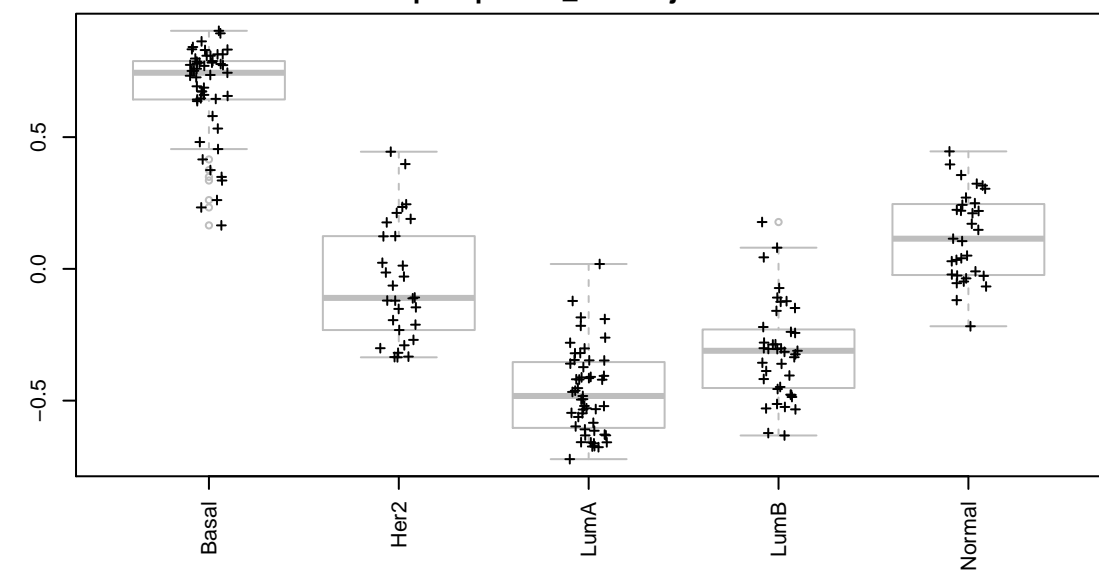
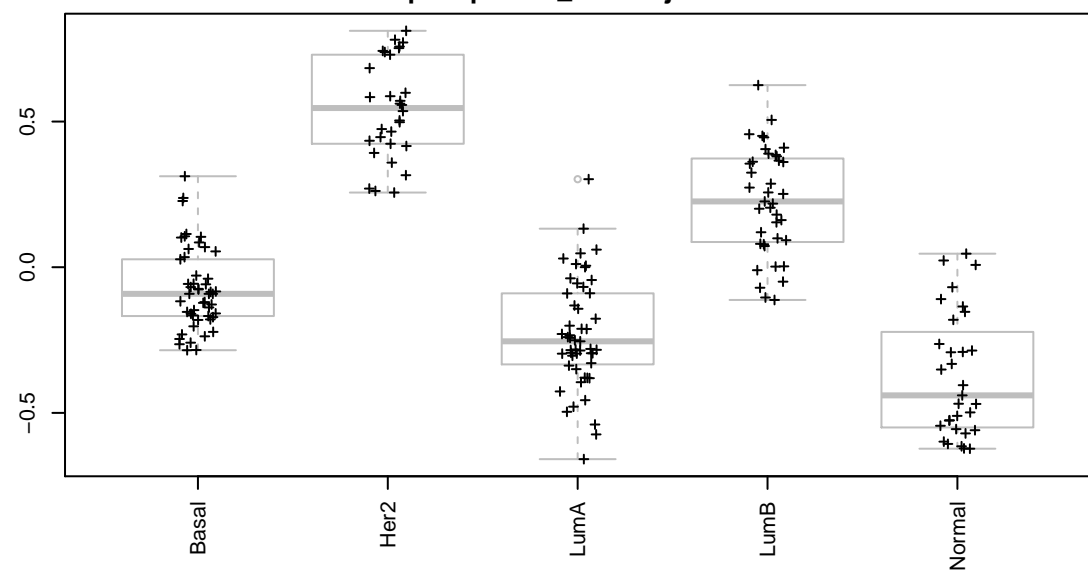


