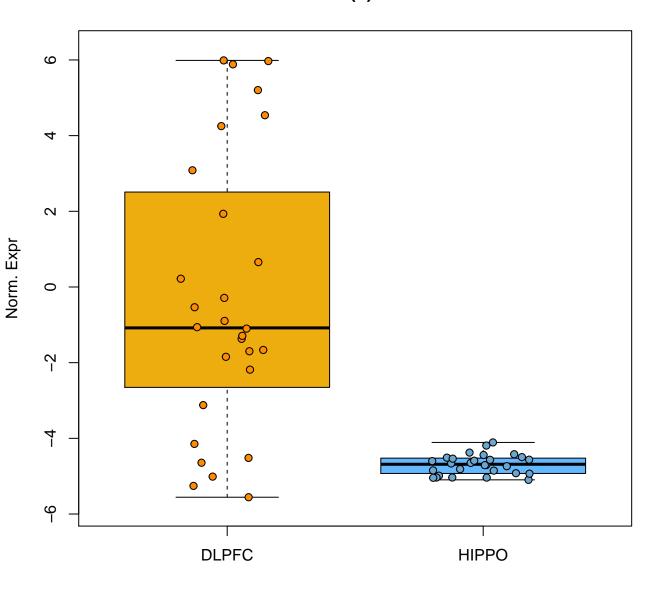
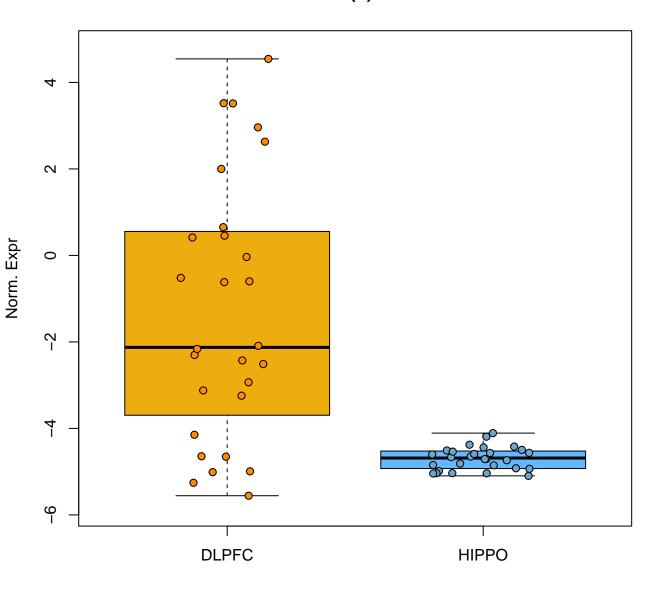
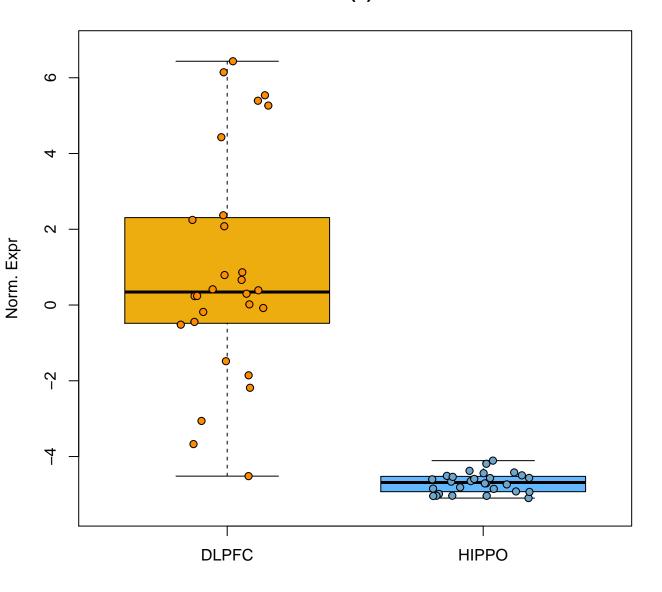
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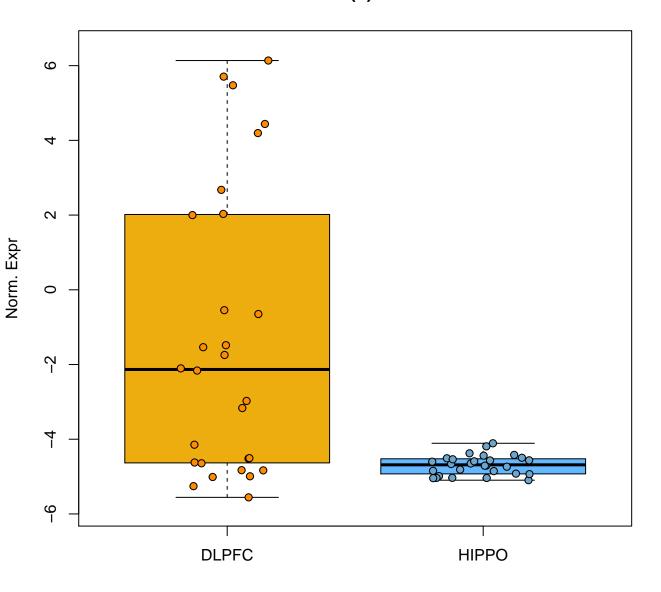
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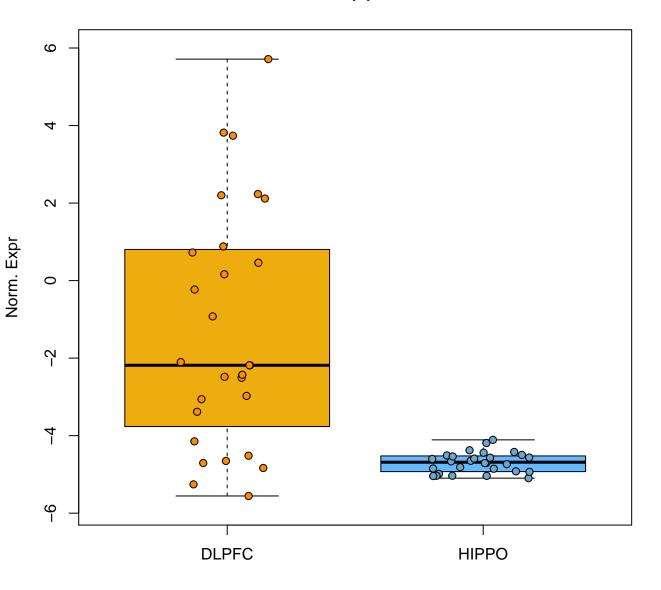
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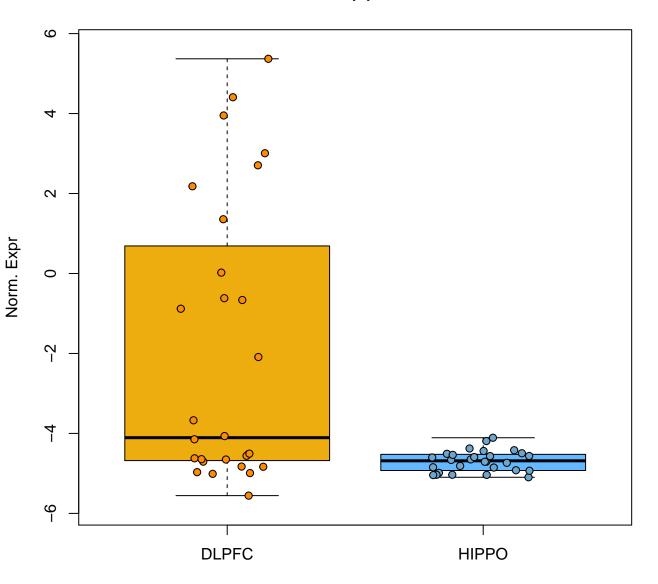
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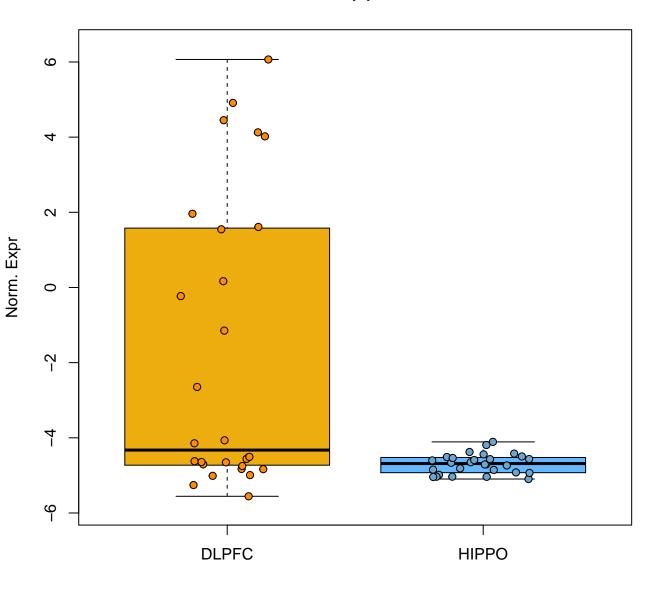
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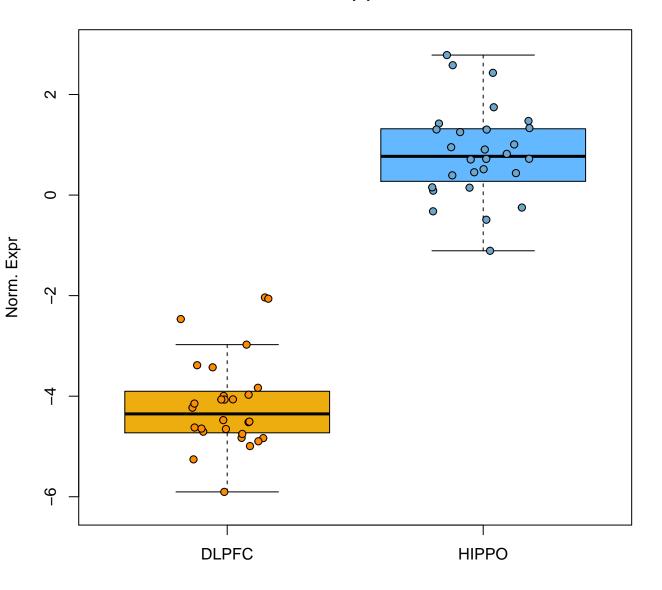
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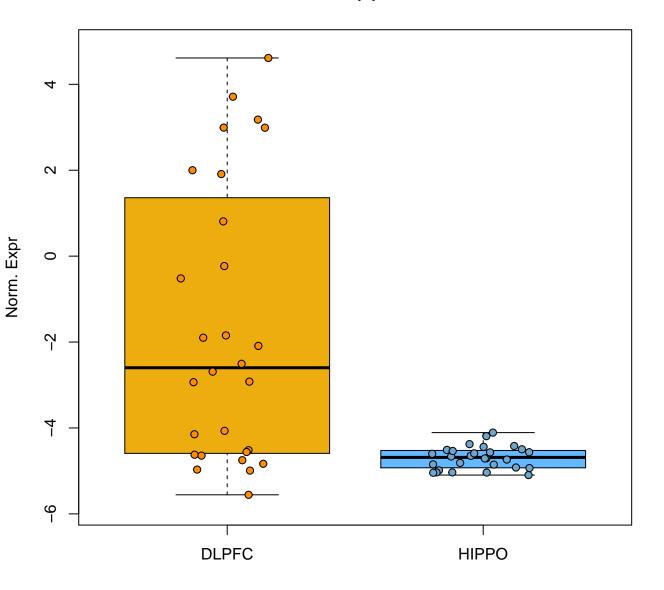
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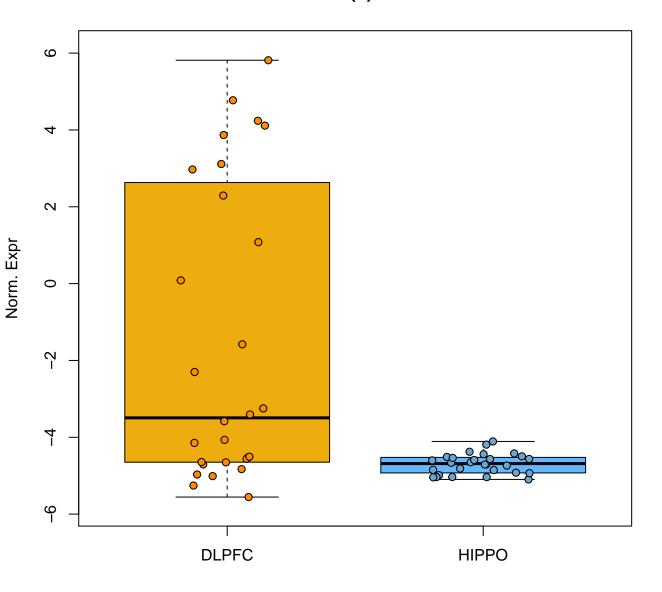
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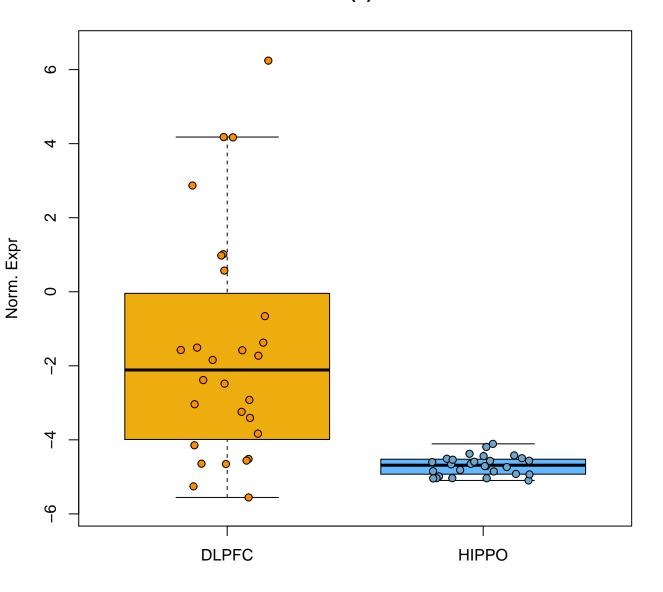
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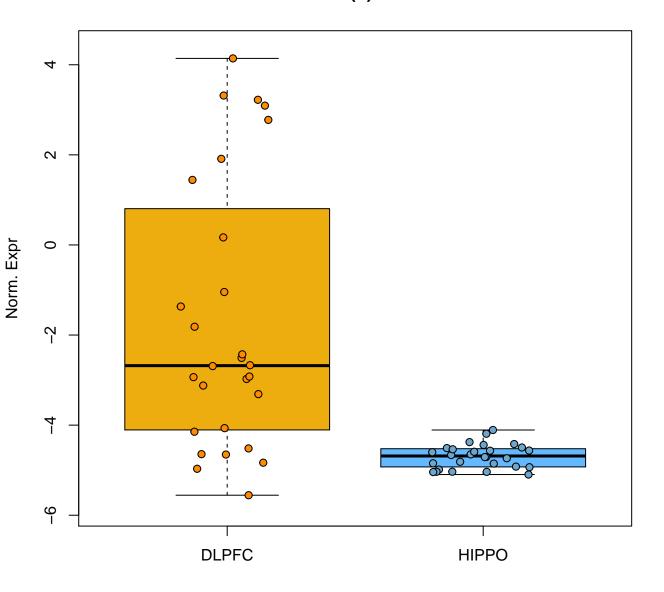
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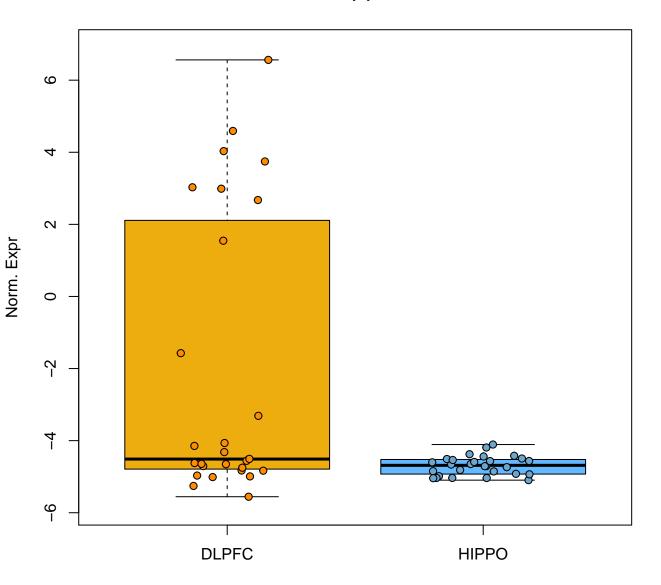
