

Number.of.Reads

9e+08

6e+08

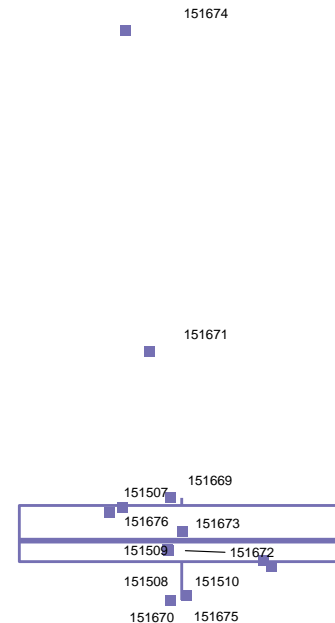
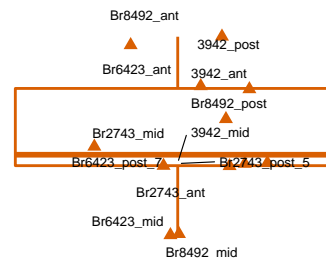
3e+08

VisiumIF

spatialDLPFC

spatialLIBD

study



Mean.Reads.per.Spot

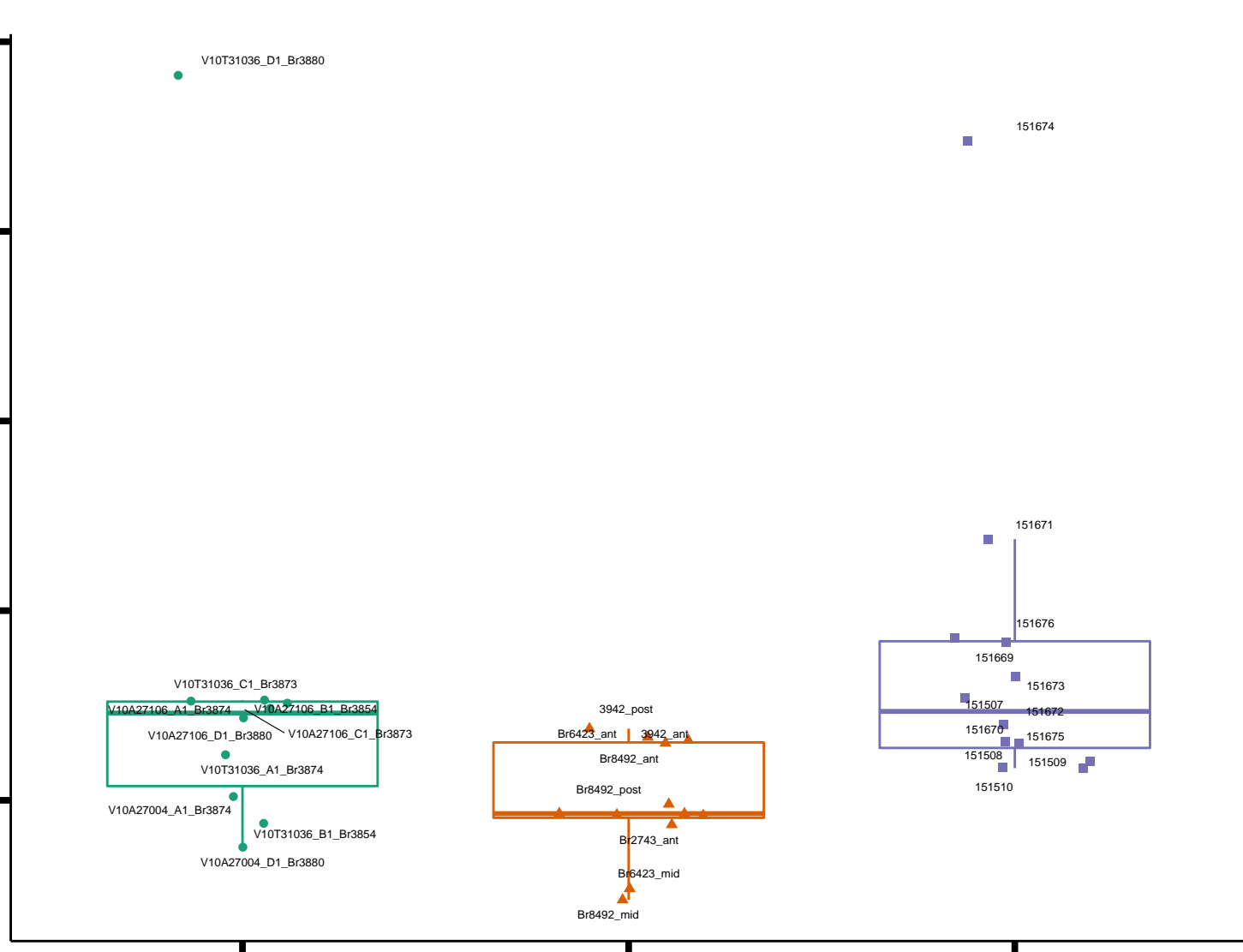
250000
200000
150000
100000
50000

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spatialLIBD

study



Median.Genes.per.Spot

2500

2000

1500

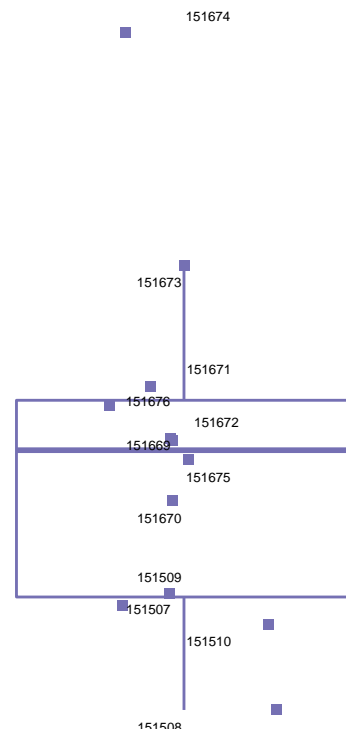
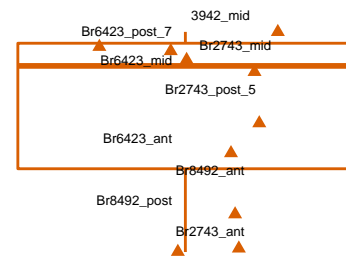
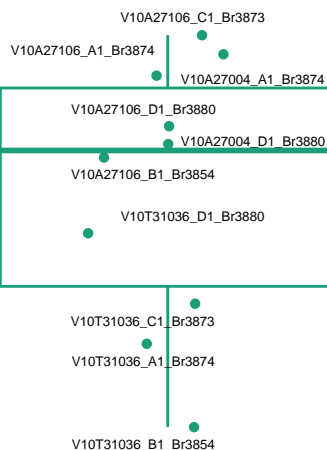
1000

