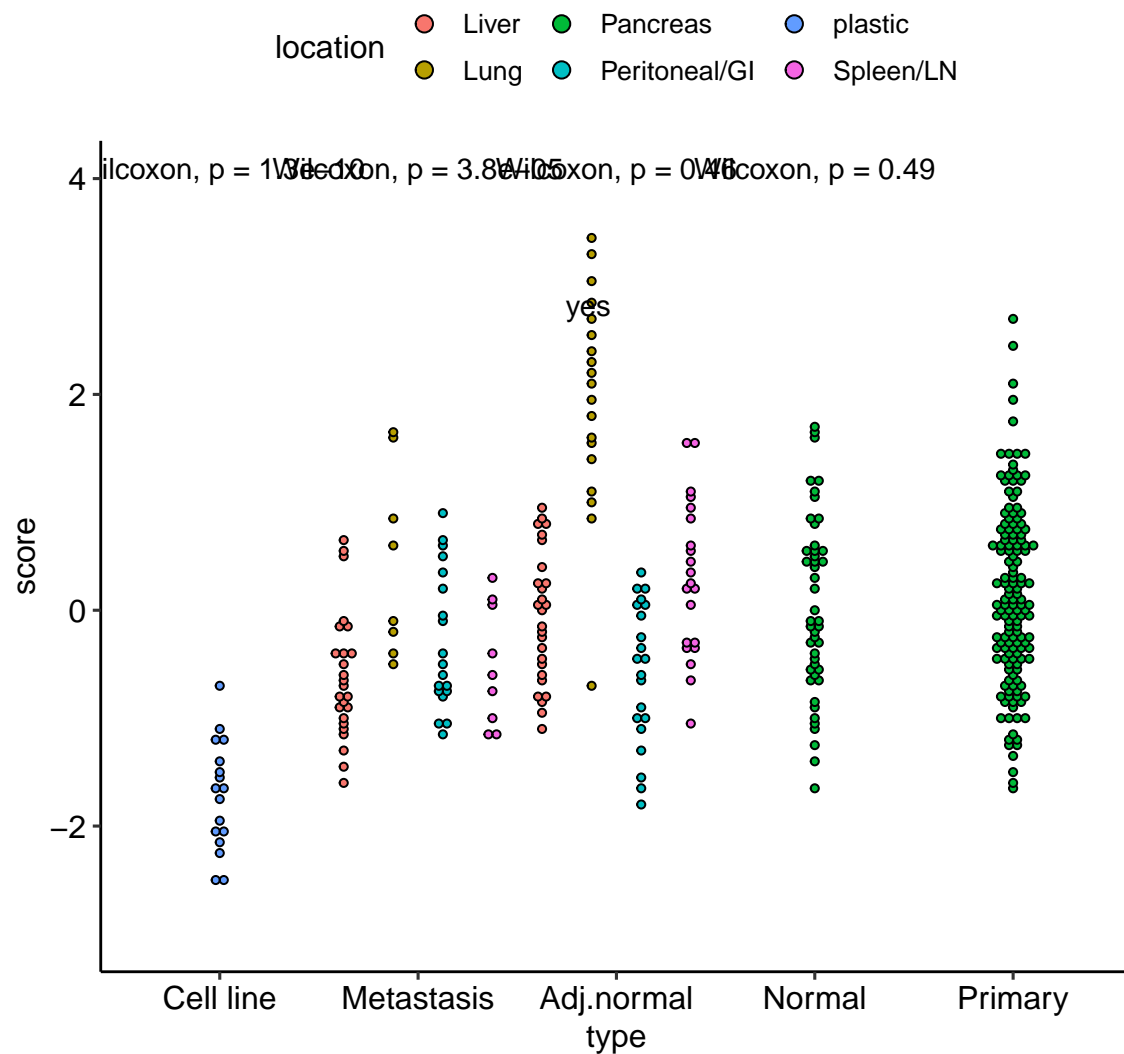
# CIBERSORT.Dendritic.cells.activated score



# CIBERSORT.Mast.cells.resting score



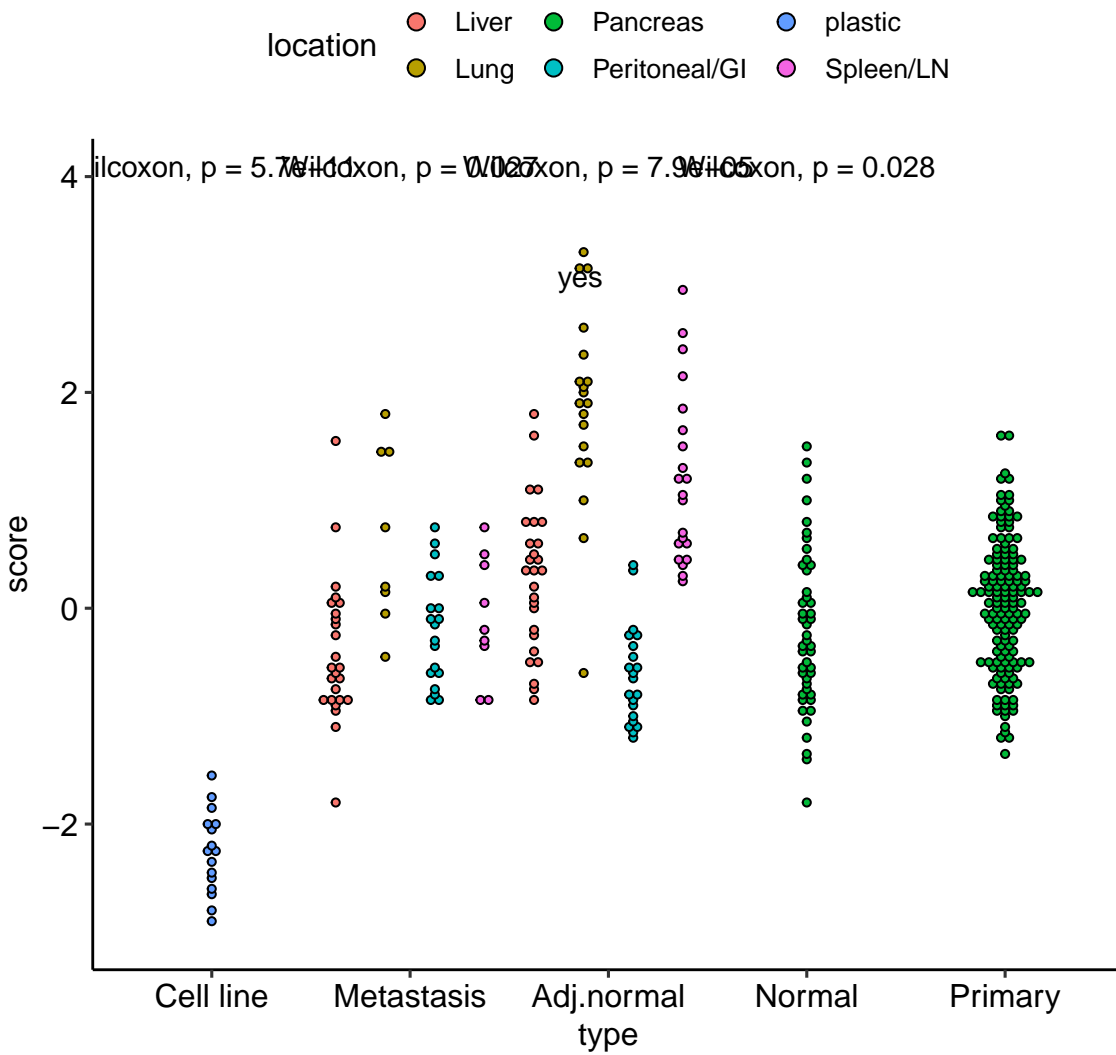
# CIBERSORT.Mast.cells.activated score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