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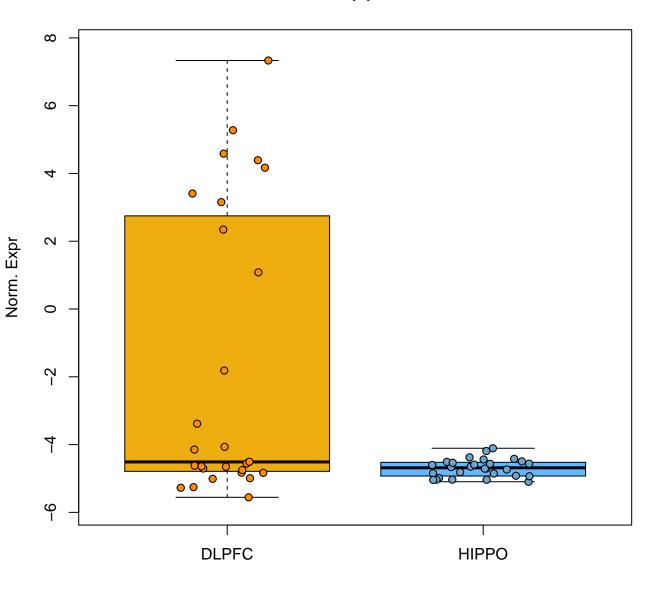
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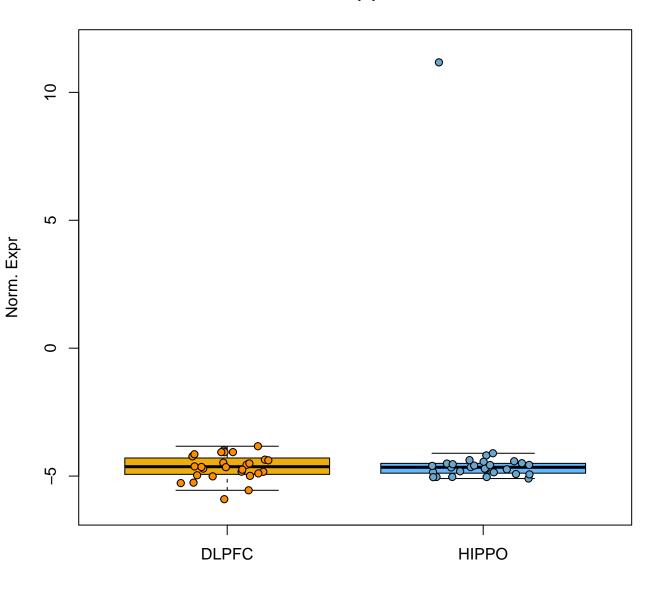
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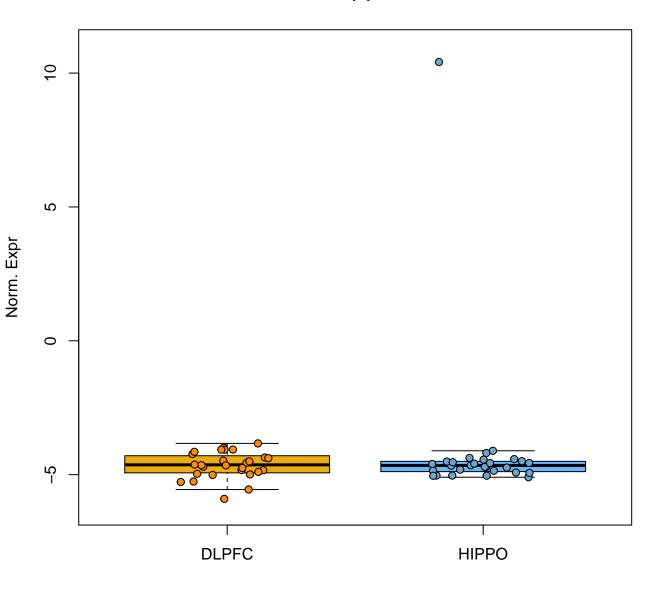
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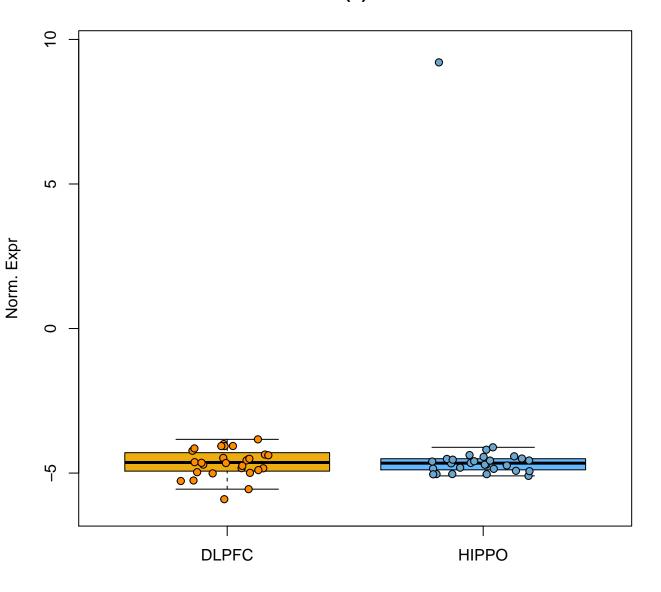
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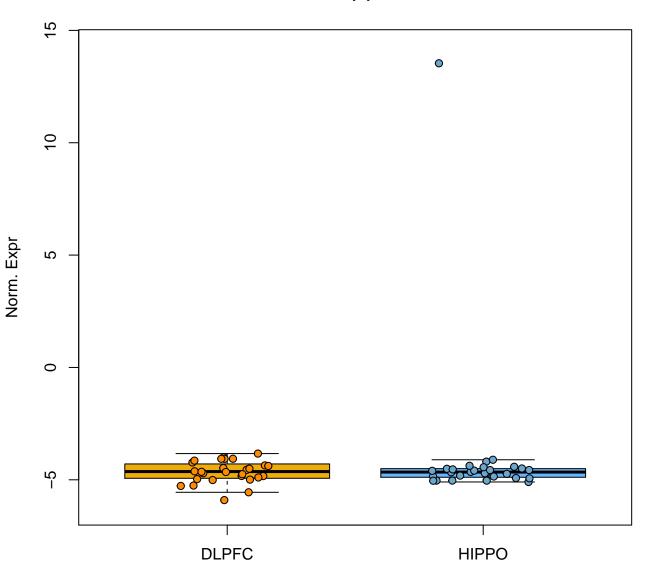
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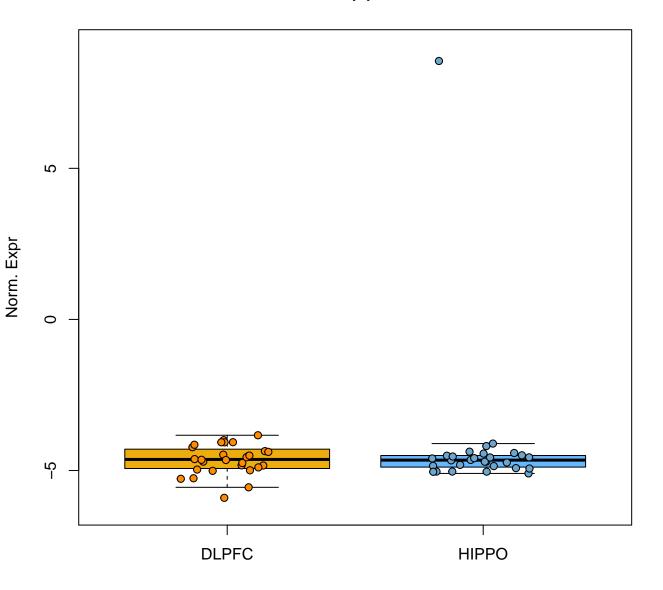
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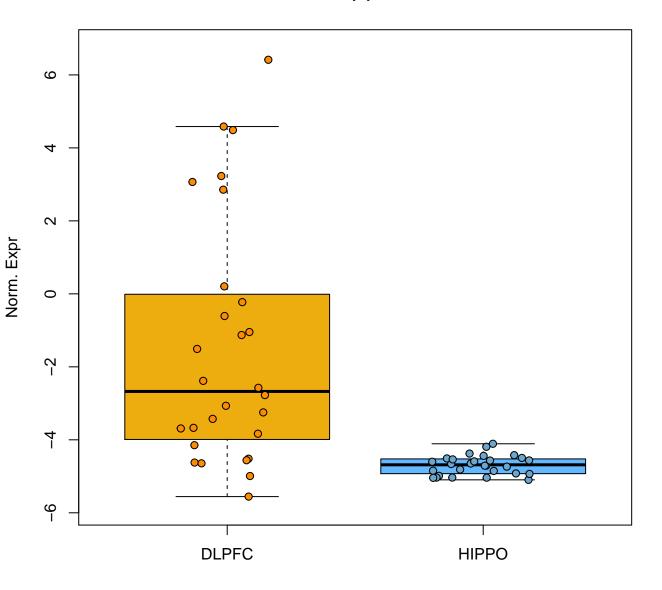
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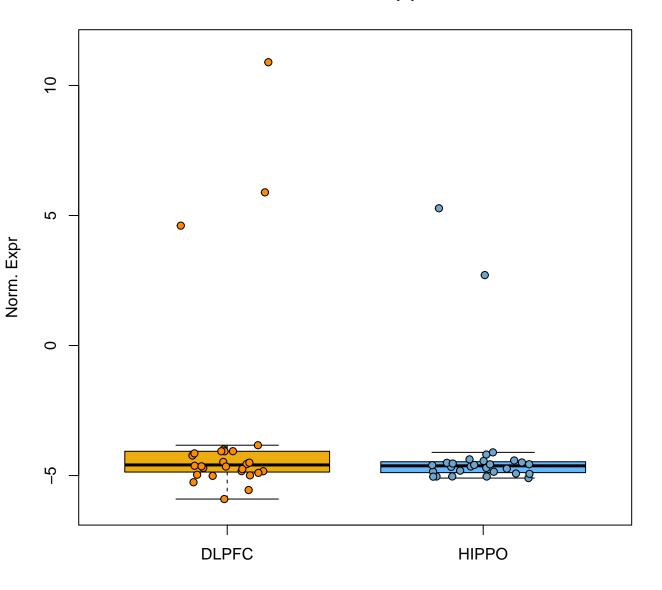
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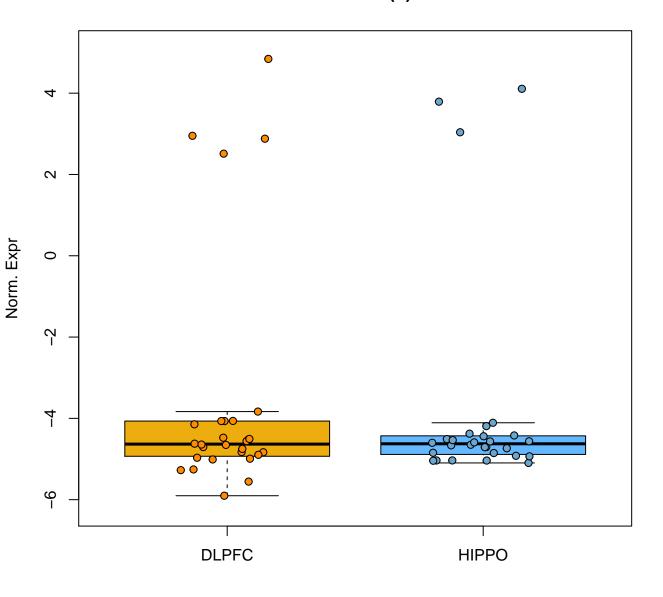
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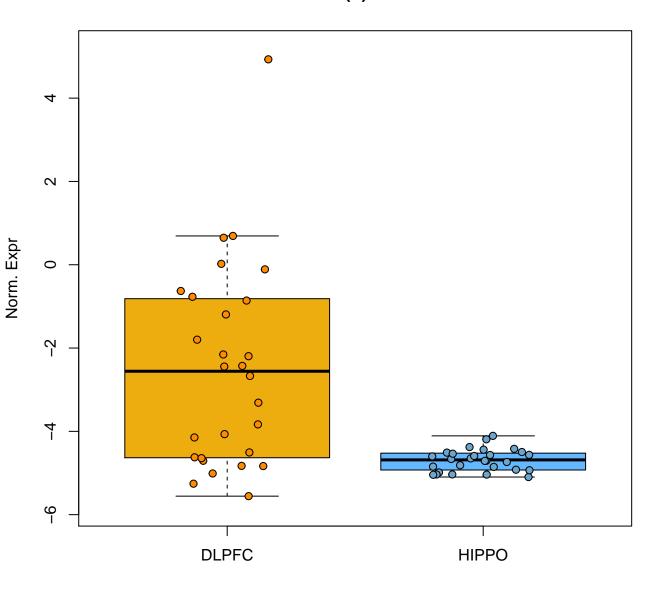
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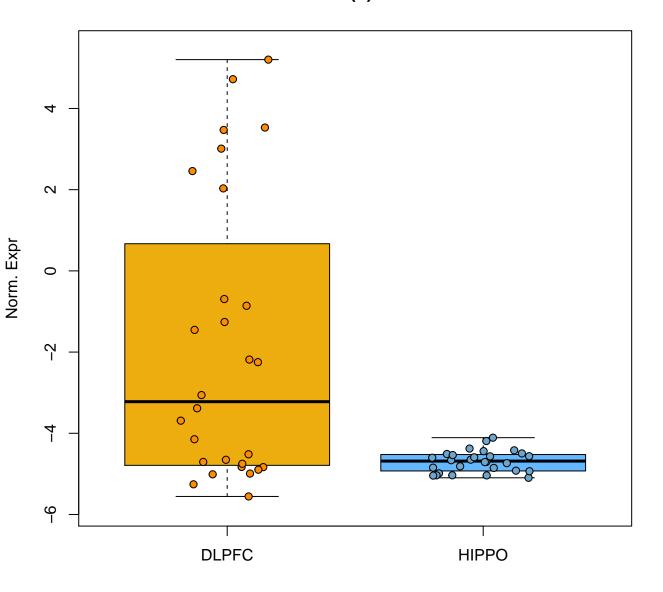