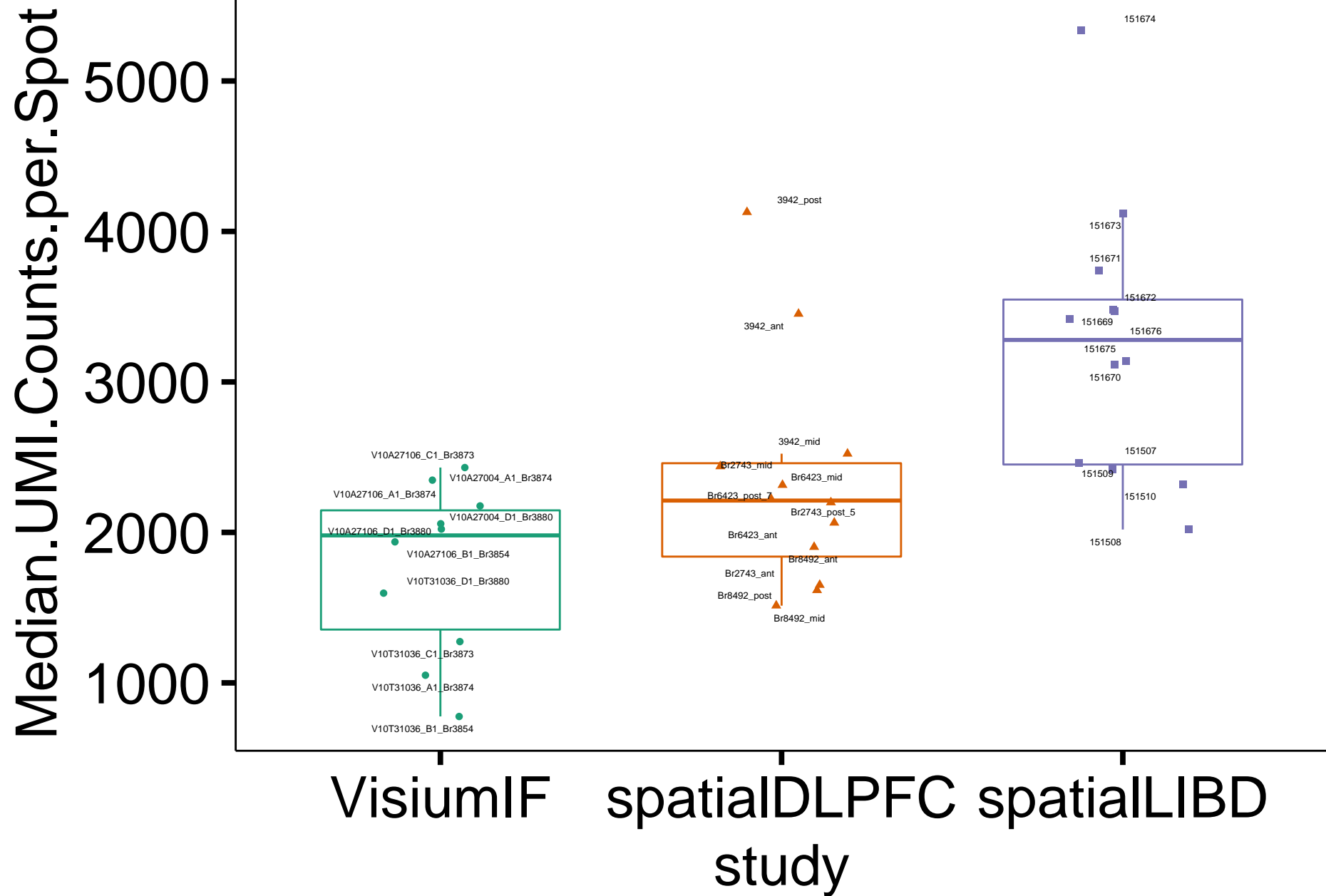
500

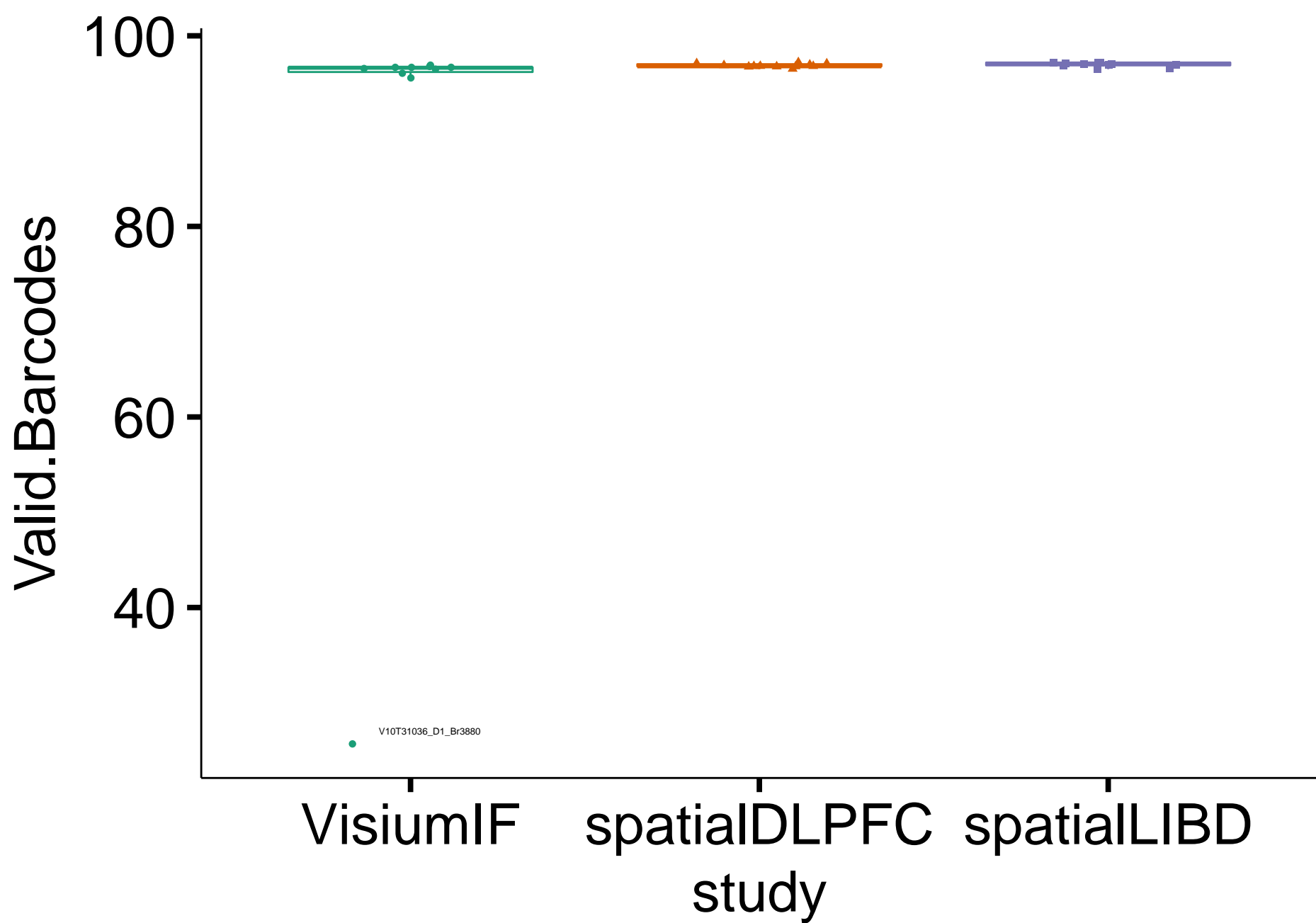
VisiumIF

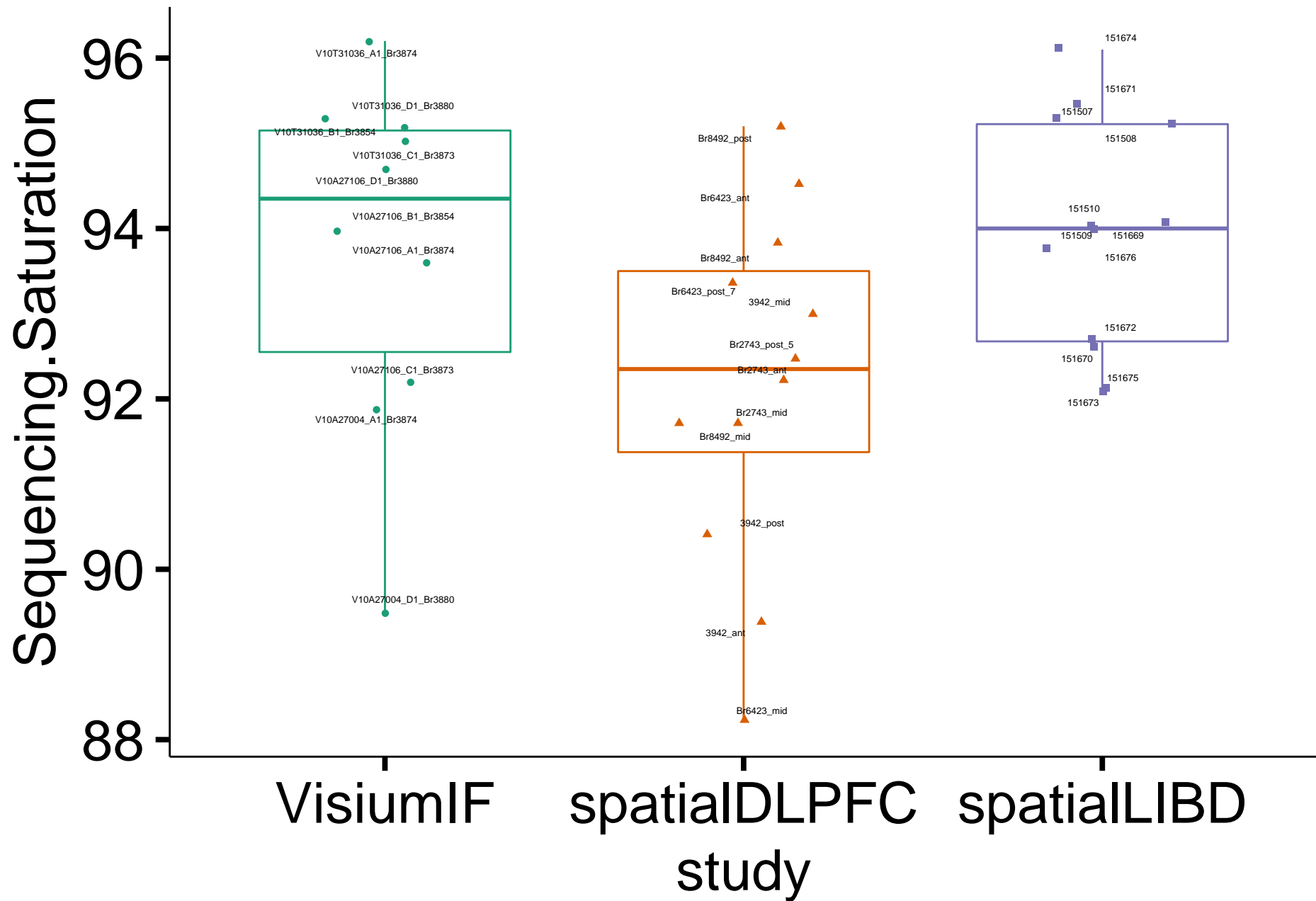
spatialDLPFC
study

spatialLIBD









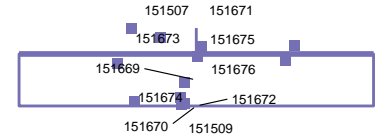
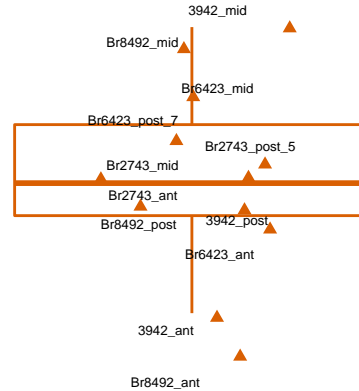
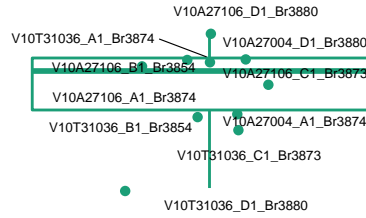