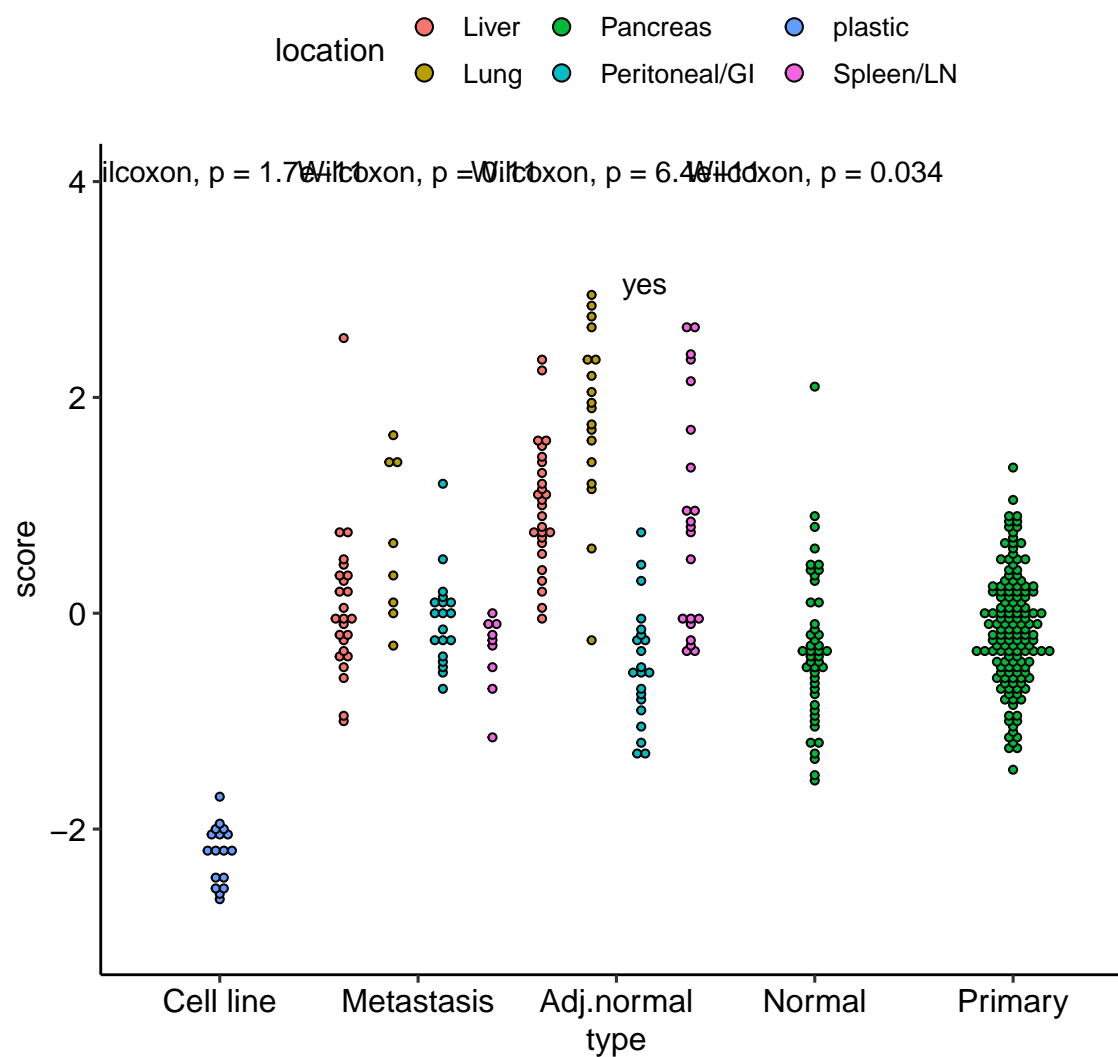
## CIBERSORT.Eosinophils score



## Warning: Removed 1 rows containing non-finite values (stat\_bindot).

## Warning: Removed 1 rows containing non-finite values (stat\_compare\_means).