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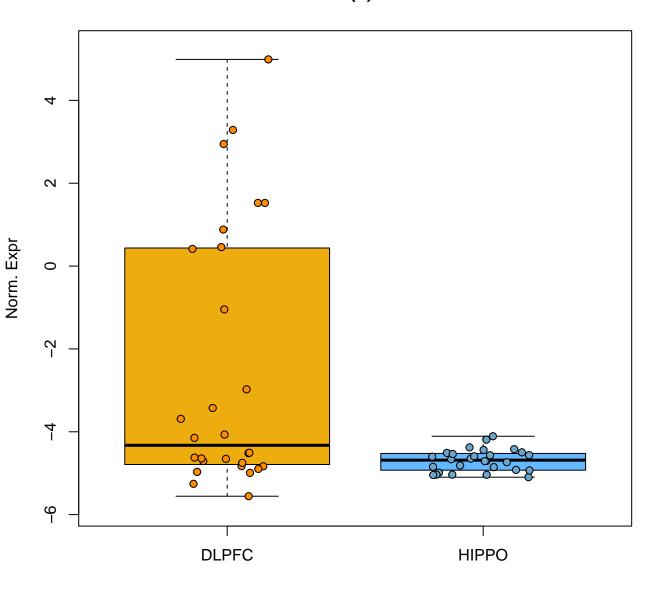
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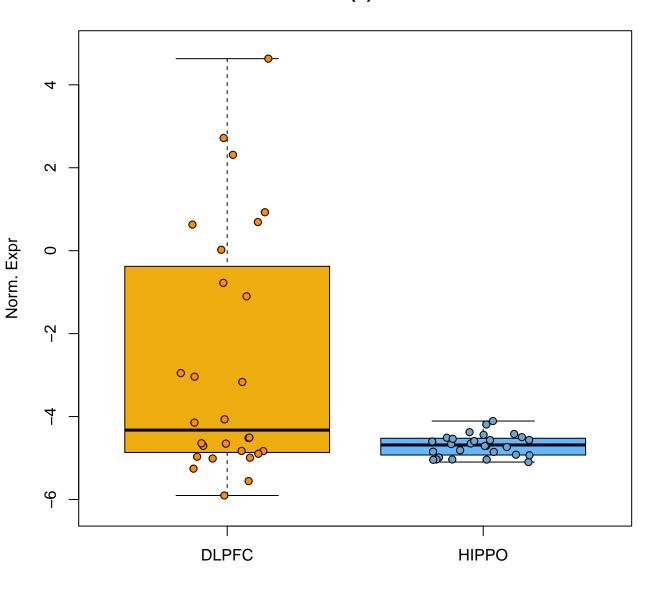
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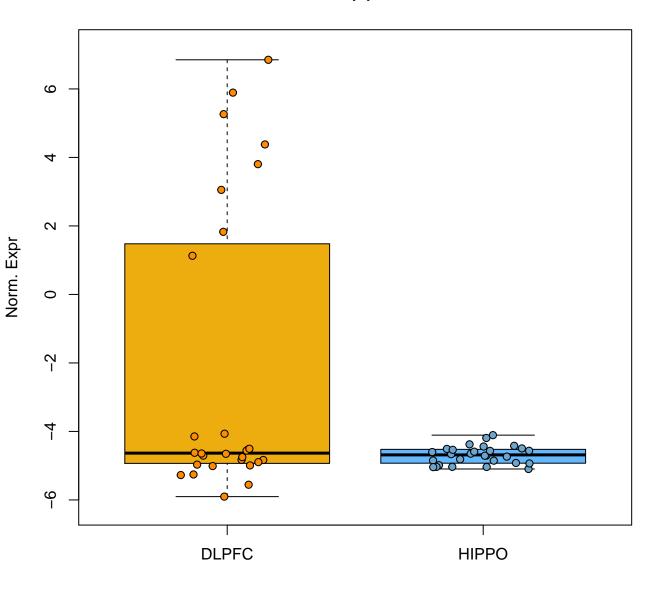
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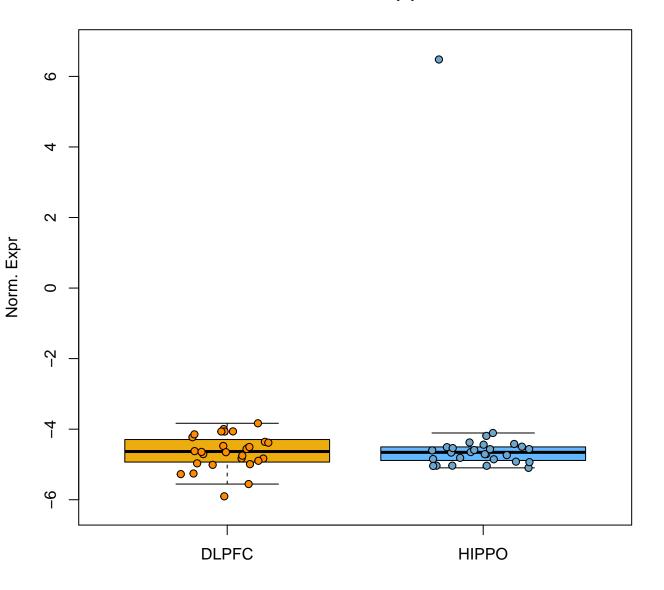
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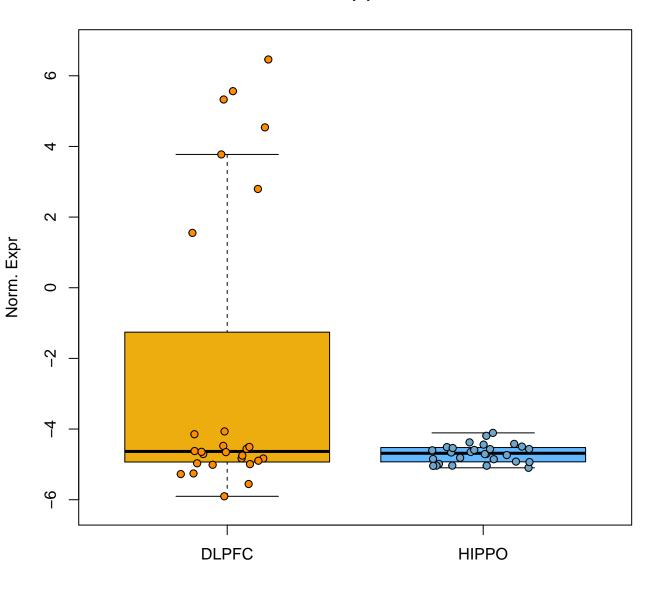
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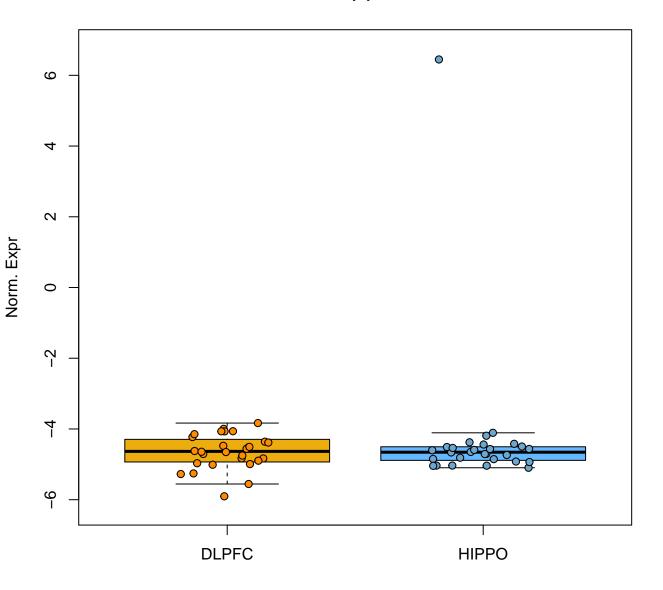
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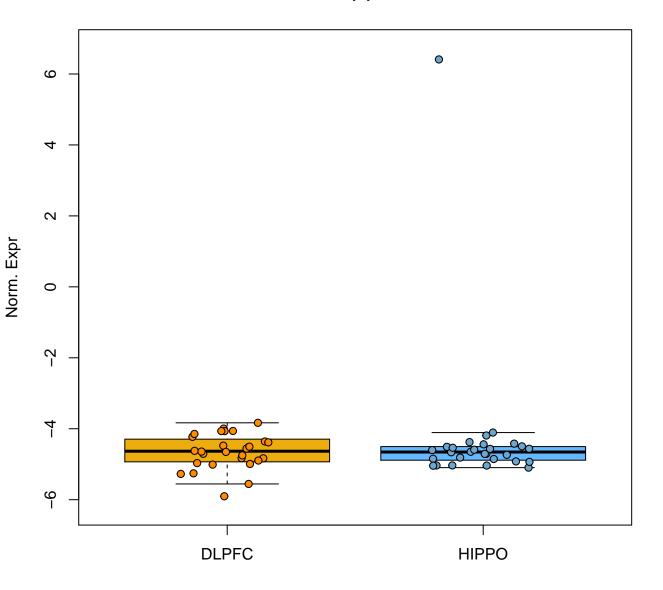
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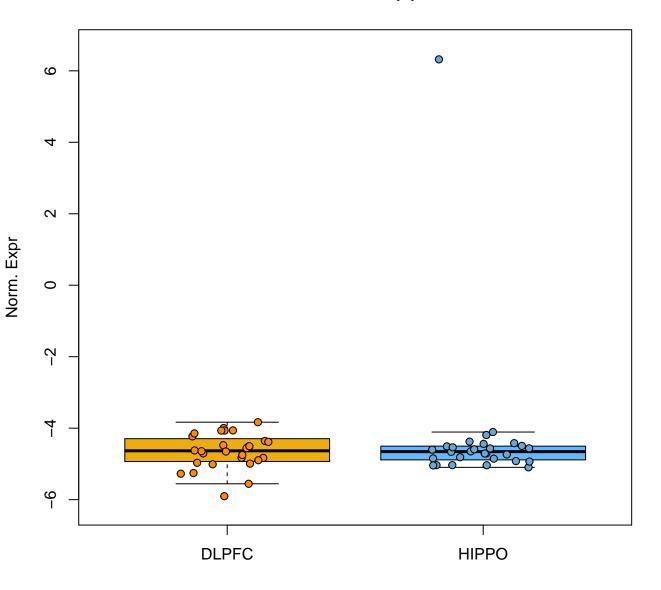
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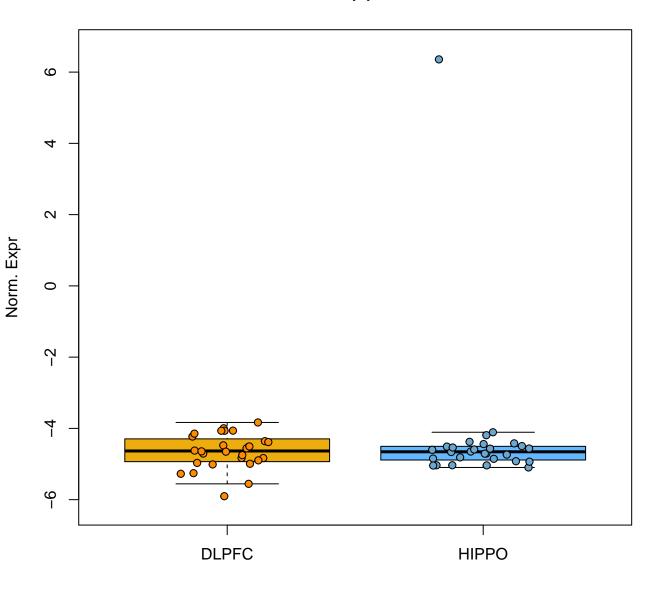
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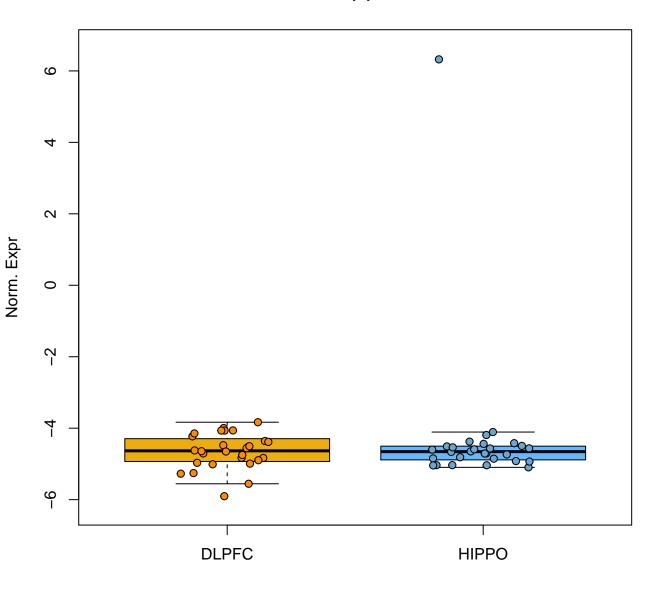
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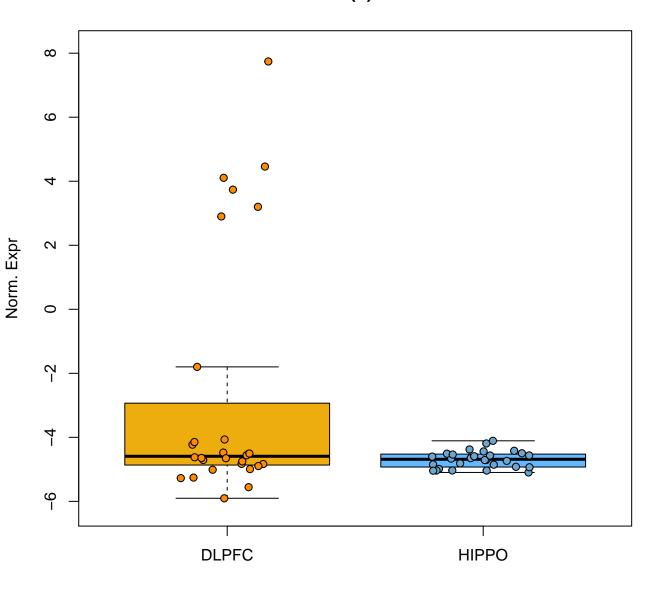