Q30.Bases.in.Barcode

97
96
95
94

VisiumIF

spatialDLPFC
study

spatialLIBD



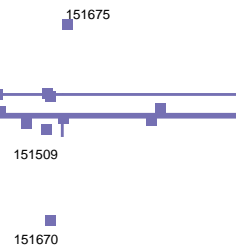
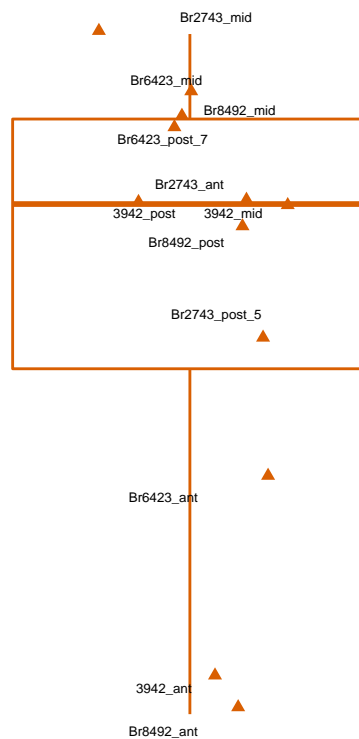
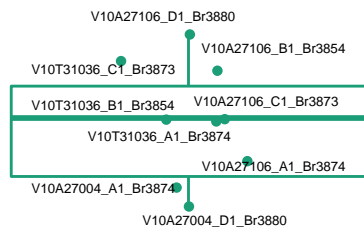
Q30.Bases.in.RNA.Read

95
94
93
92
91

VisiumIF

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study

spatialLIBD