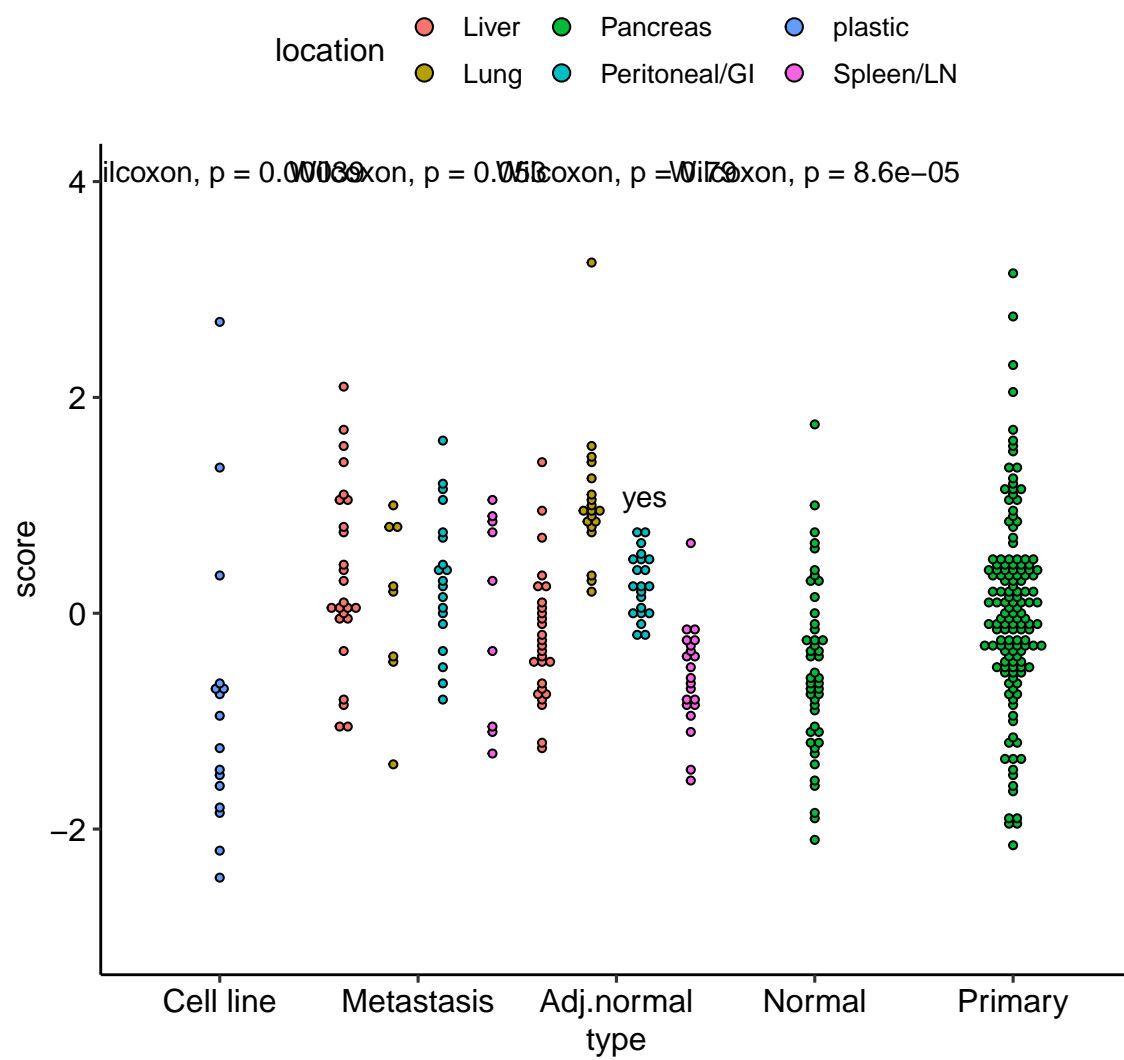
# CIBERSORT.Neutrophils score



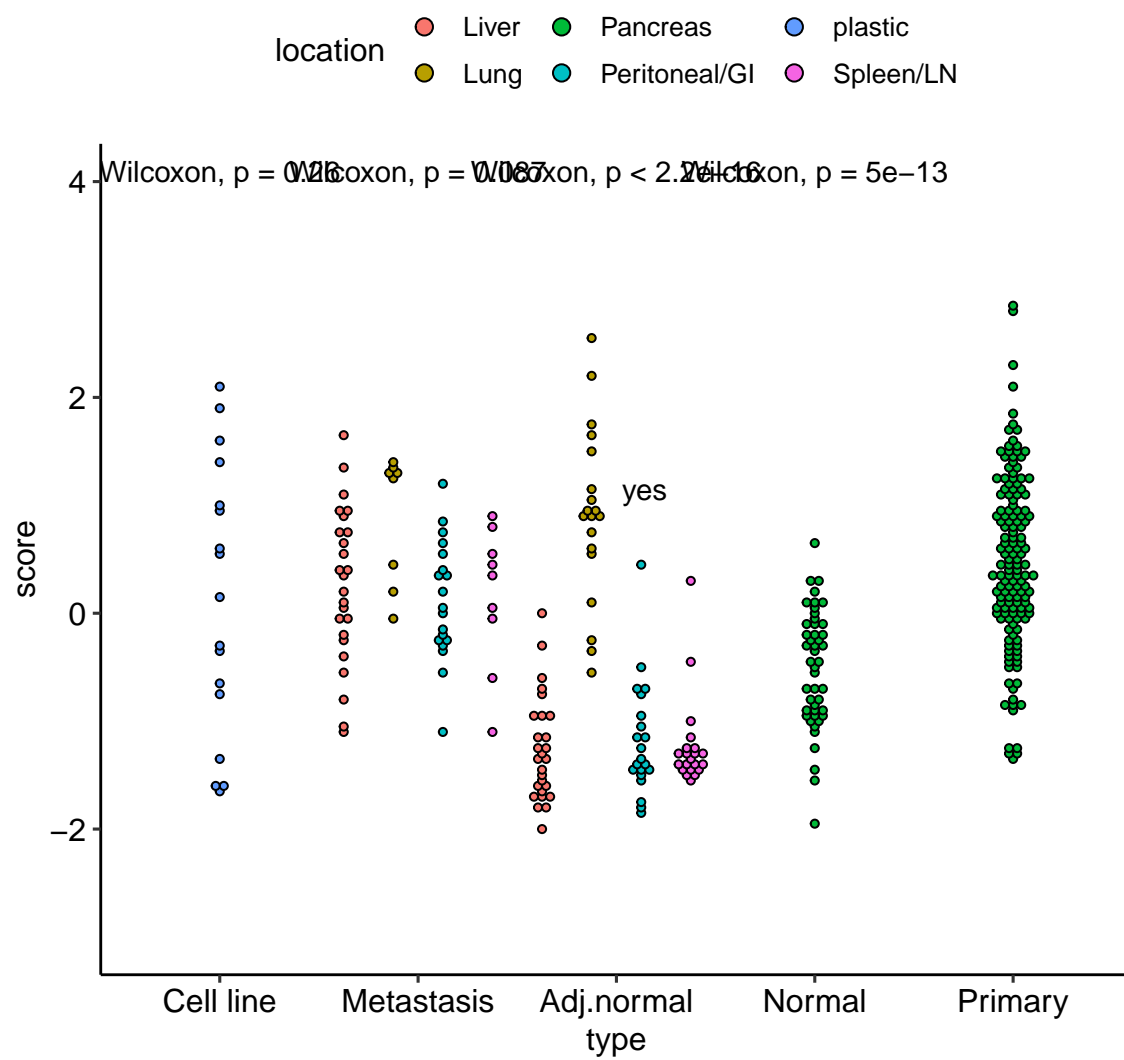
## Warning: Removed 4 rows containing non-finite values (stat\_bindot).