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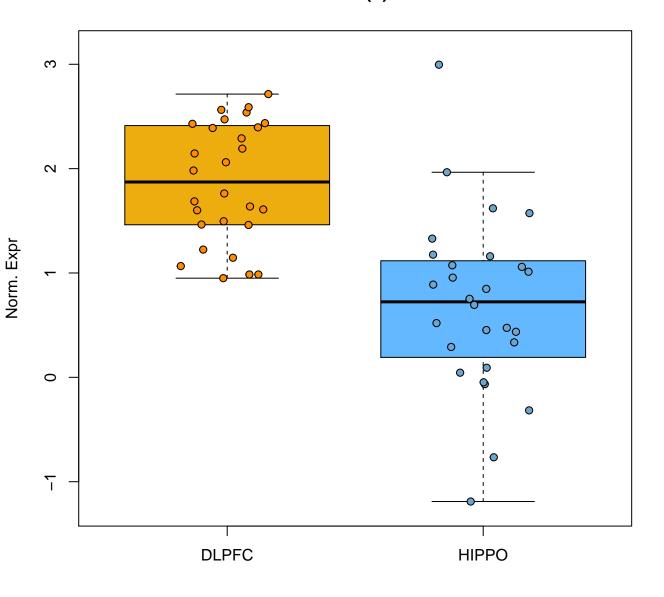
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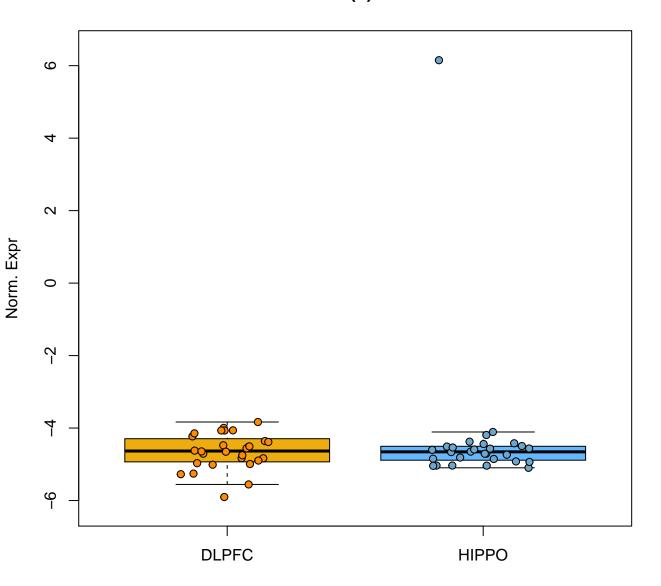
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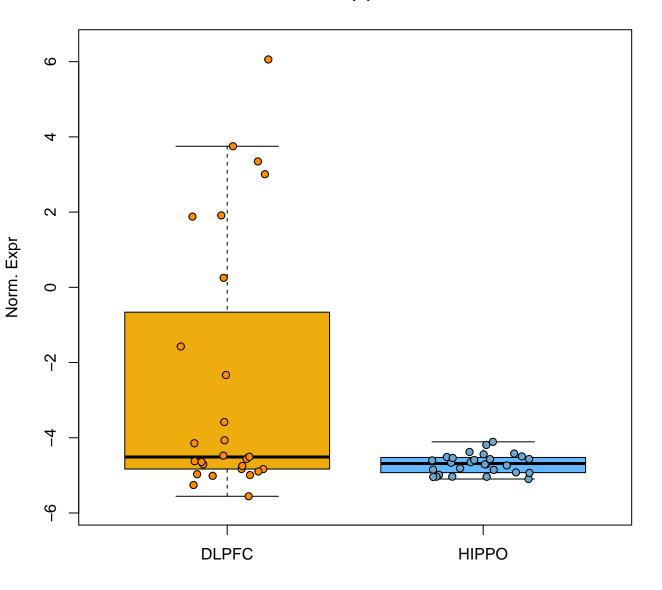
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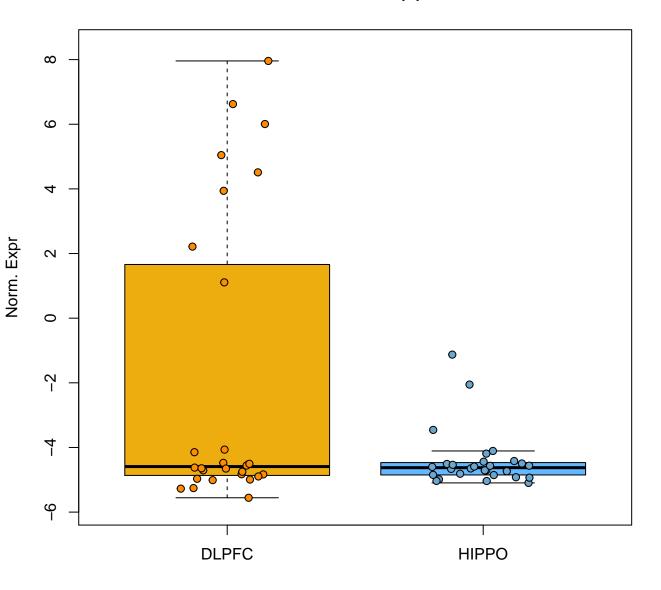
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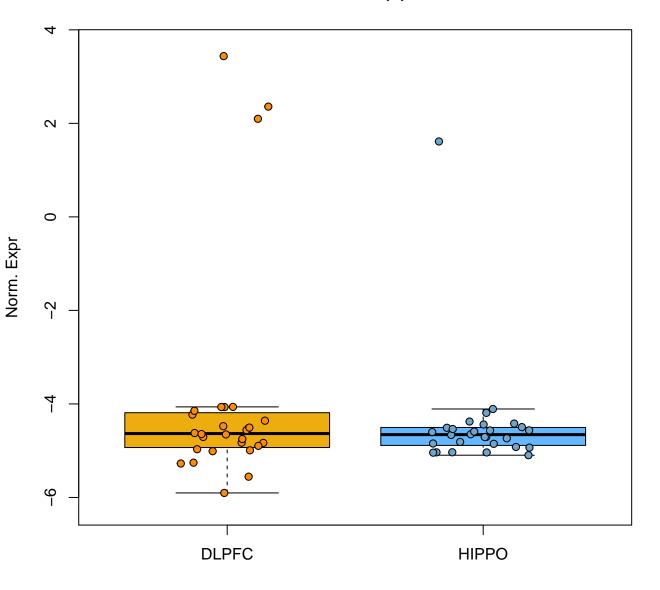
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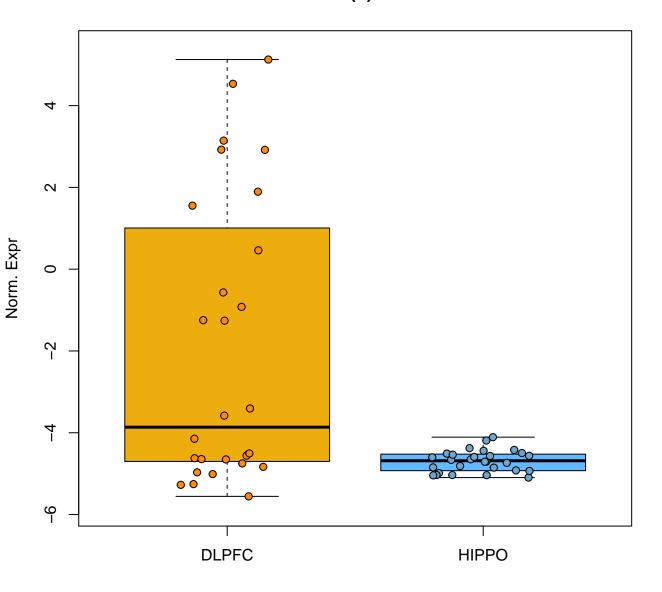
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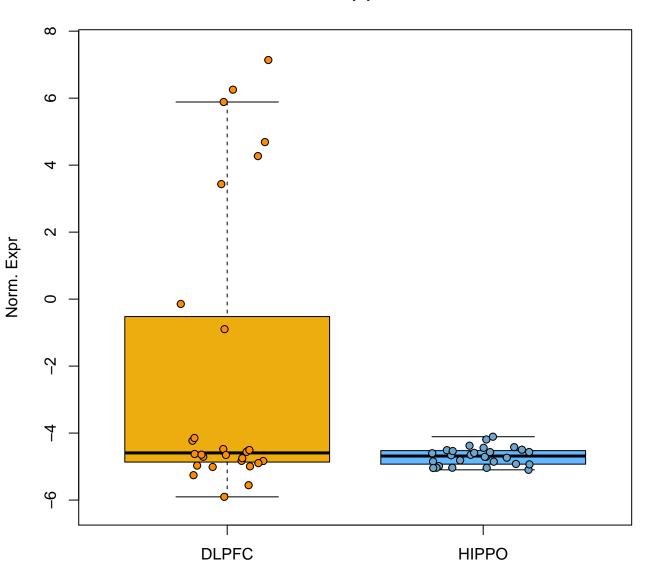
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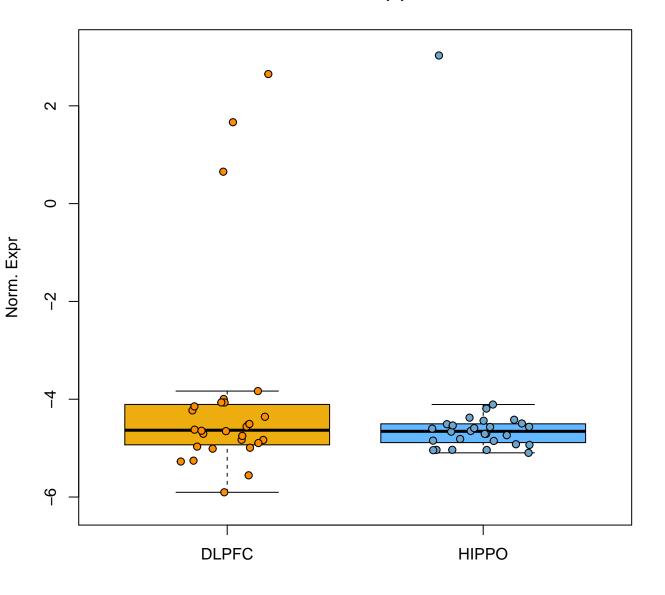
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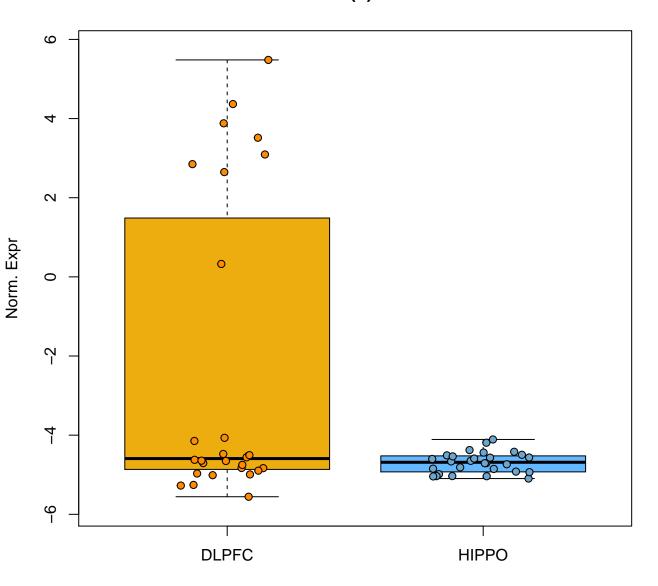
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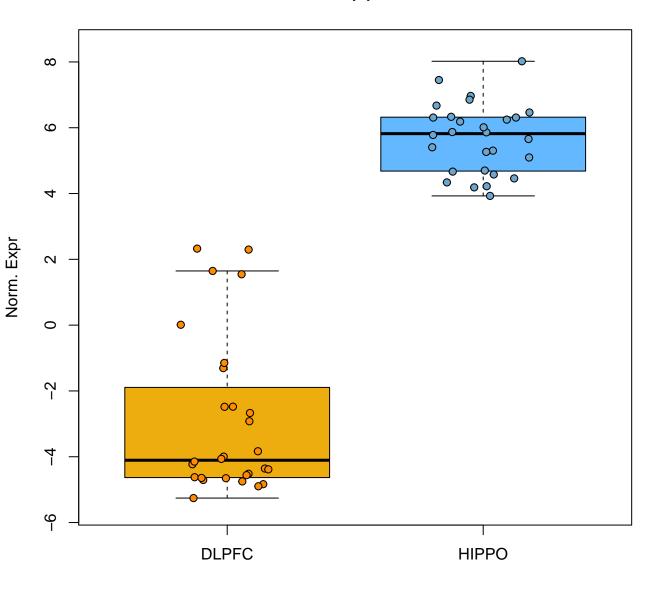
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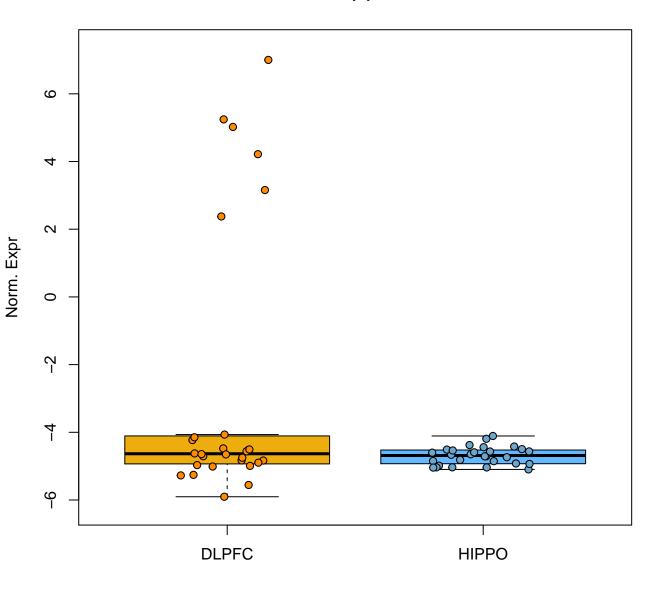