Q30.Bases.in.UMI

97

96

95

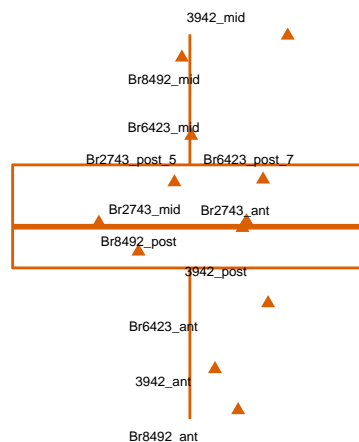
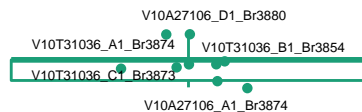
94

VisiumIF

spatialDLPFC

spatialLIBD

study



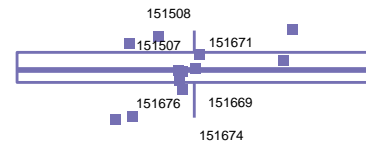
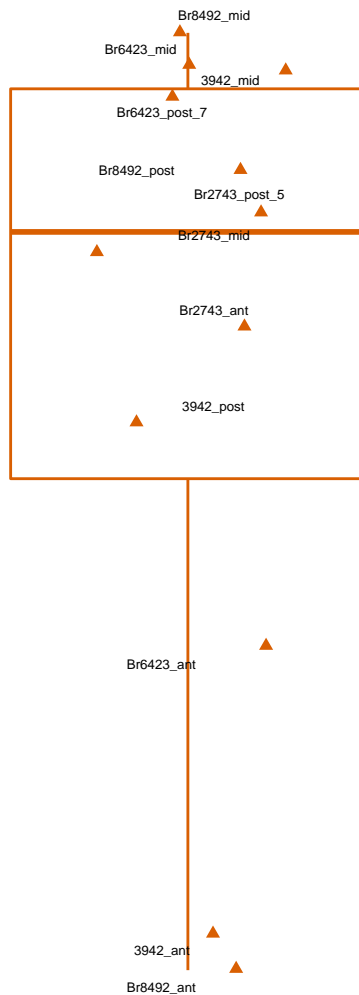
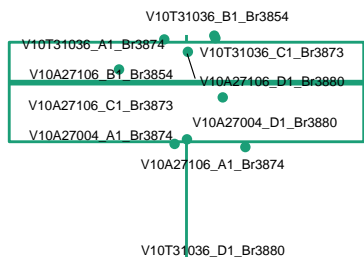
Reads.Mapped.to.Genome