## Warning: Removed 4 rows containing non-finite values (stat\_compare\_means).

# Notta.BasalA score

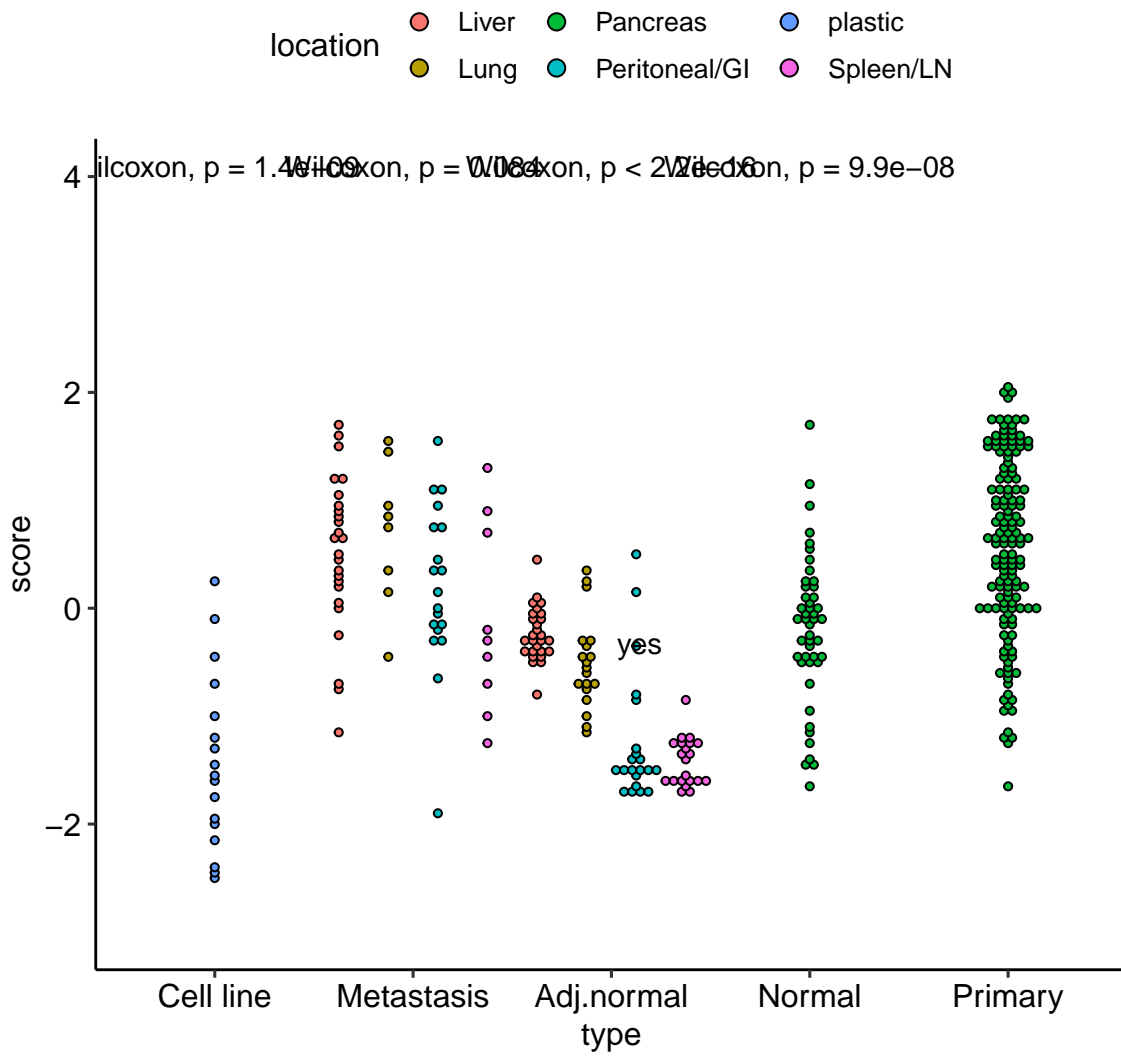


## Notta.BasalB score