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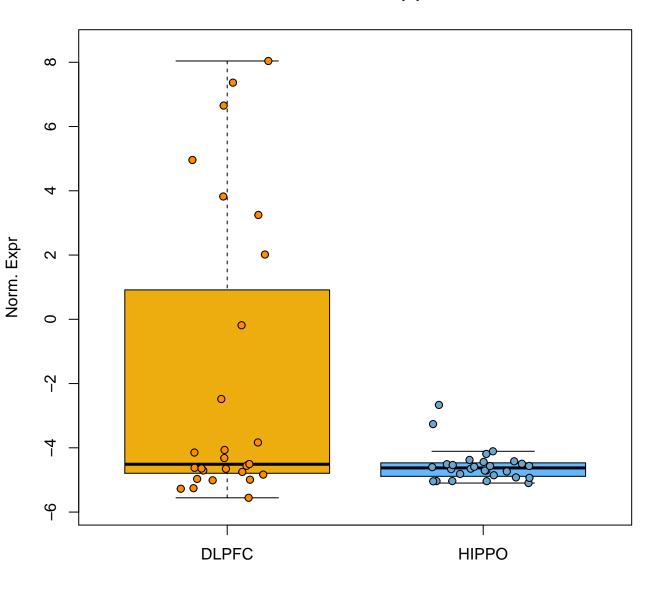
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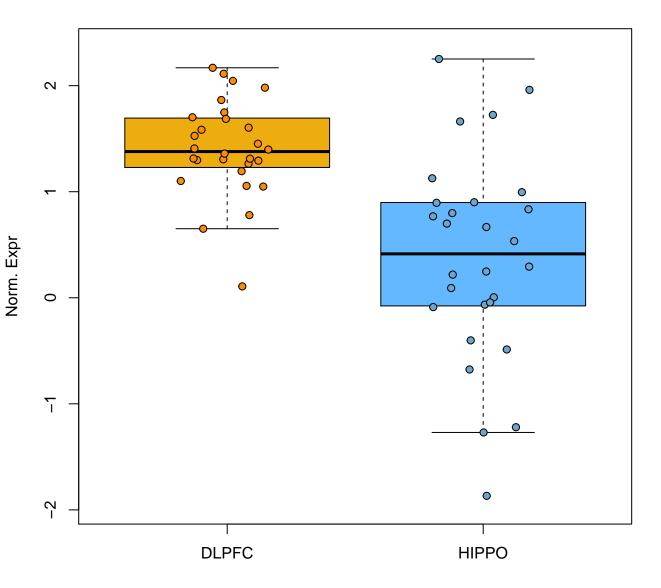
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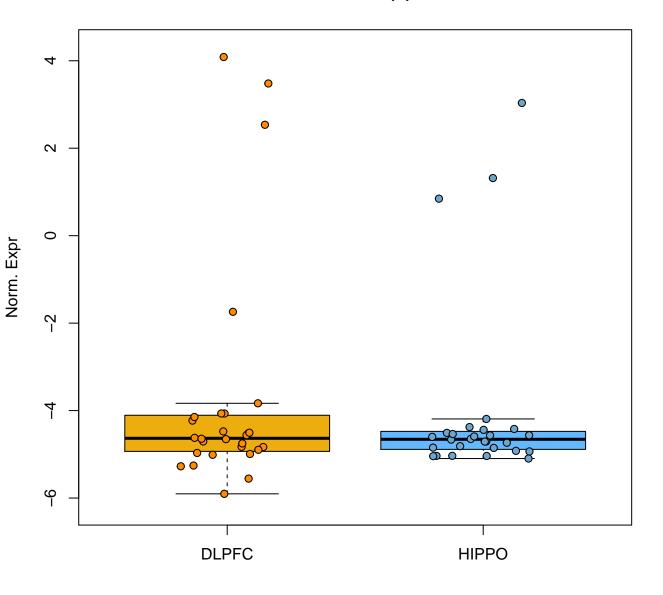
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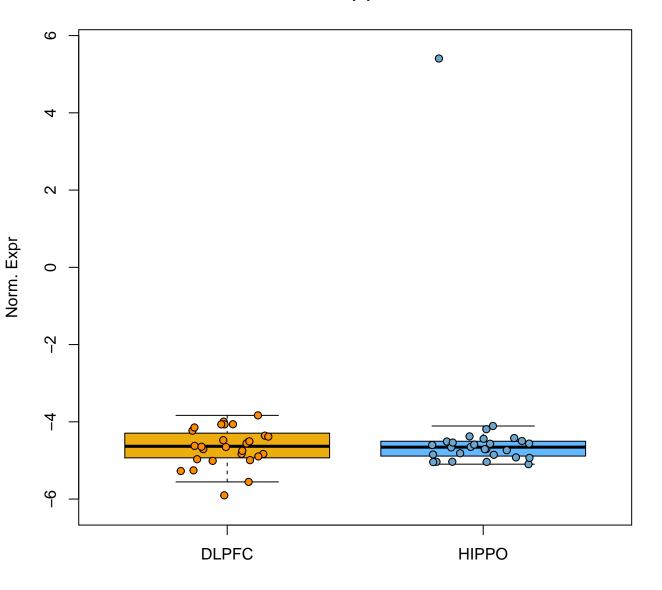
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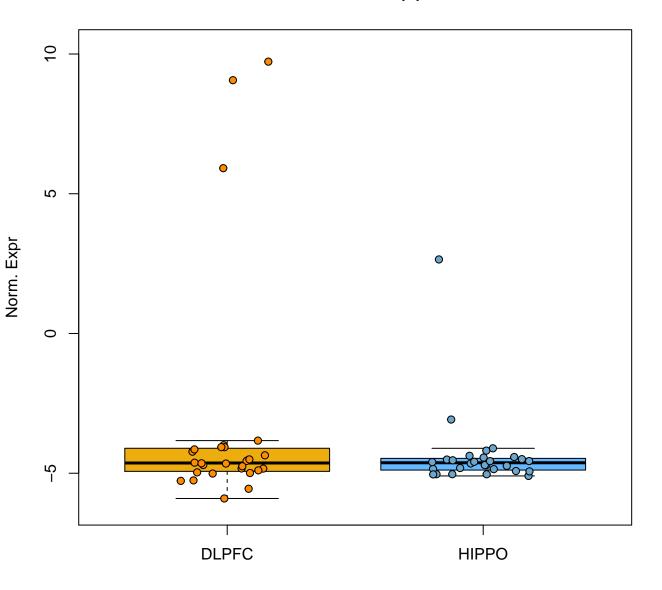
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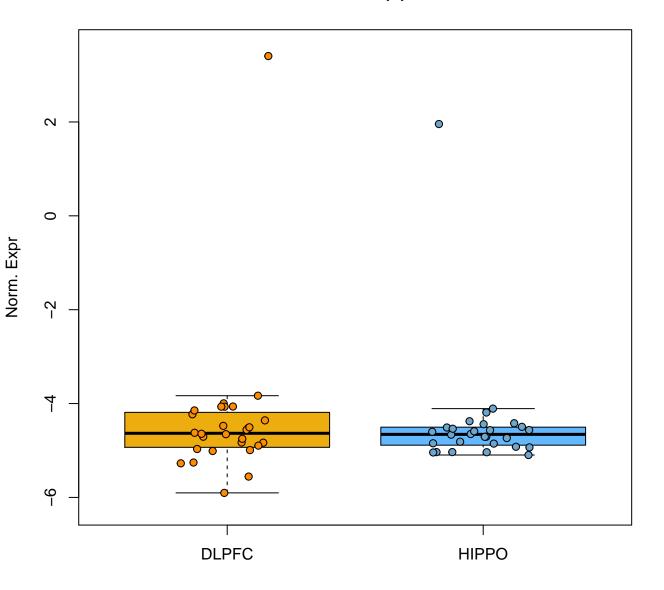
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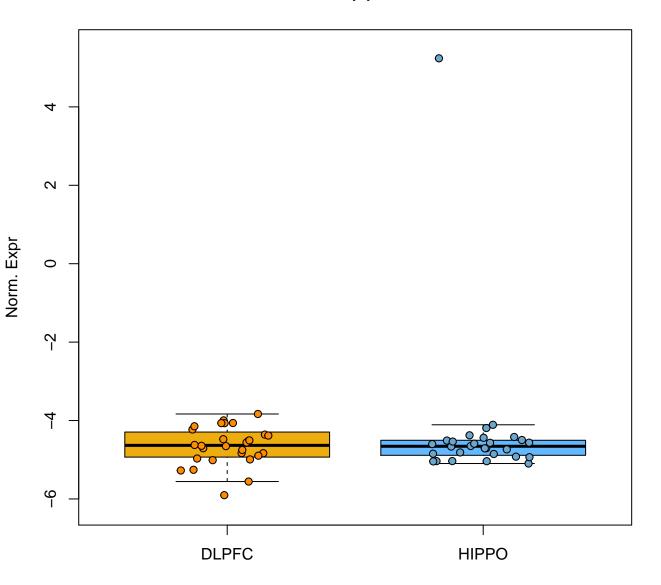
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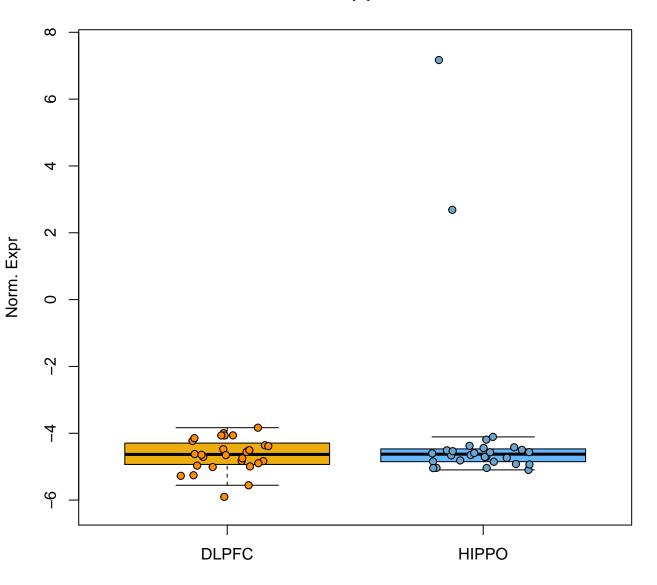
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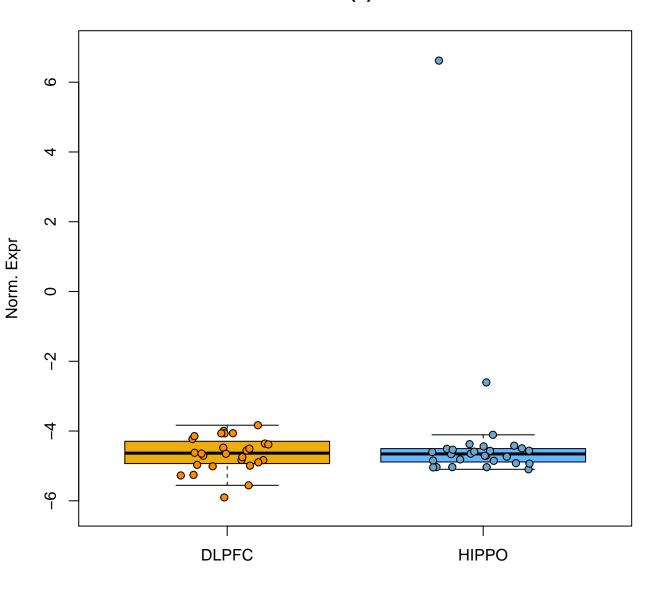
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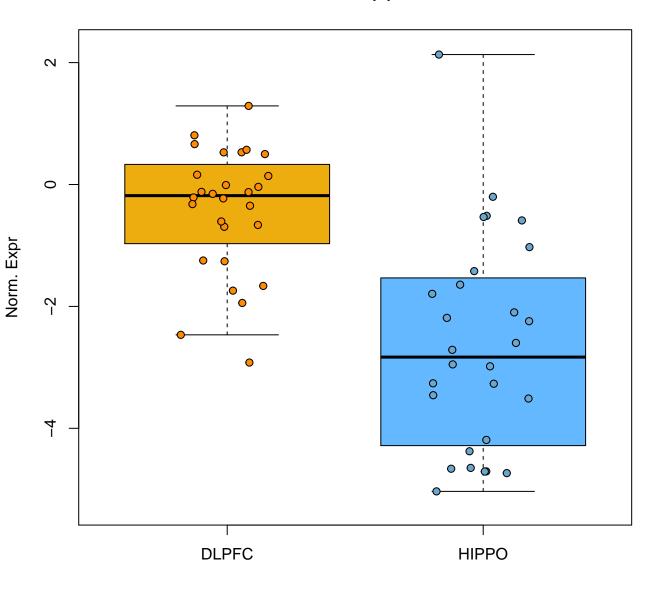
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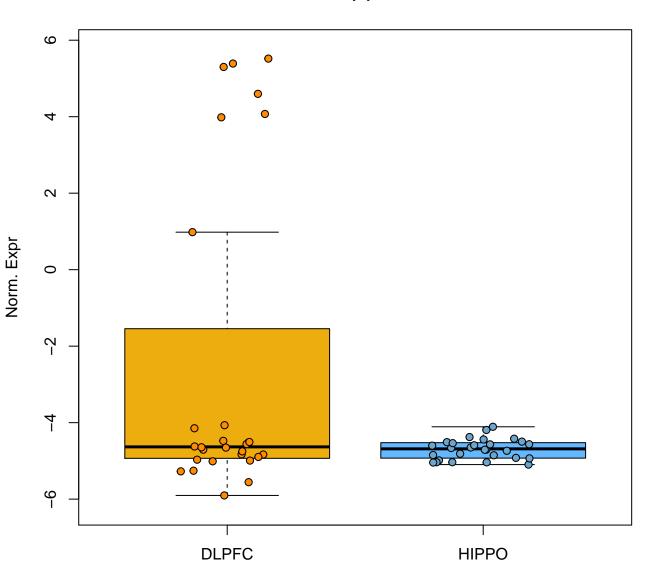