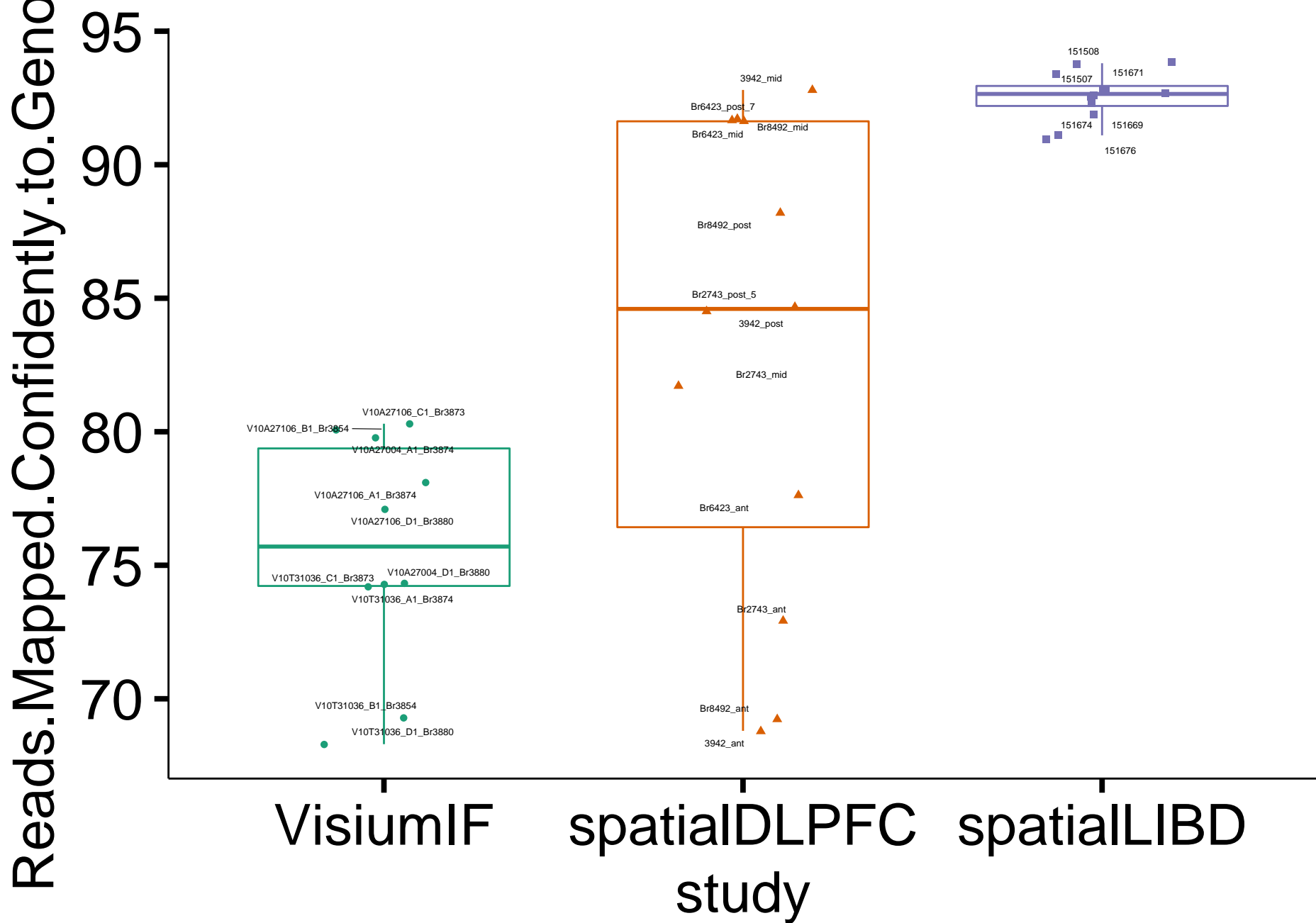
95
90
85
80
75

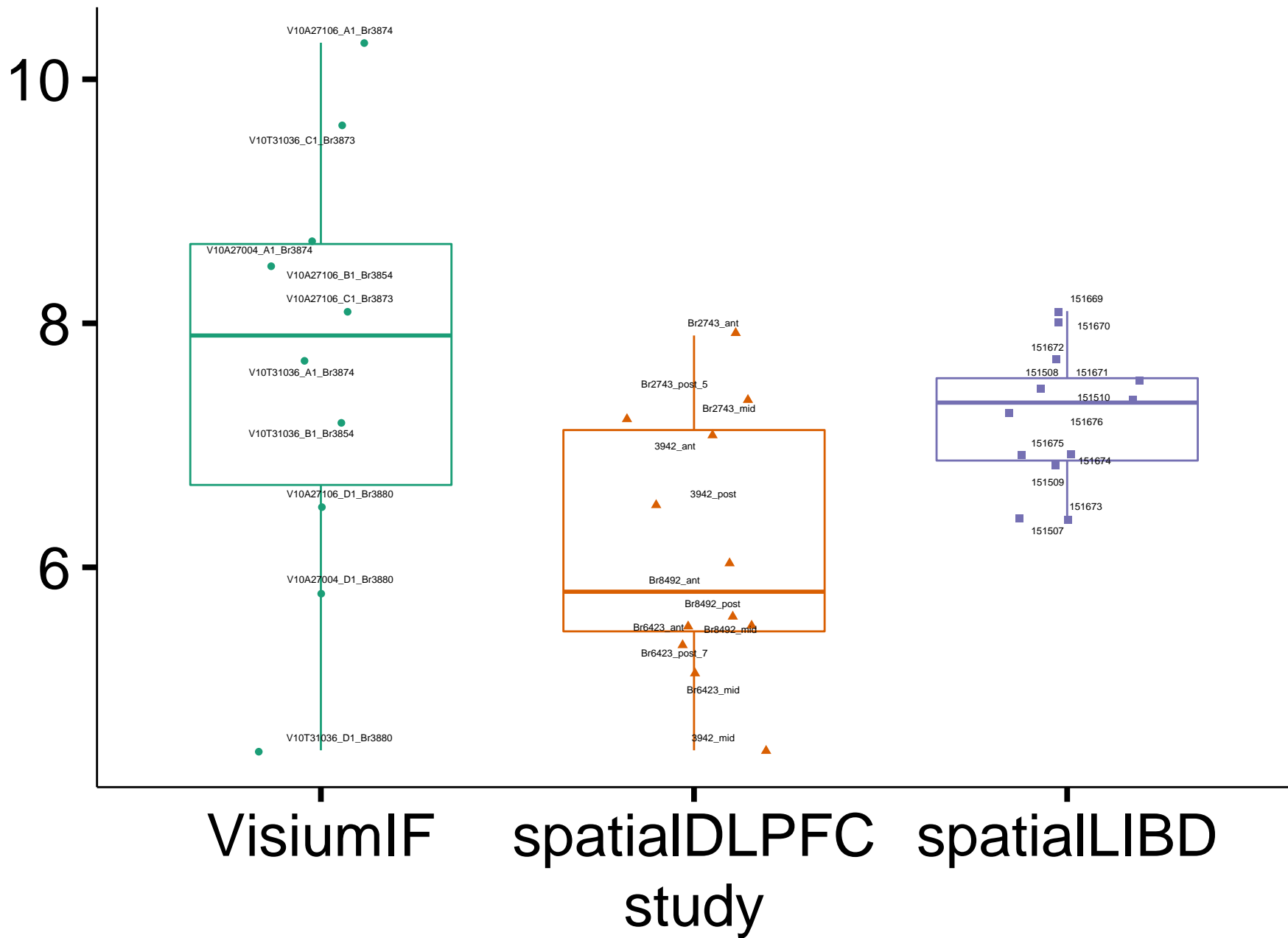
VisiumIF