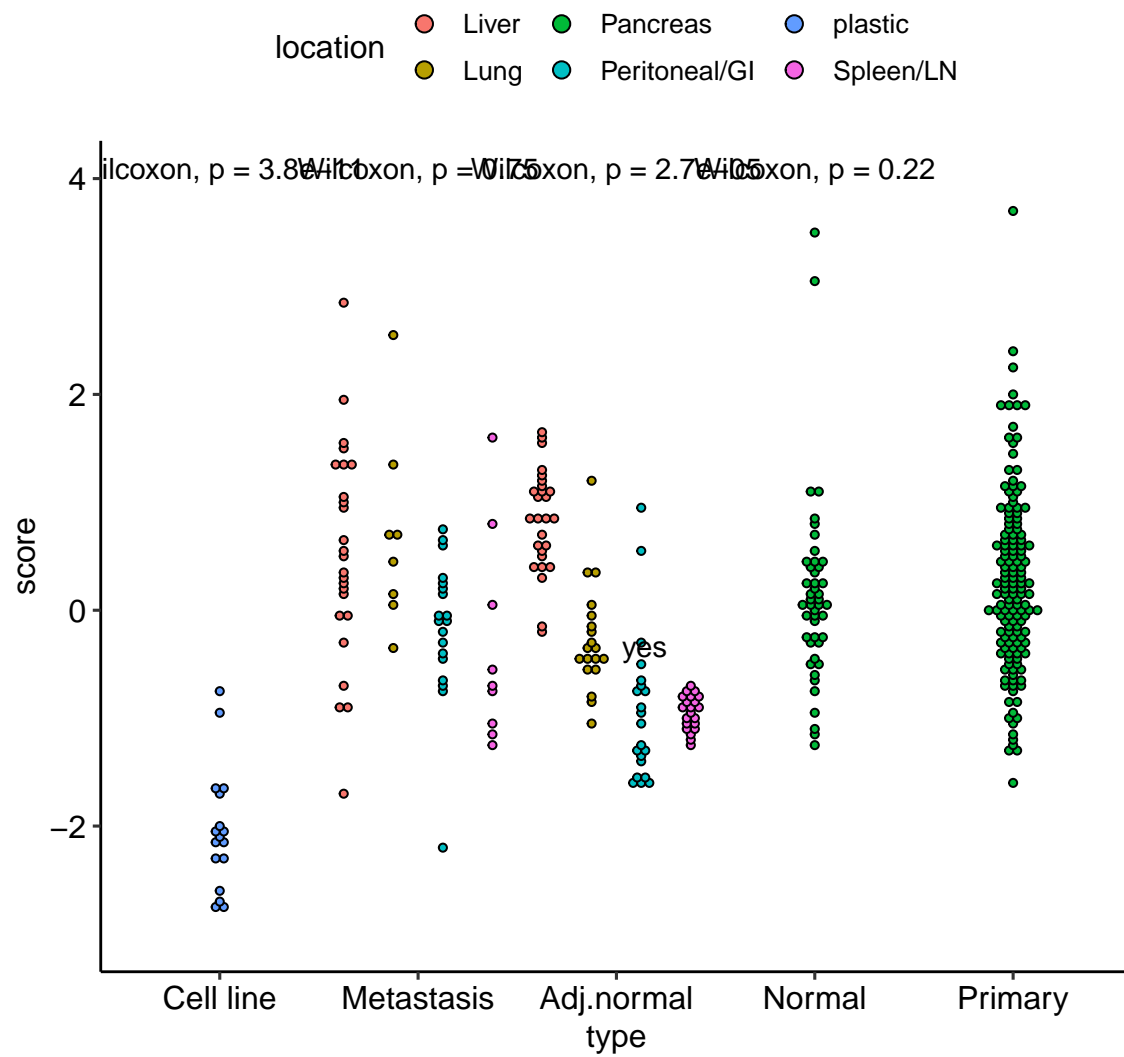




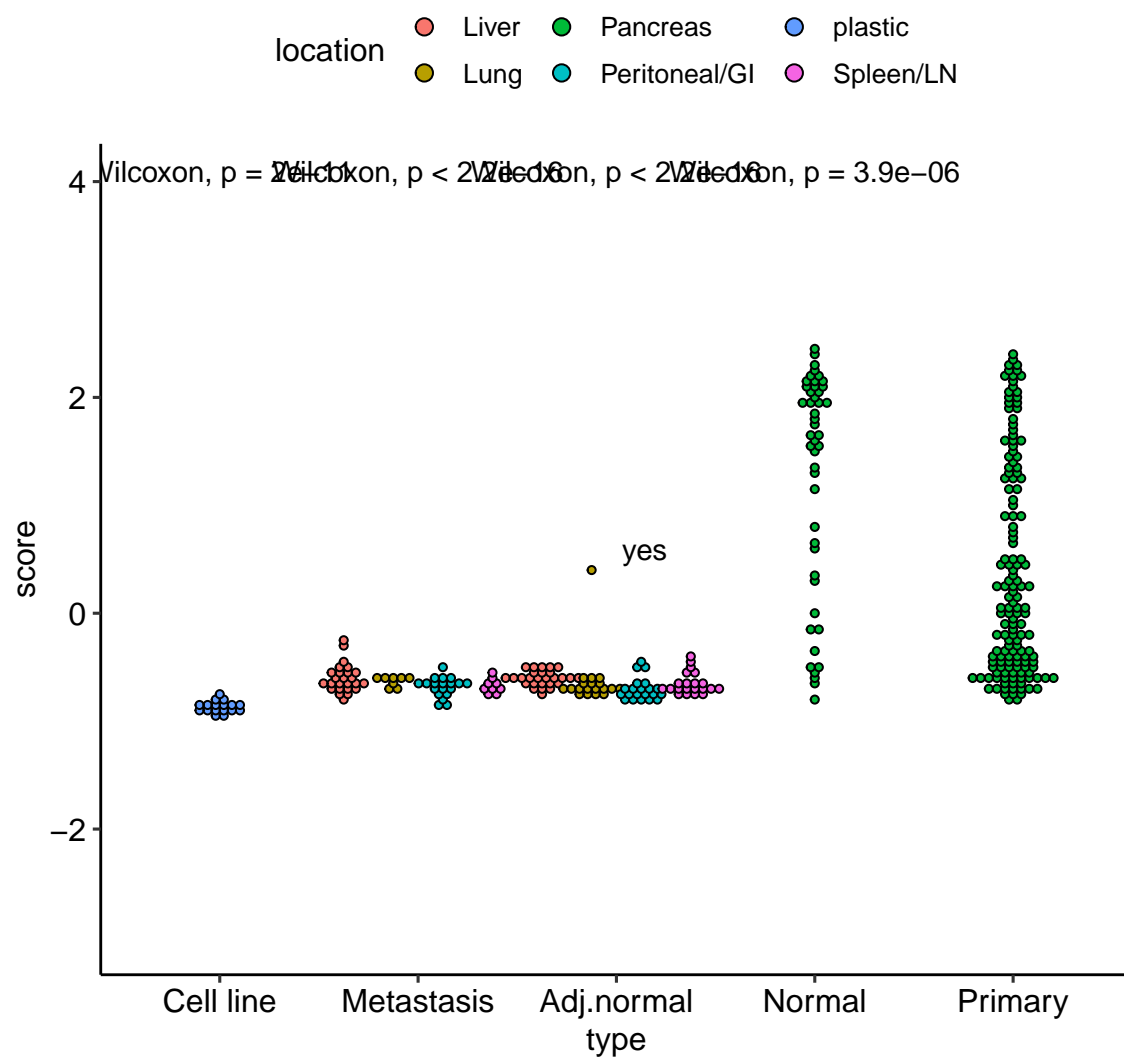
# Notta.ClassicalA score



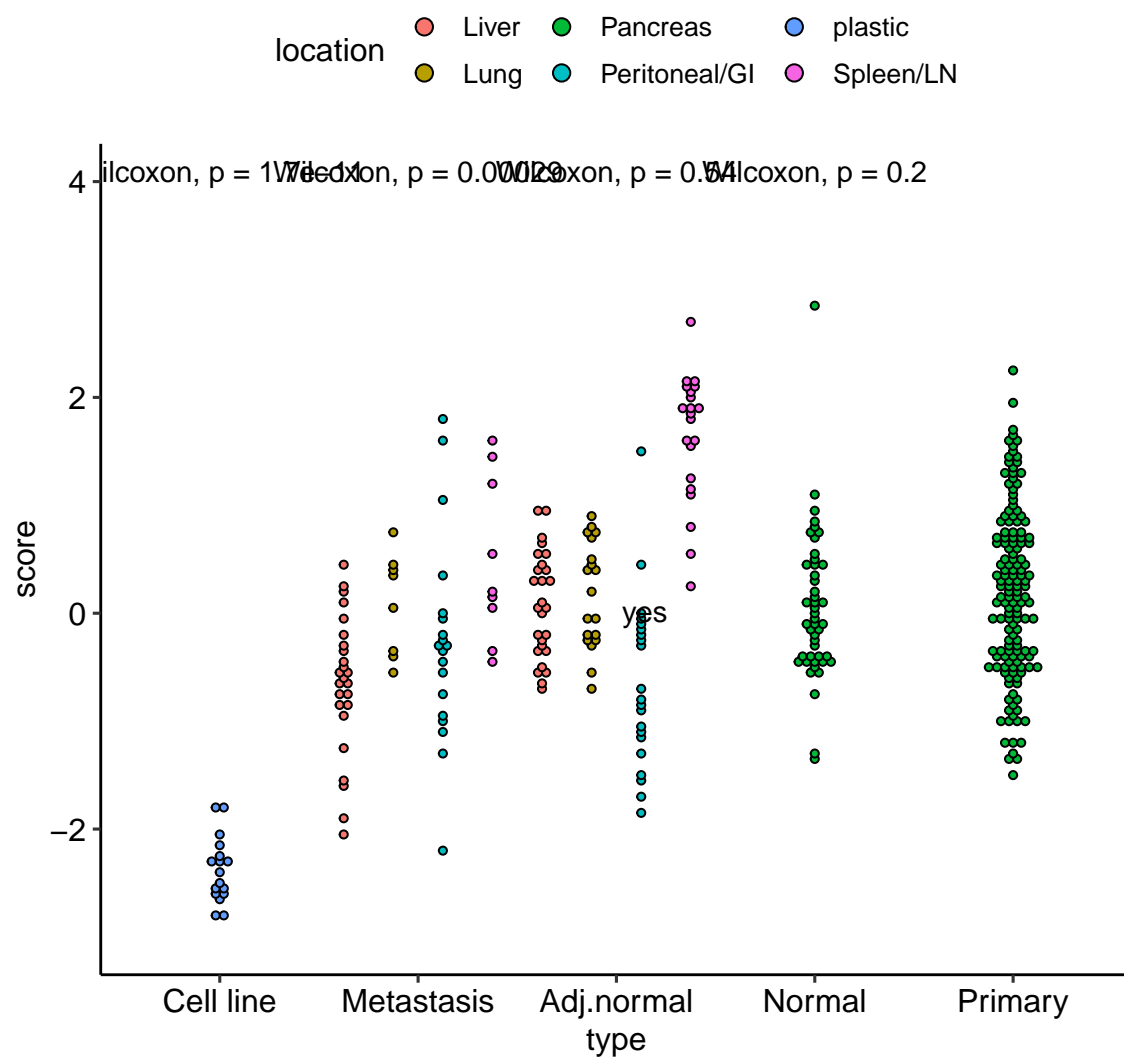
# Notta.ClassicalB score