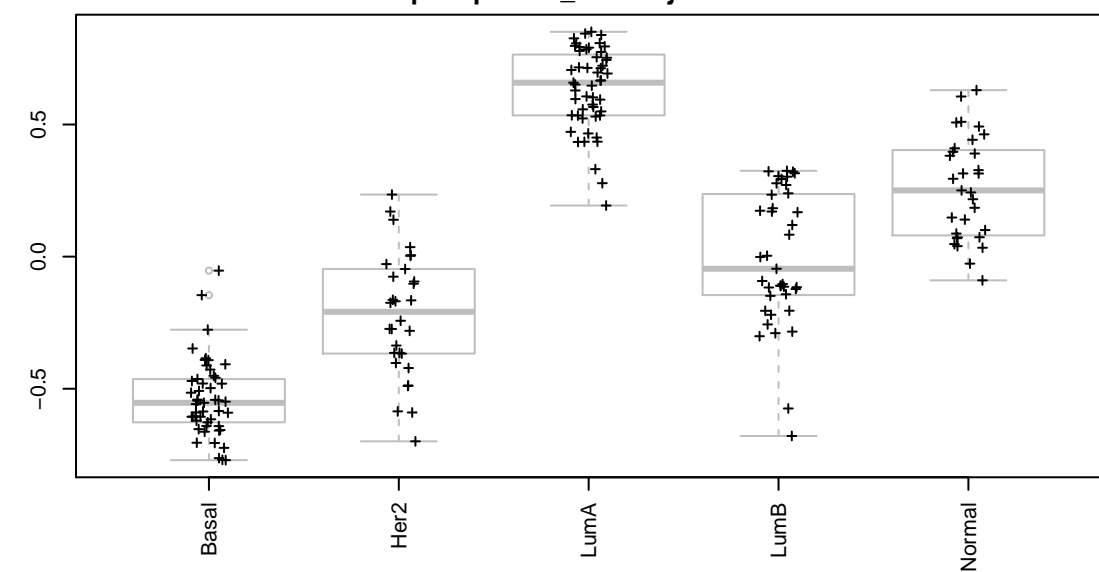
sampleInputFile_200subjects: Basal



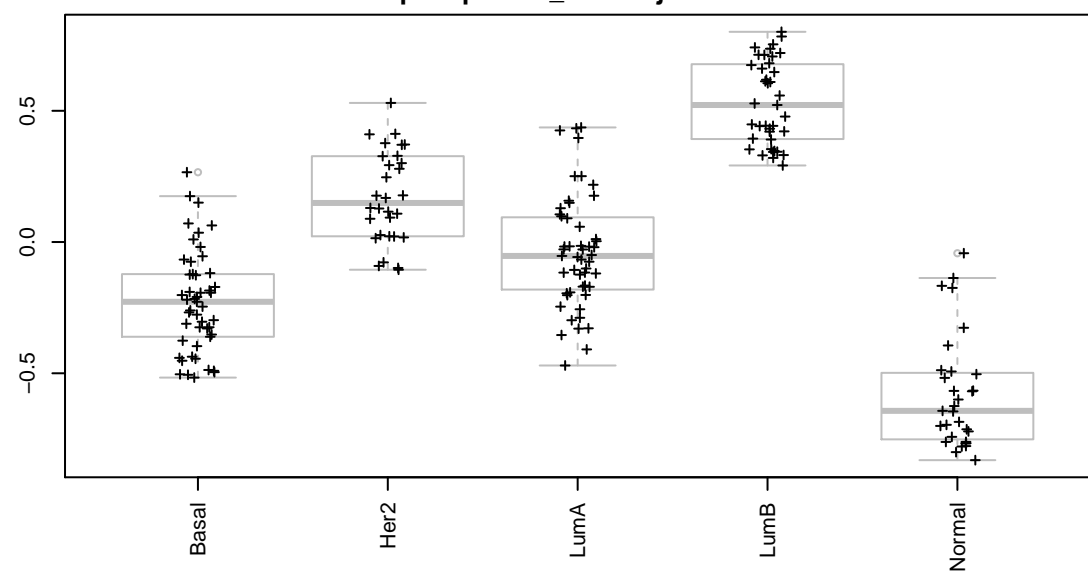
sampleInputFile_200subjects: Her2



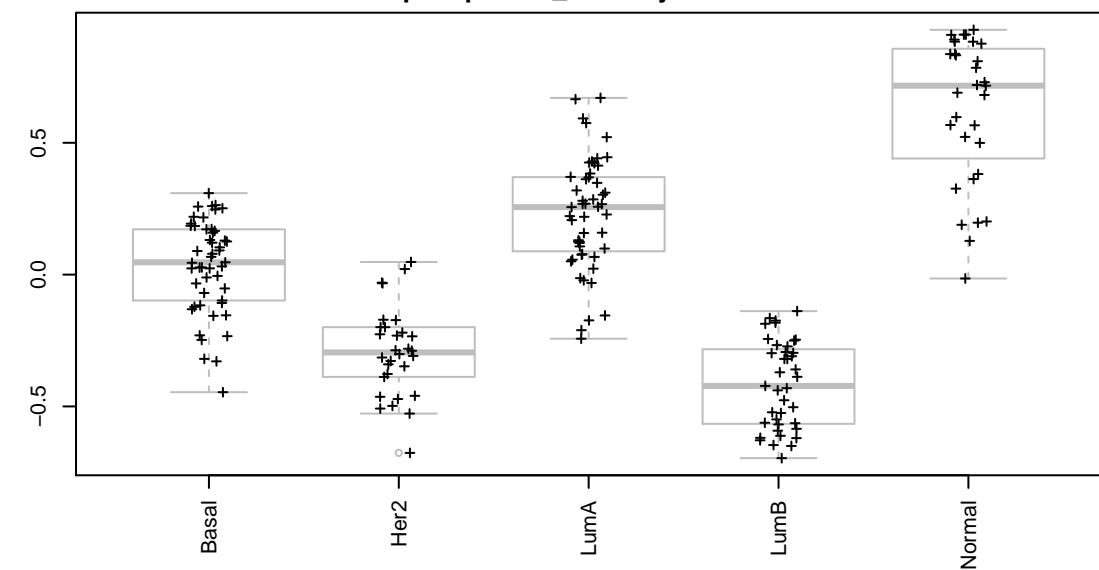
sampleInputFile_200subjects: LumA



sampleInputFile_200subjects: LumB



sampleInputFile_200subjects: Normal



sampleInputFile_200subjects: Proliferation Index