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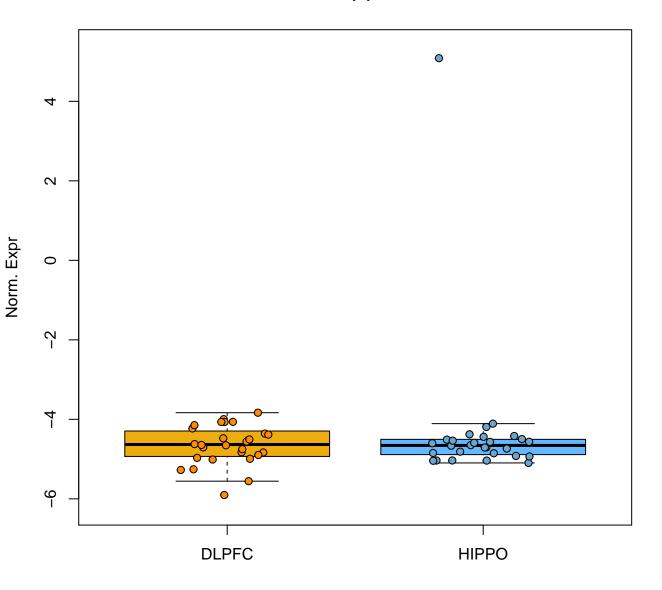
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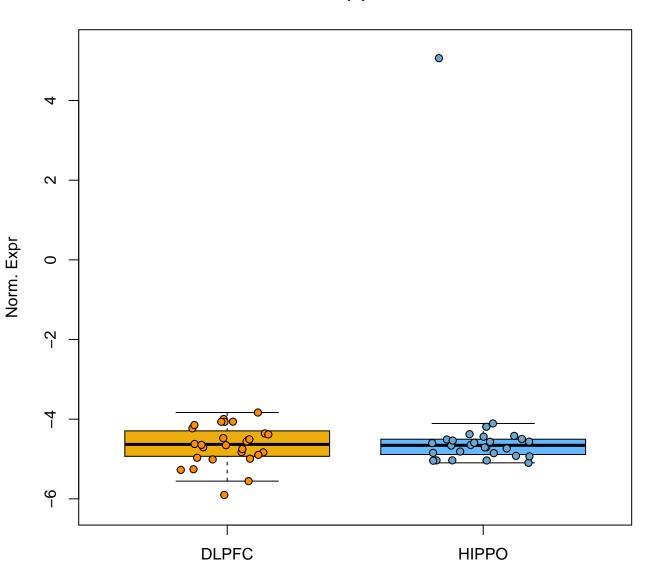
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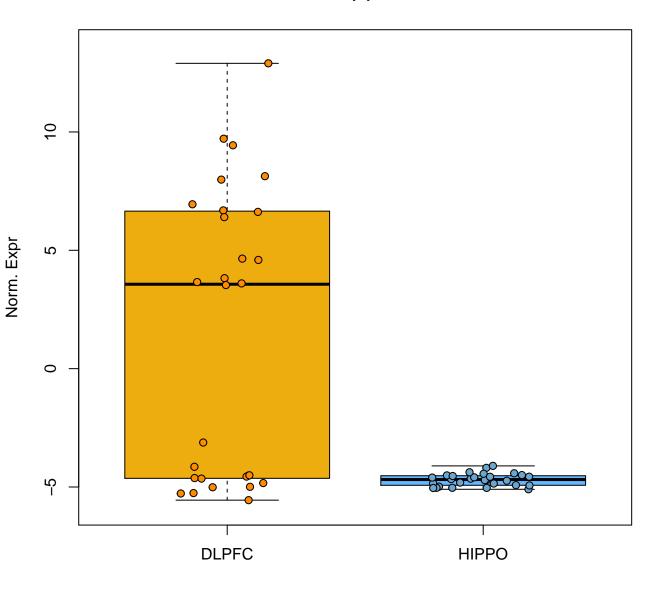
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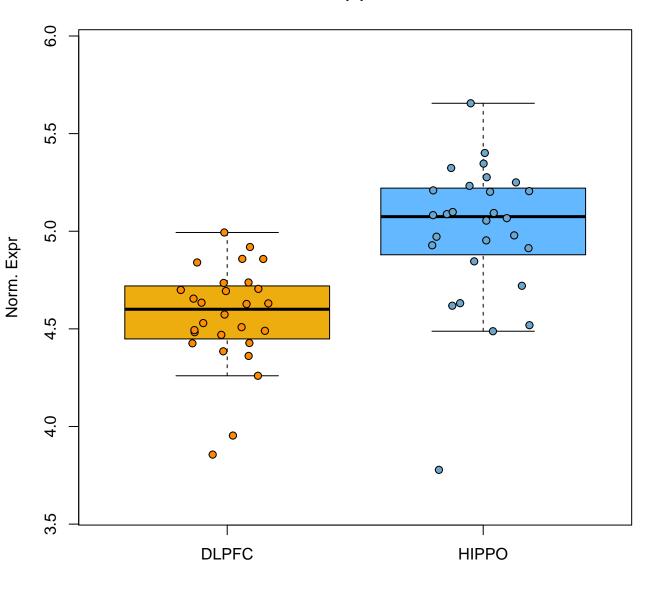
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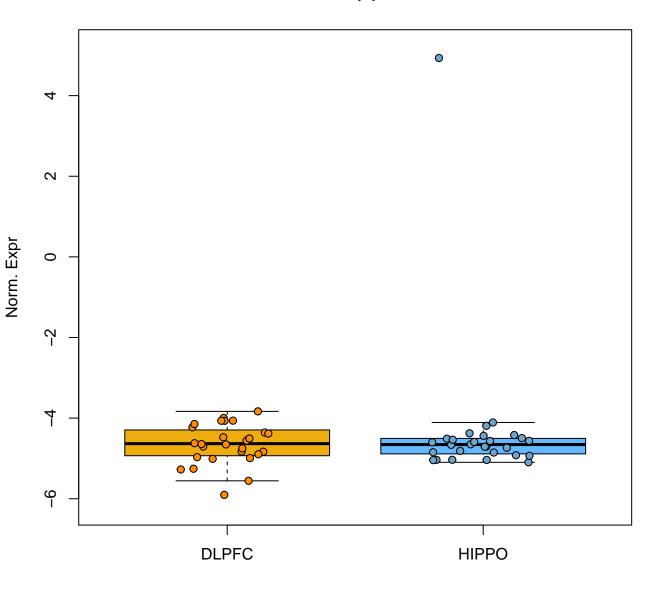
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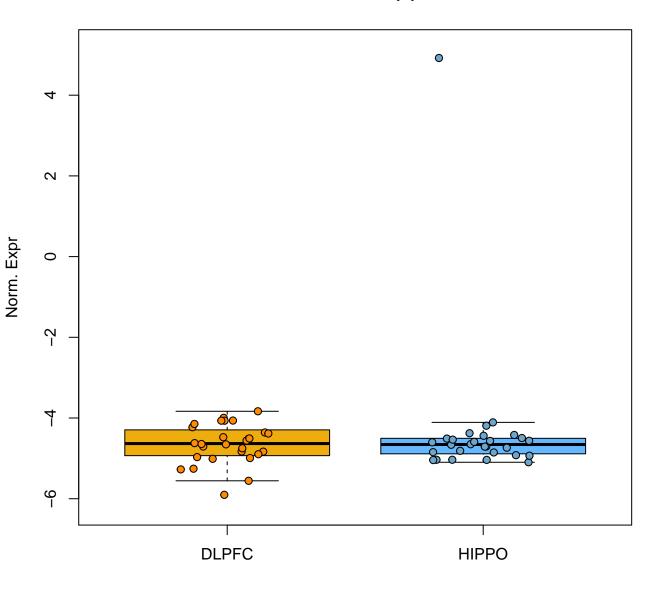
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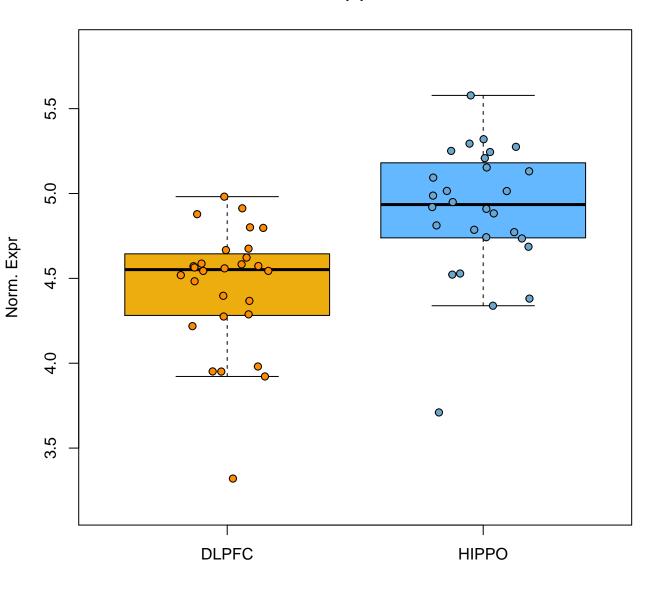
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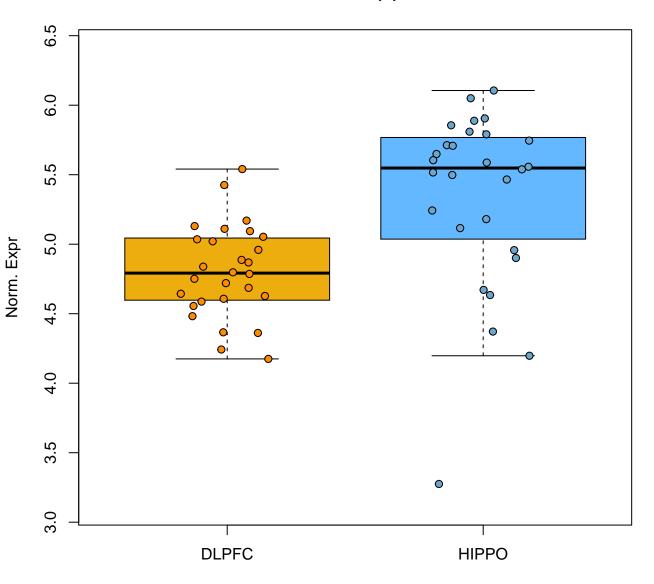
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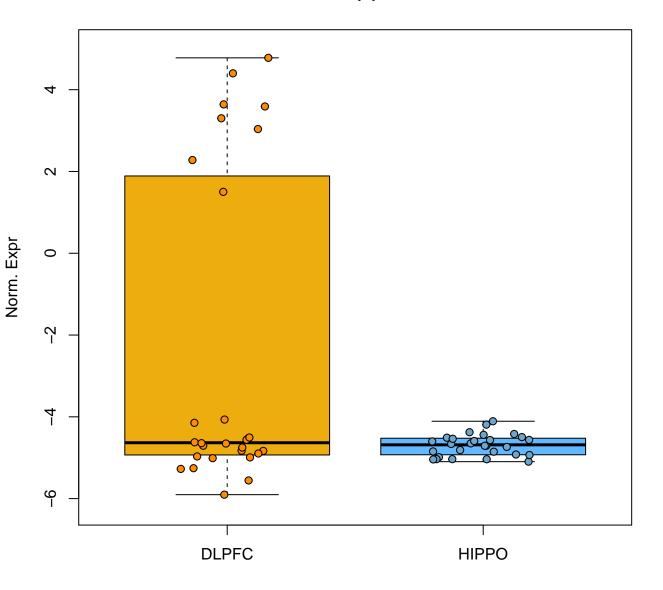
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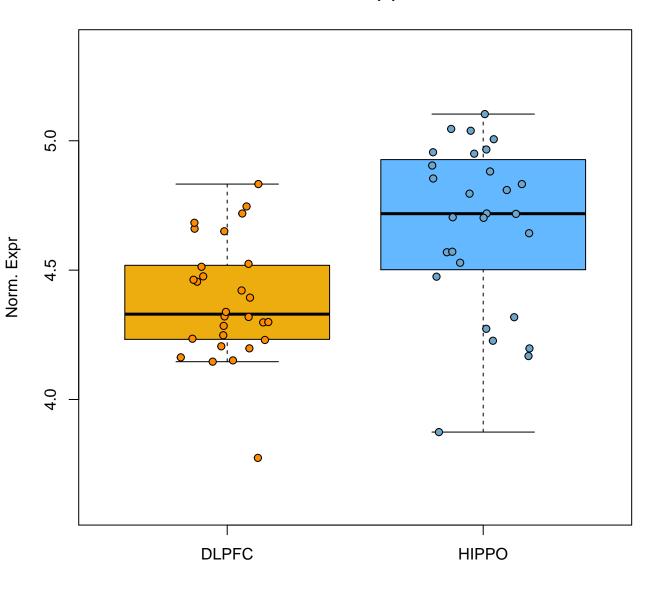
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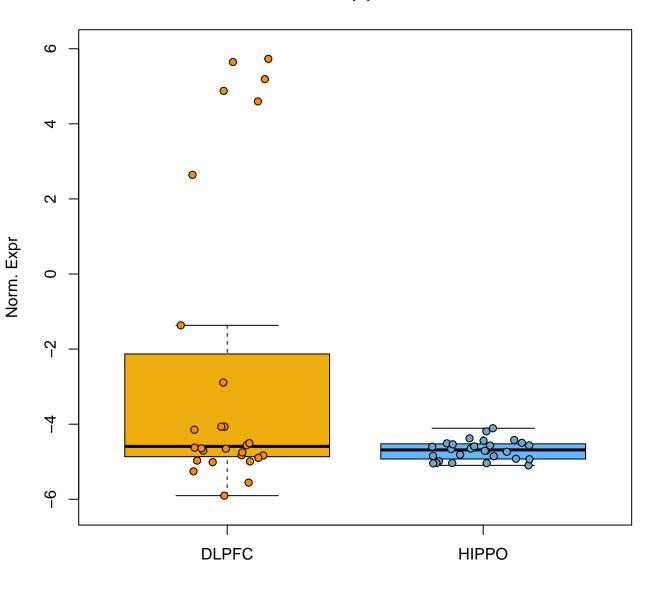