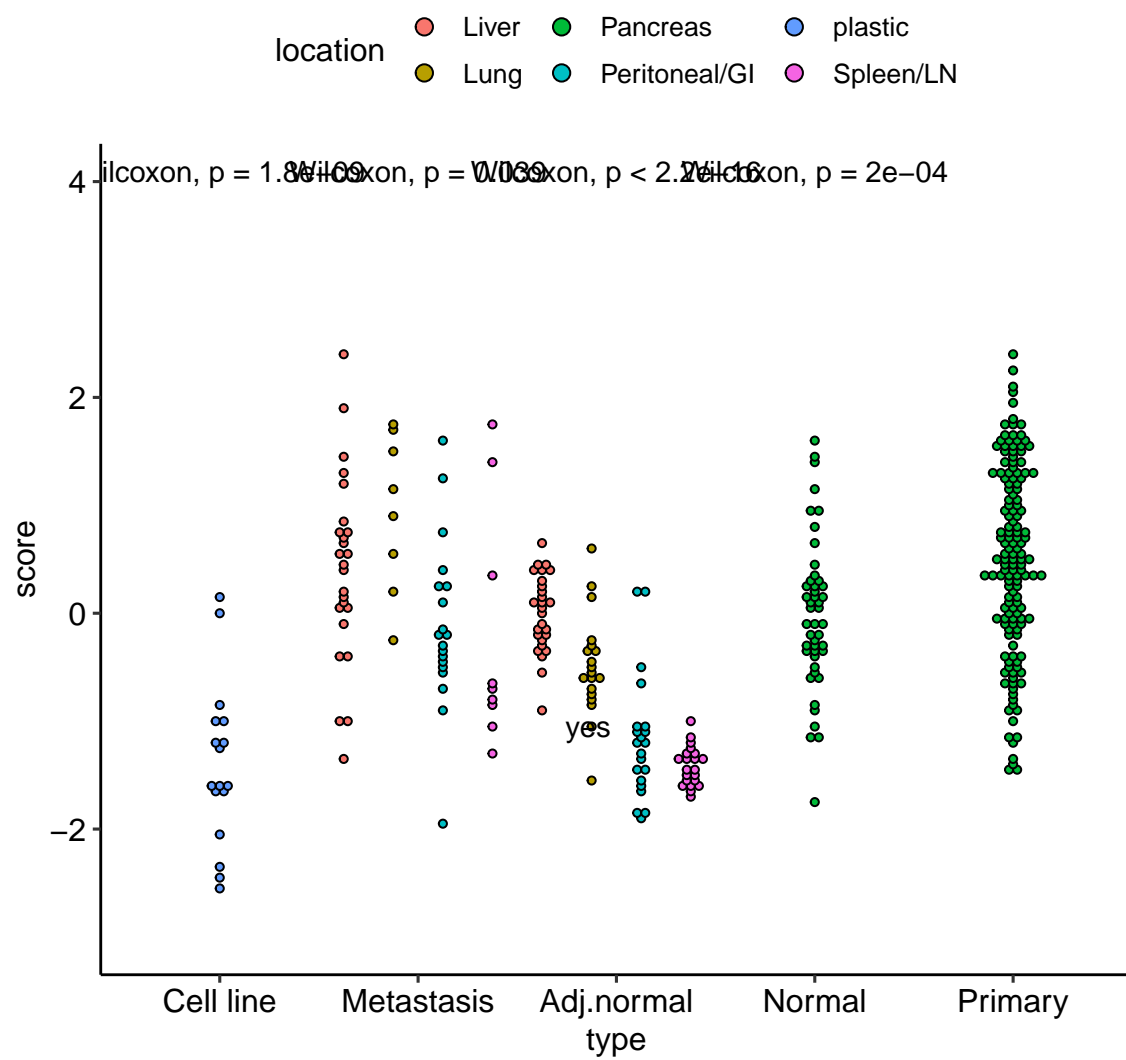
## ADEX\_unique score



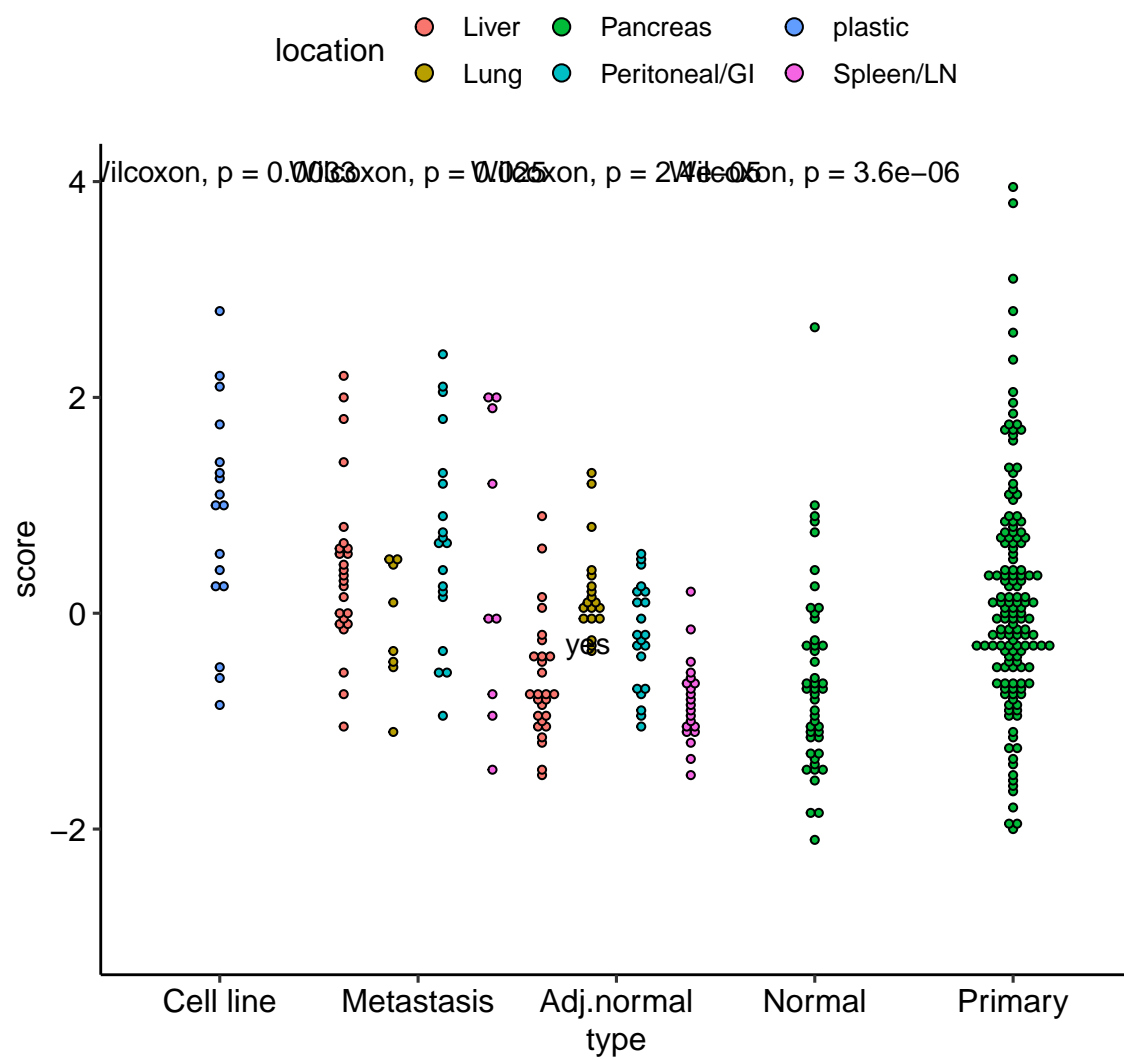
# Immunogenic\_unique score



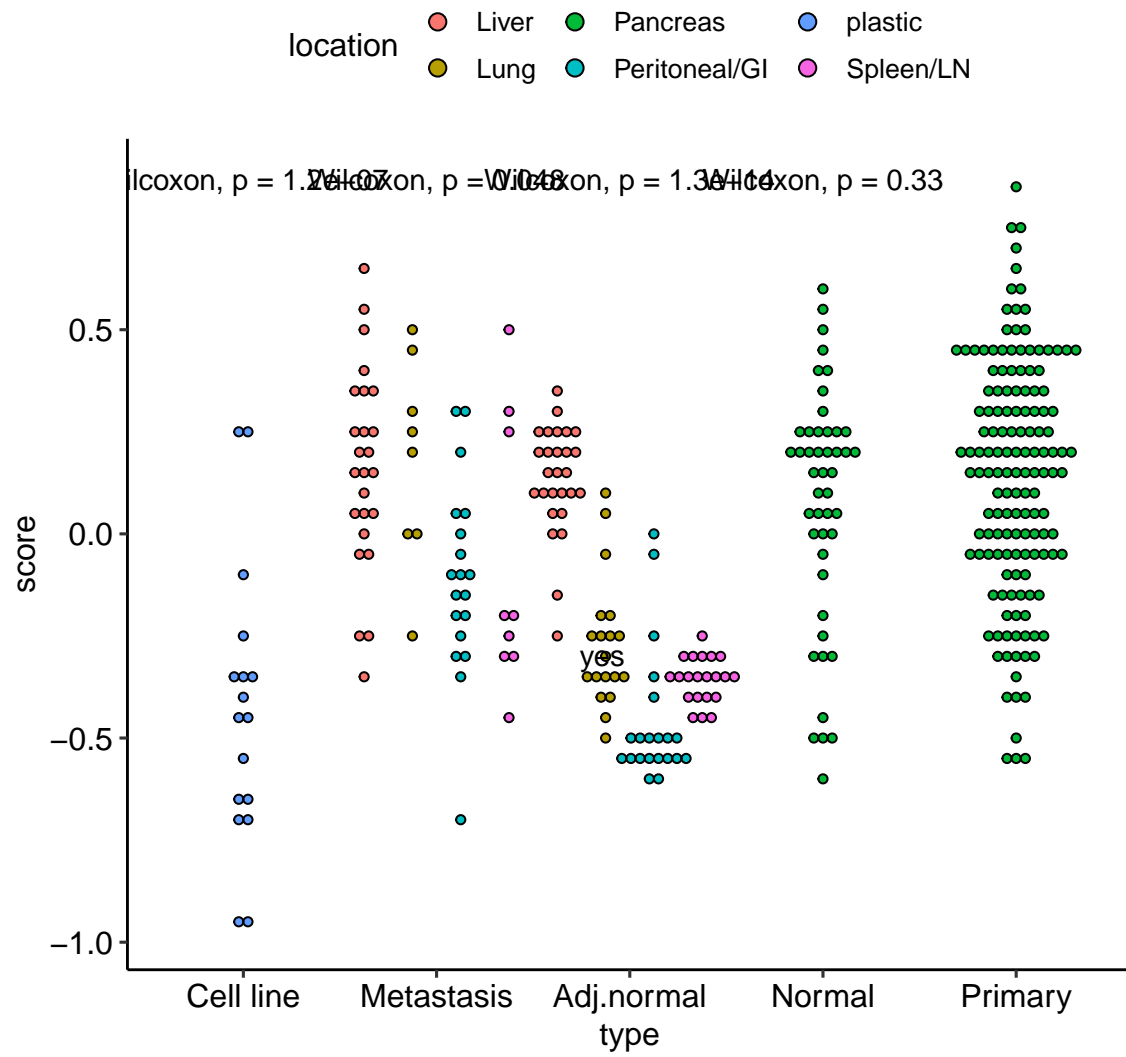