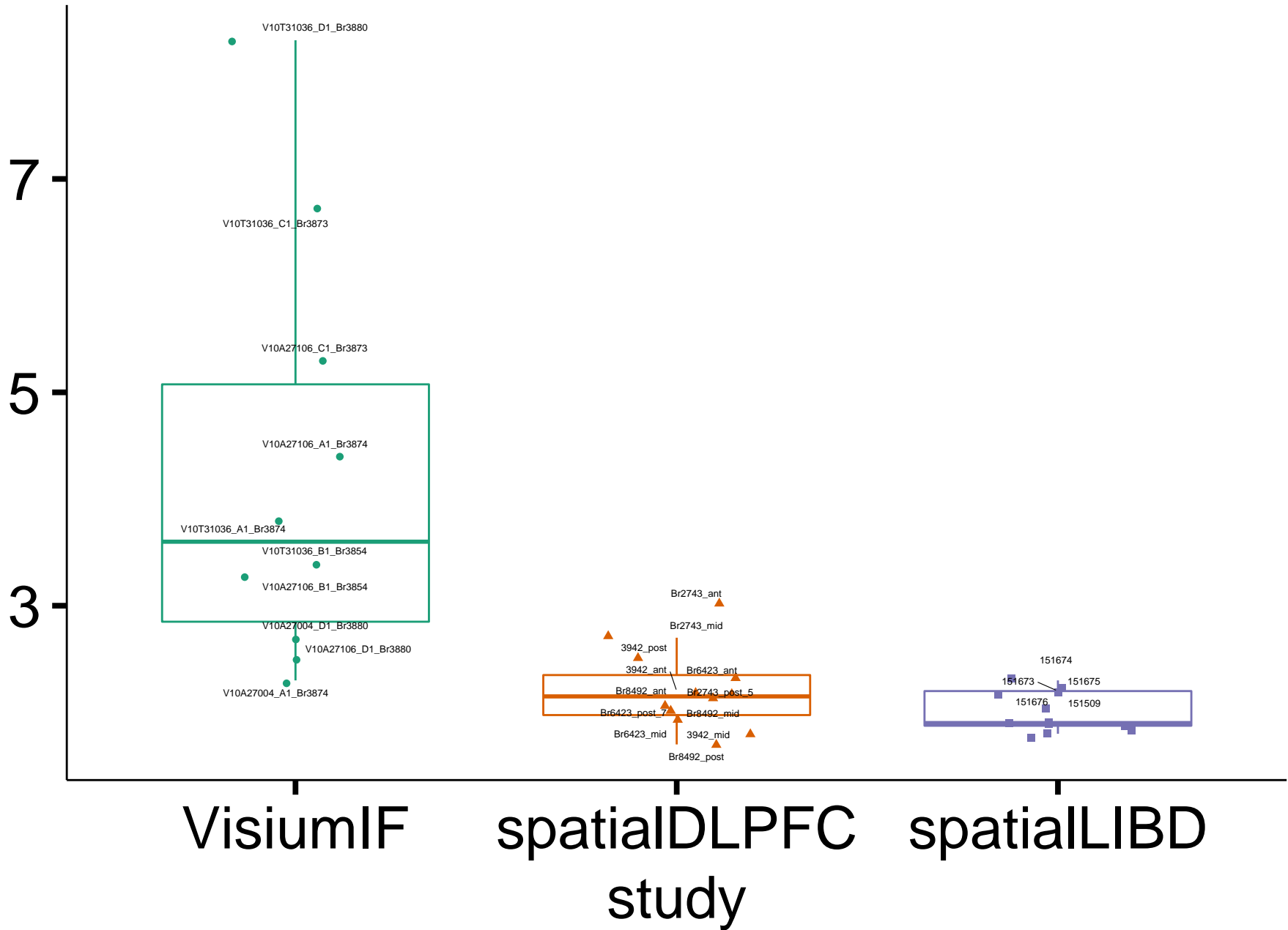
spatialDLPFC
study

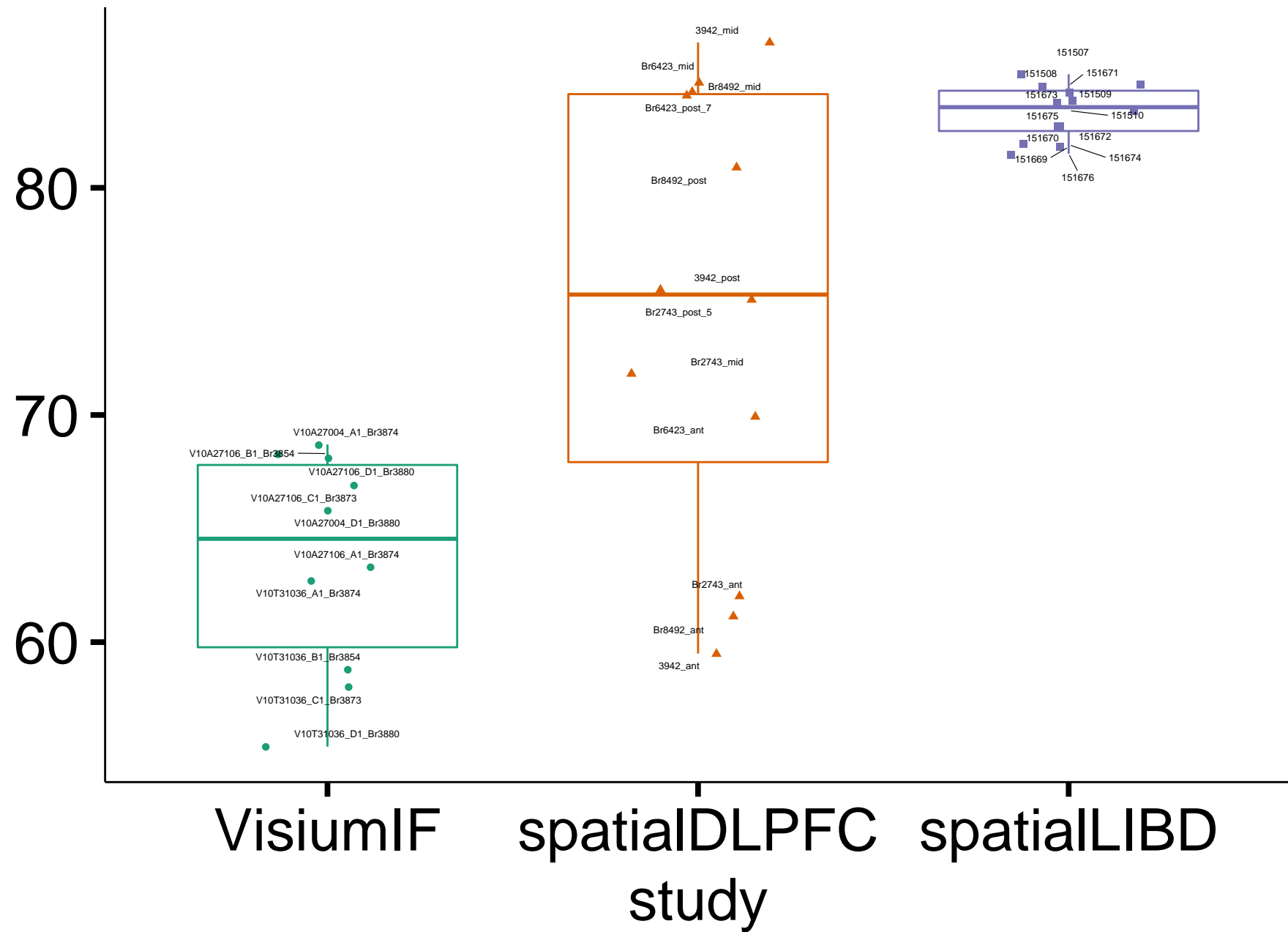
spatialLIBD