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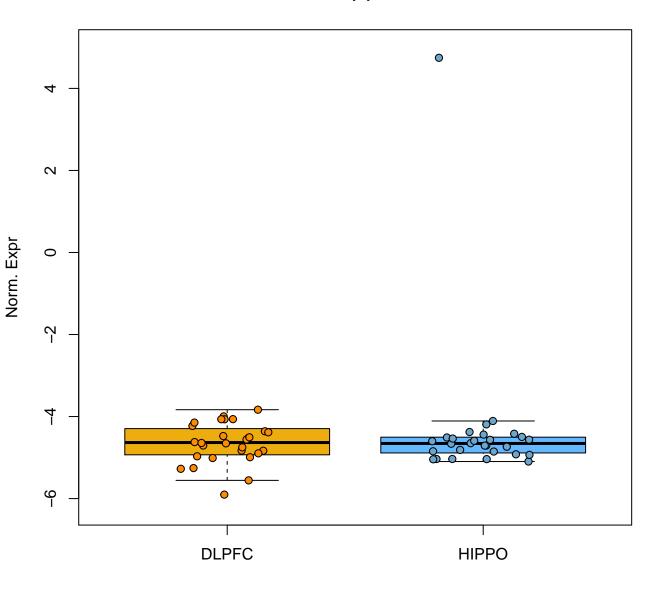
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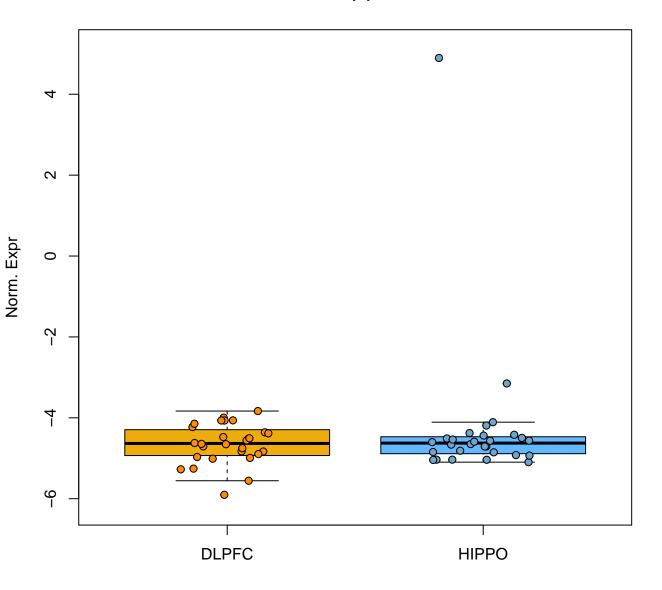
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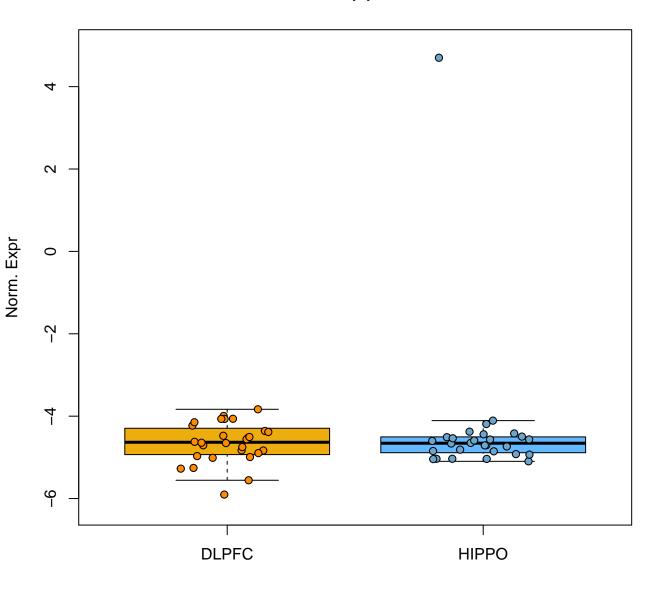
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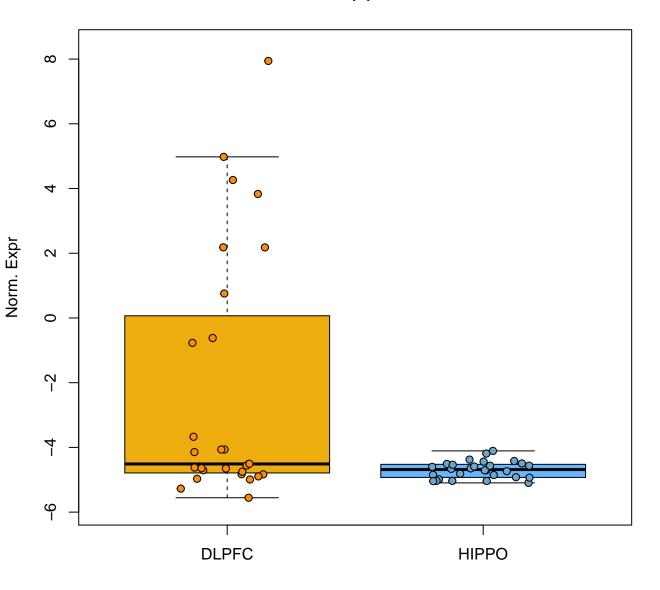
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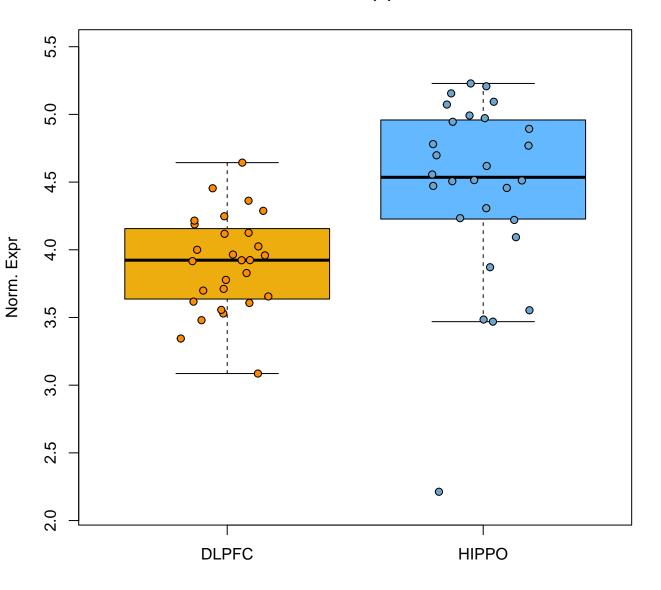
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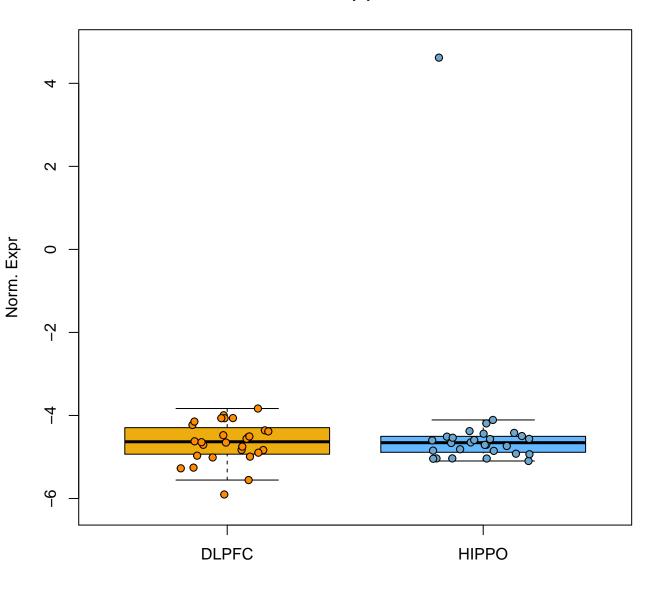
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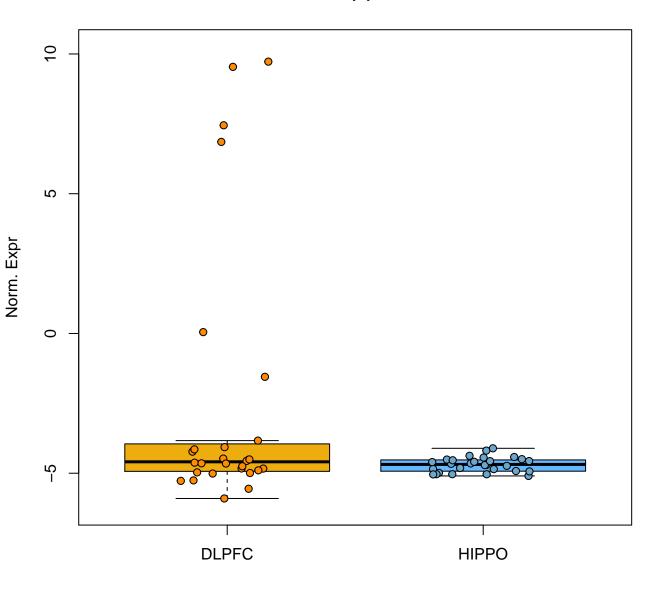
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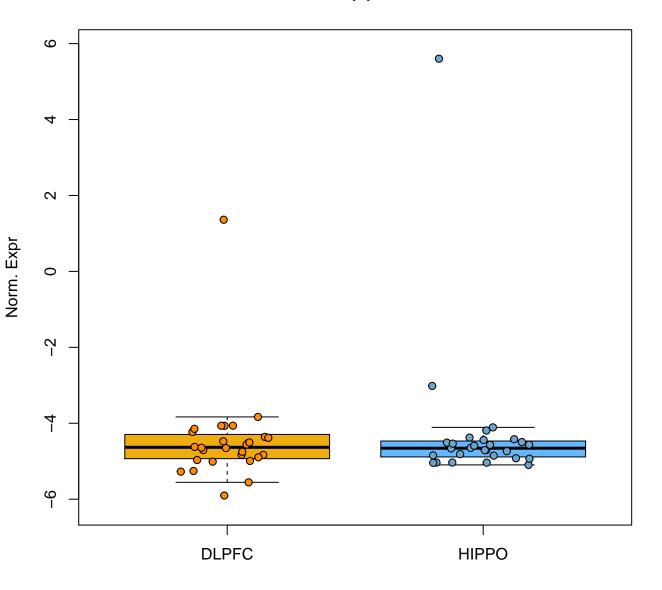
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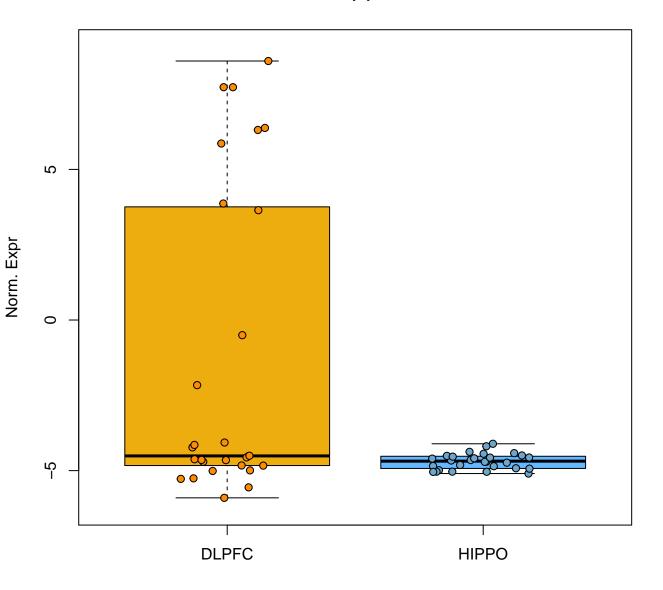
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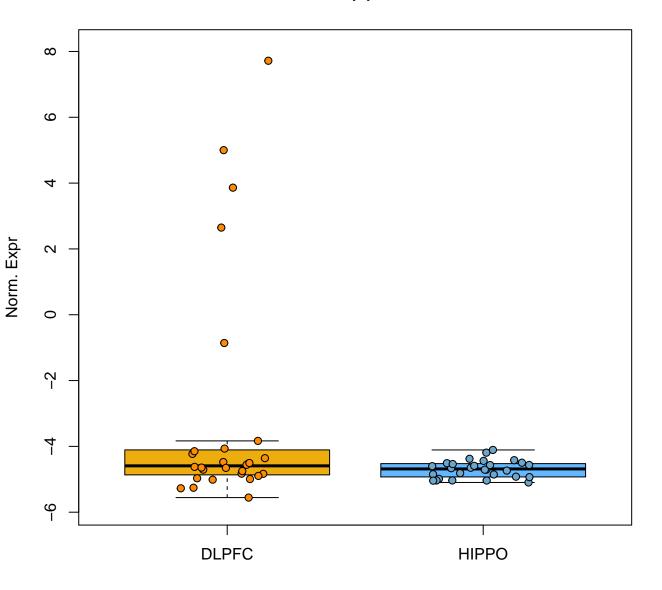
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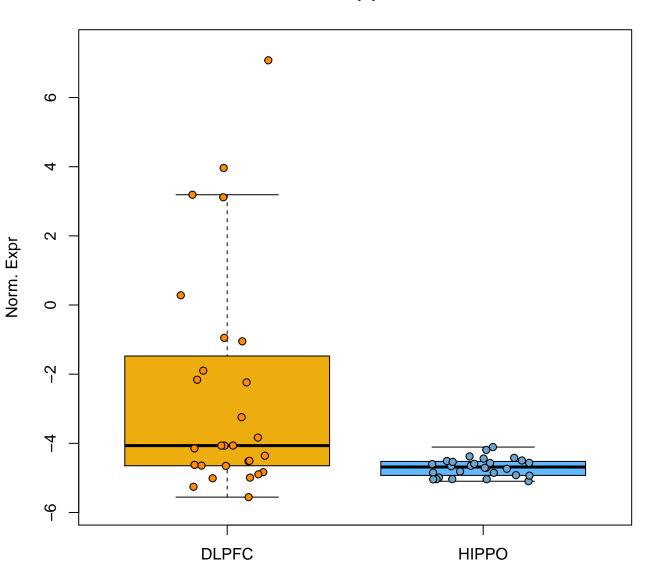