# Progenitor\_unique score



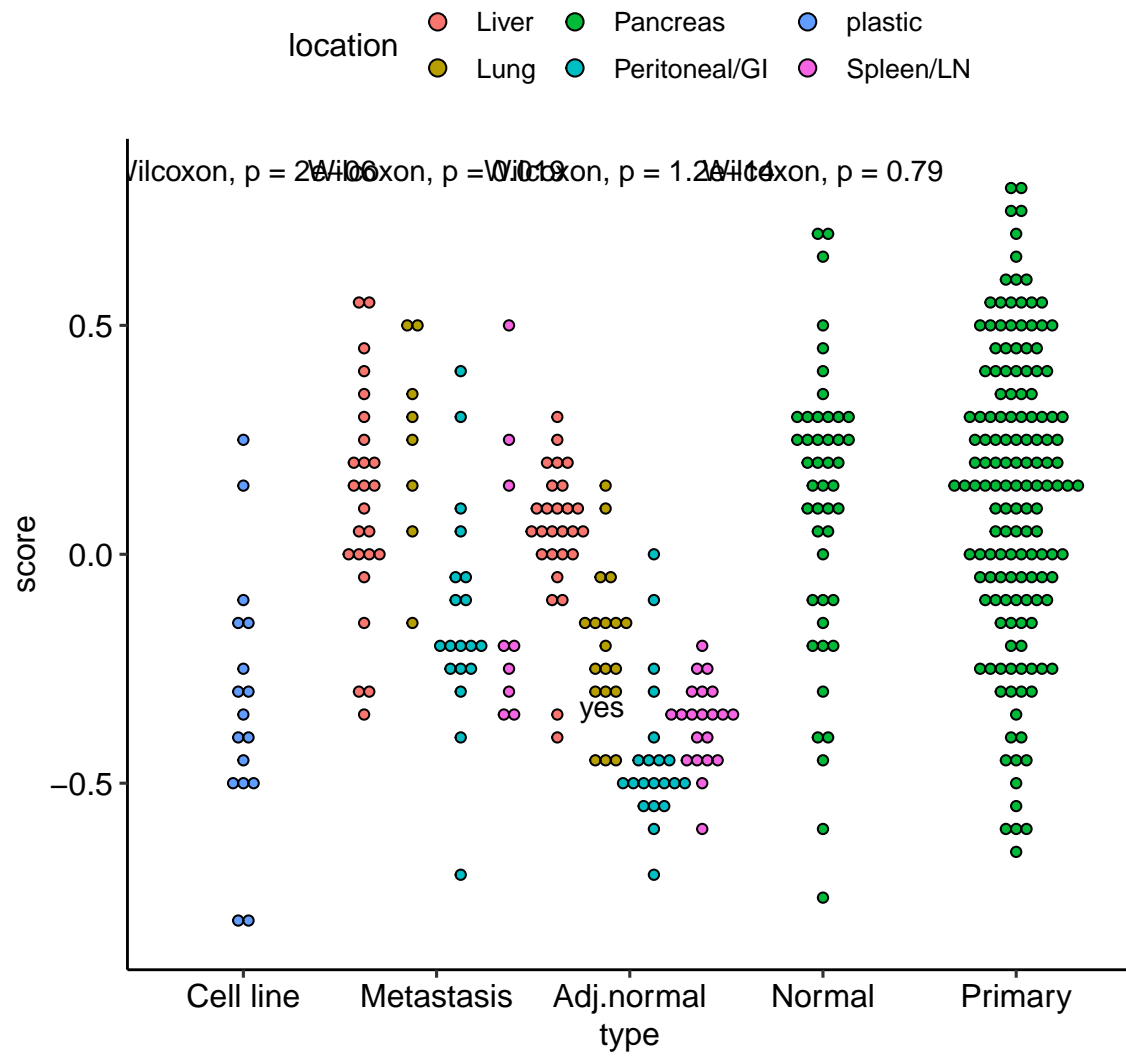
## Squamous\_unique score



# molgrad\_PDX score

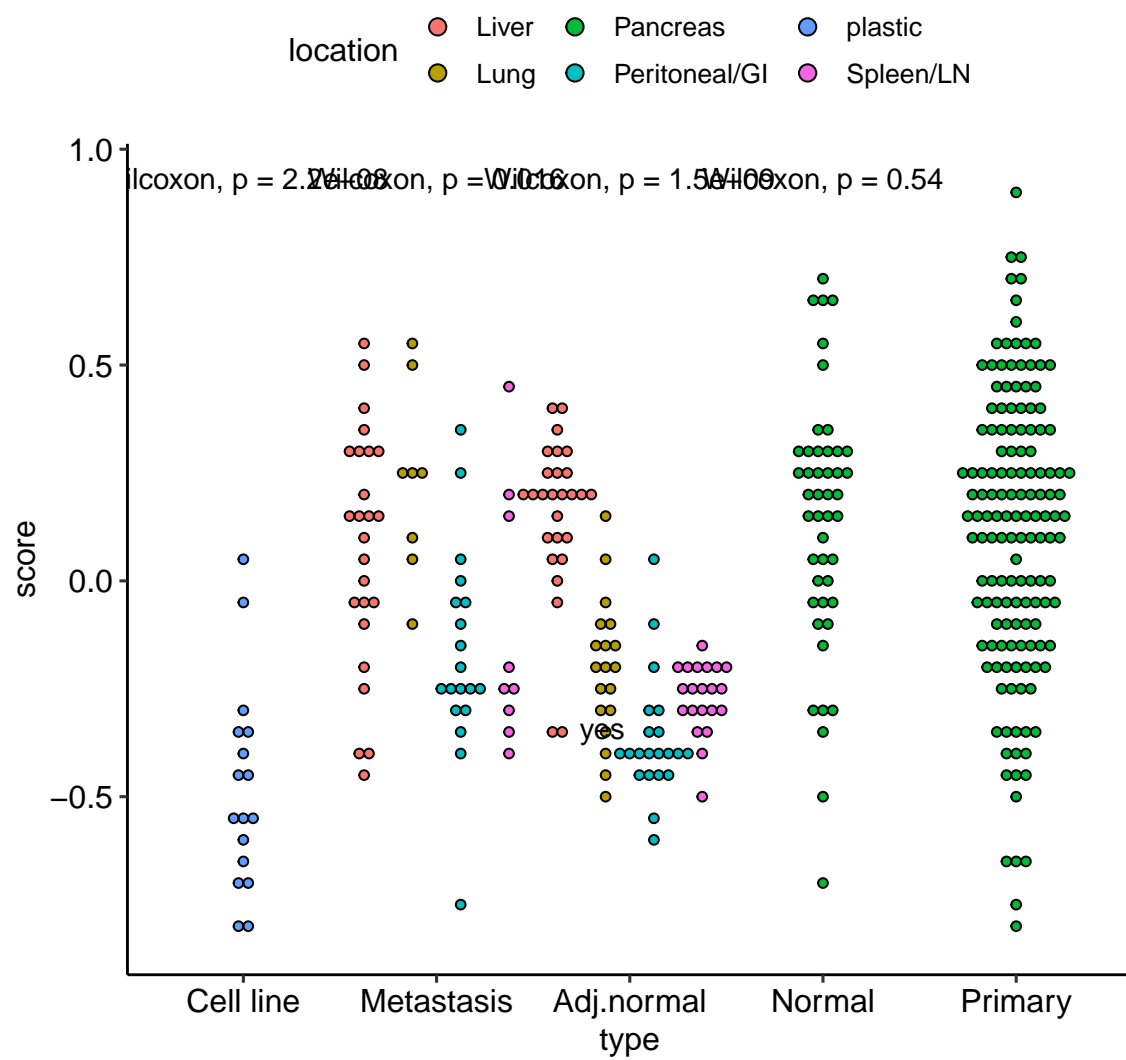


# molgrad\_Puleo score