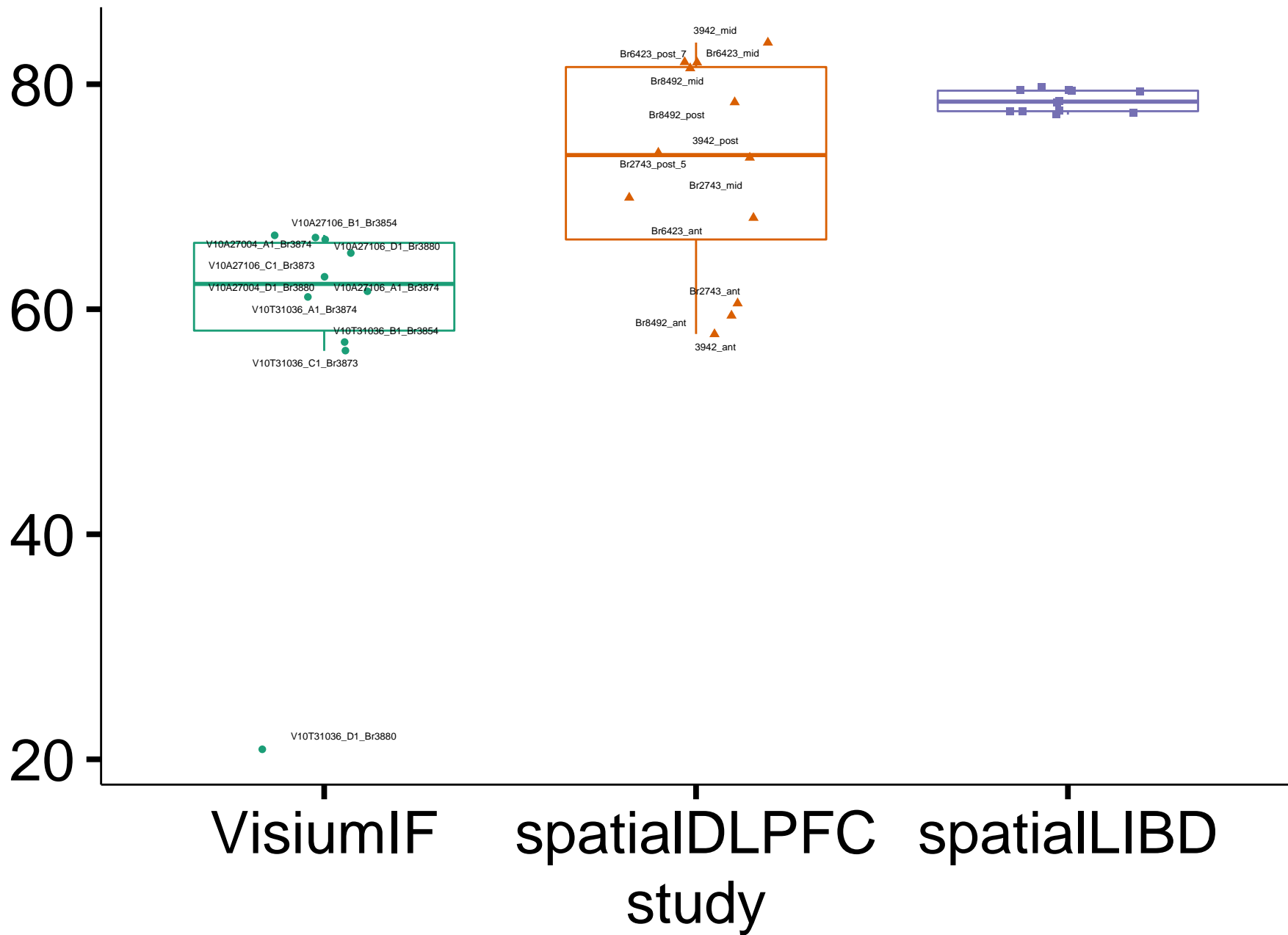


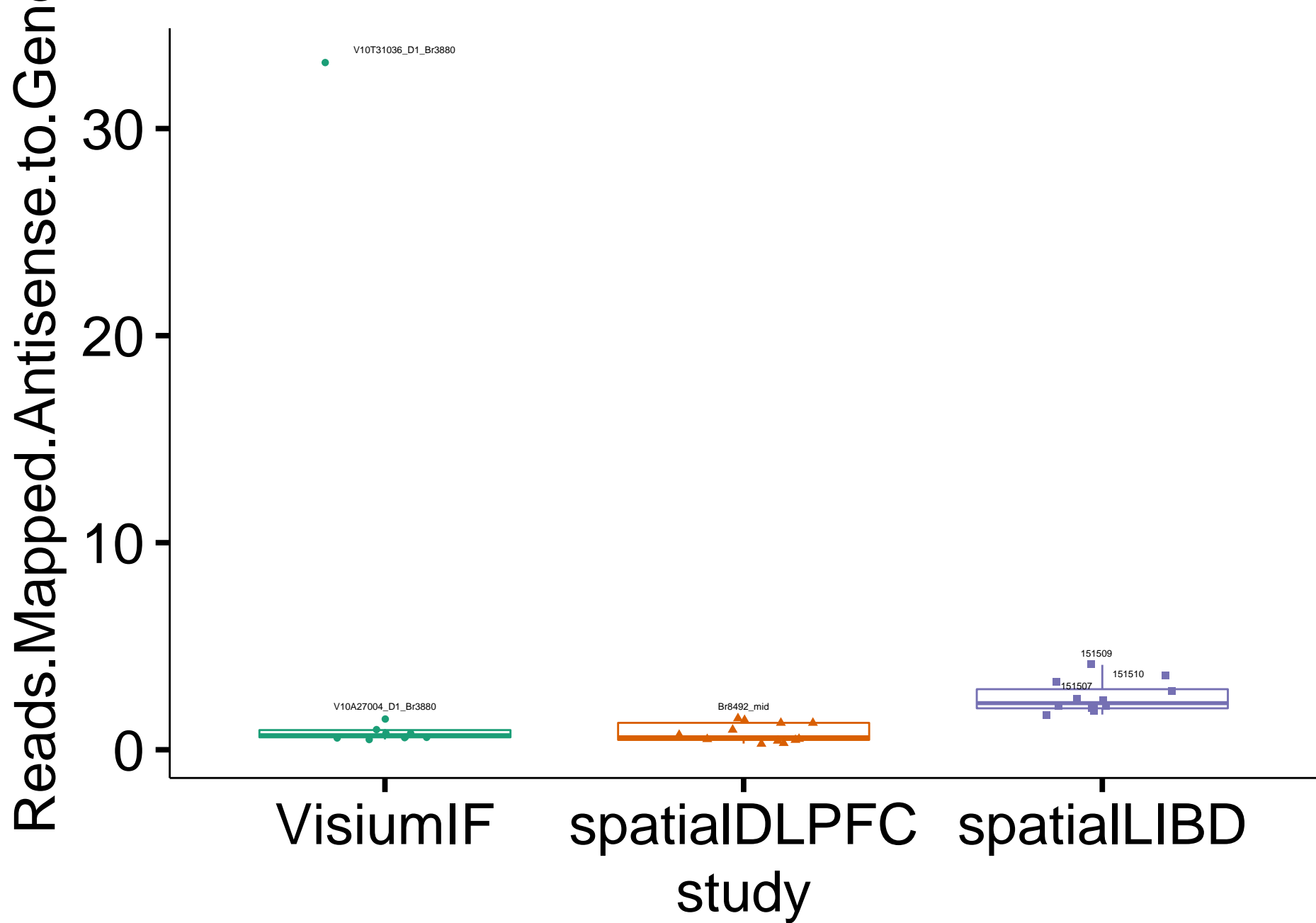






Reads.Mapped.Confidently.to.Transcri





Total.Genes.Detected

22000

21000

20000

VisiumIF

spatialDLPFC

spatialLIBD

study