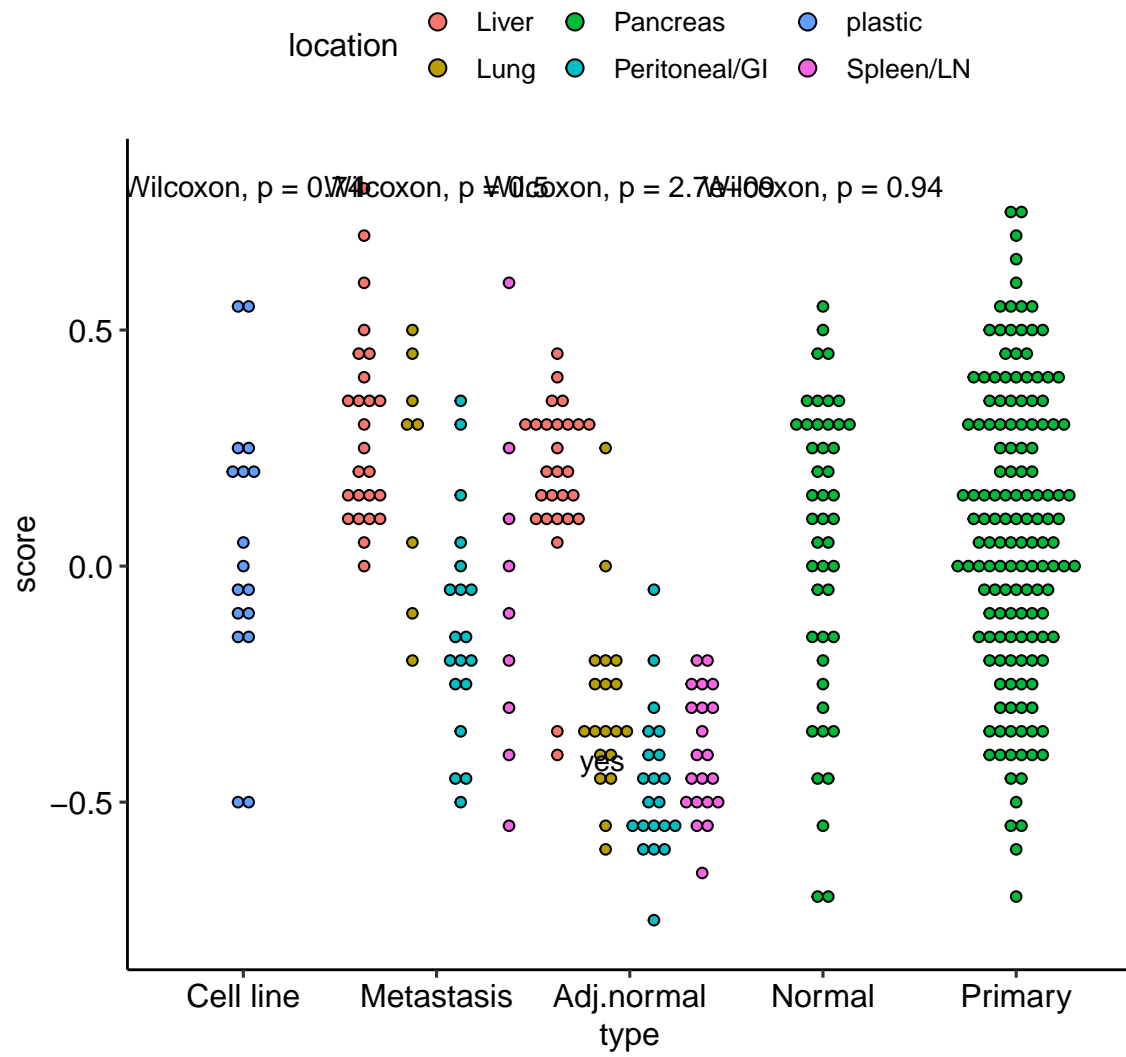




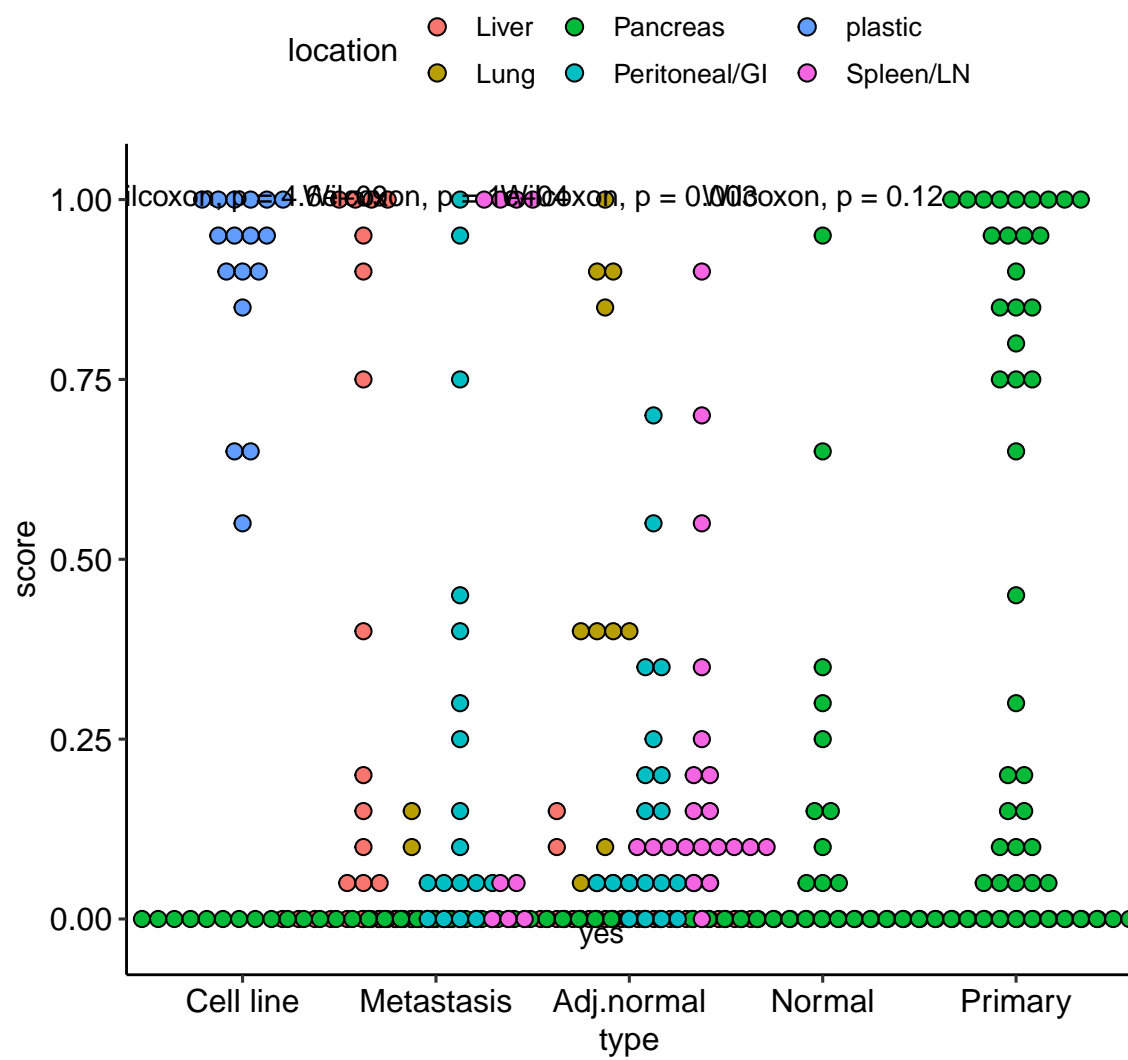
# molgrad\_ICGArray score



# molgrad\_ICGCrnaseq score



# purIST score



# molgrad\_scaled score