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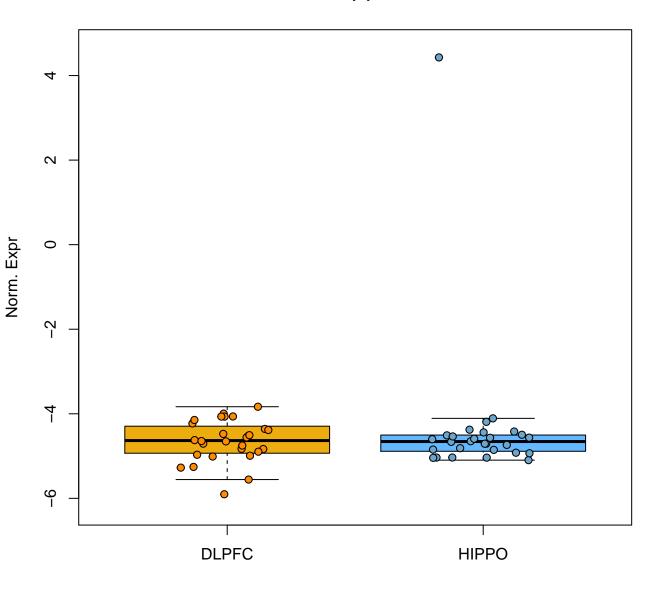
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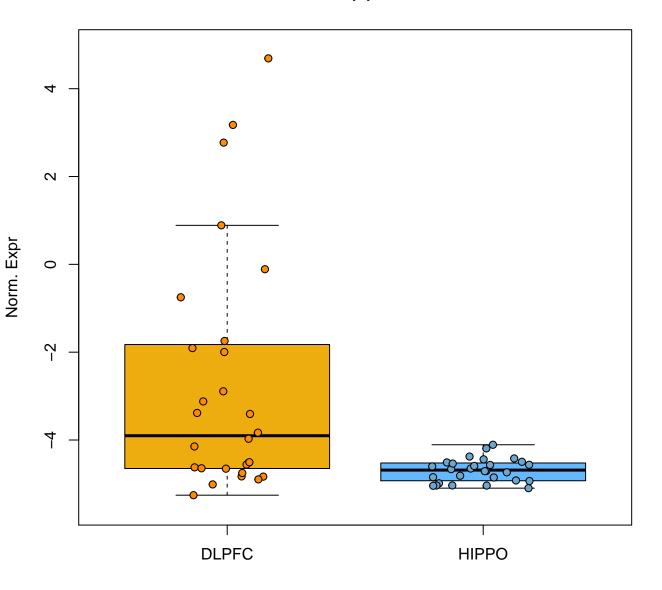
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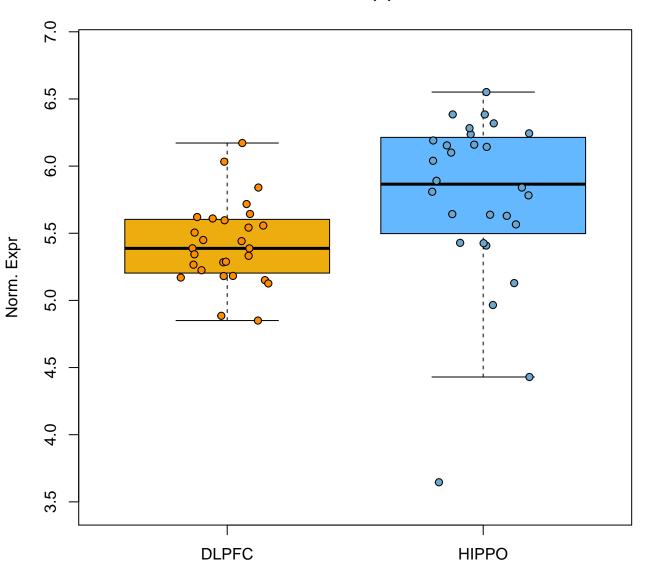
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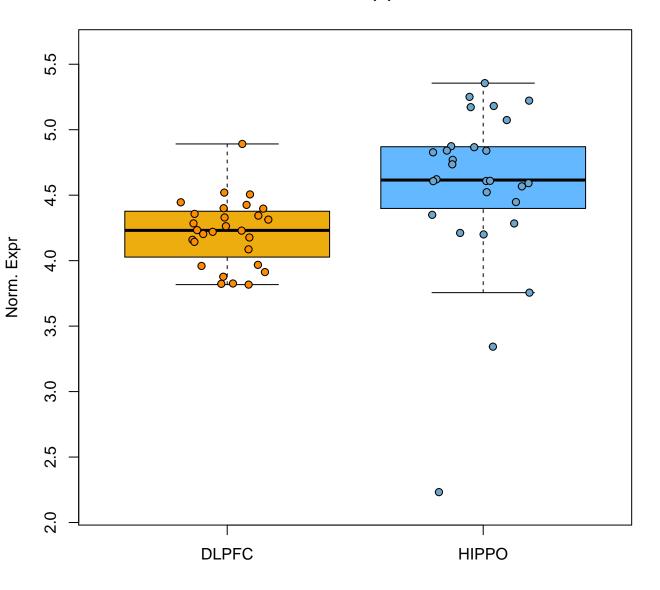
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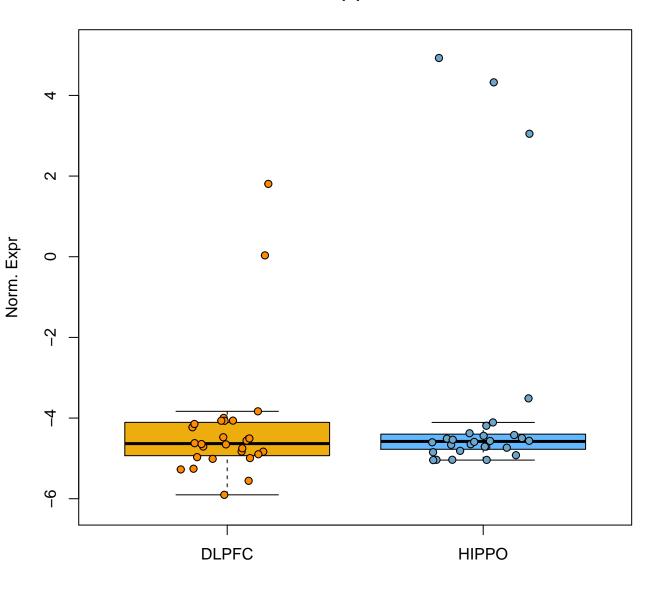
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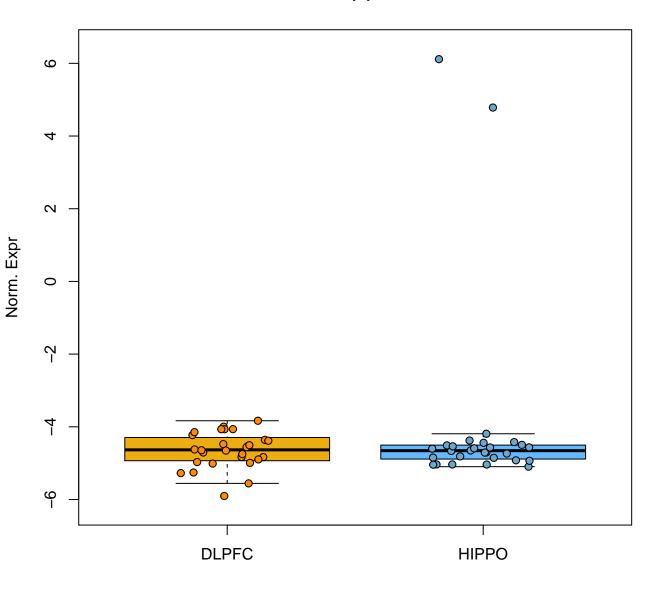
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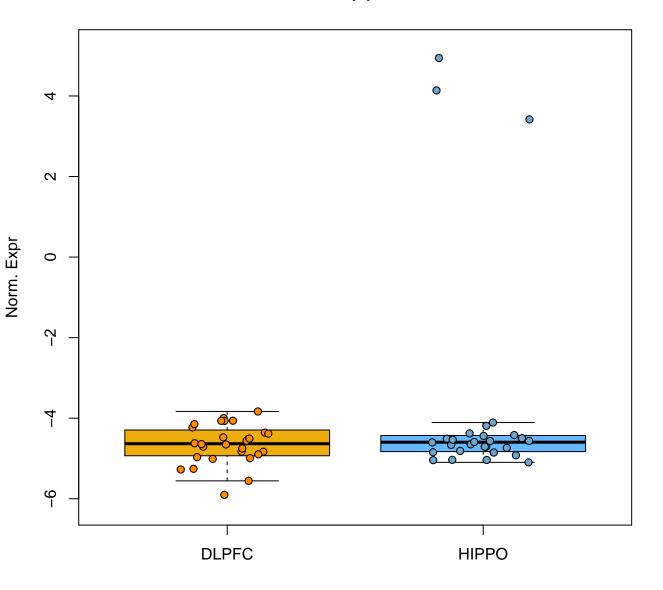
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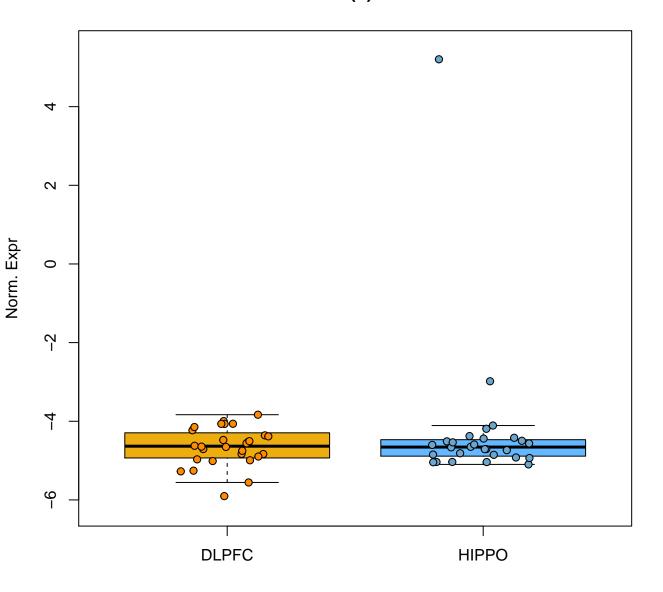
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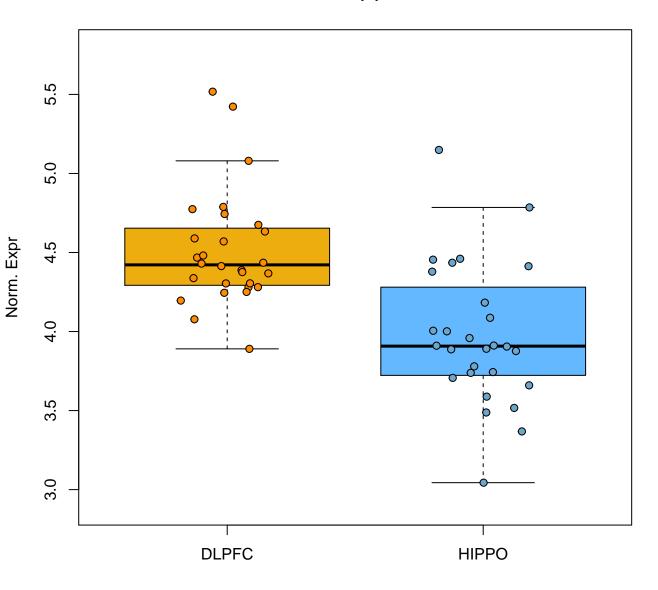
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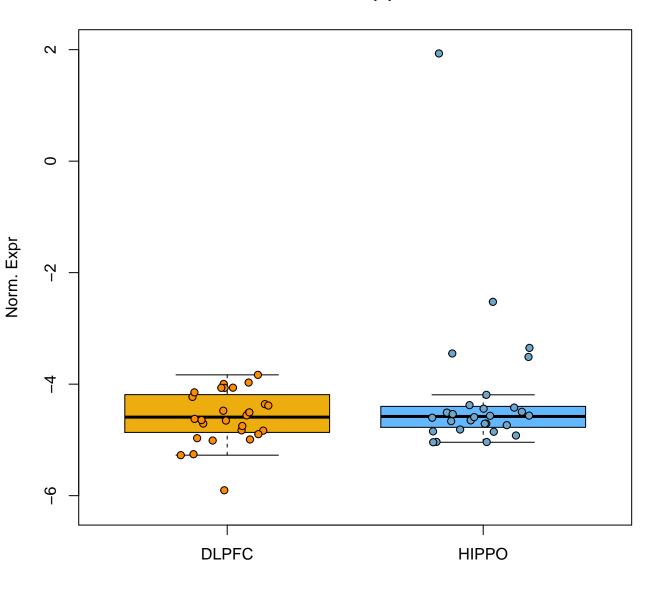
chrM:1769-2260(+) FDR 5.27e-06



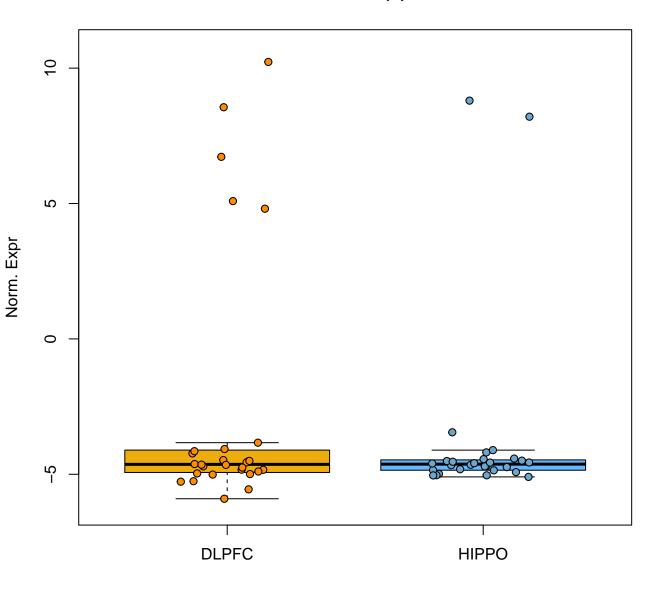
chr12:6917715-6917939(+) ENO2 FDR 5.71e-06



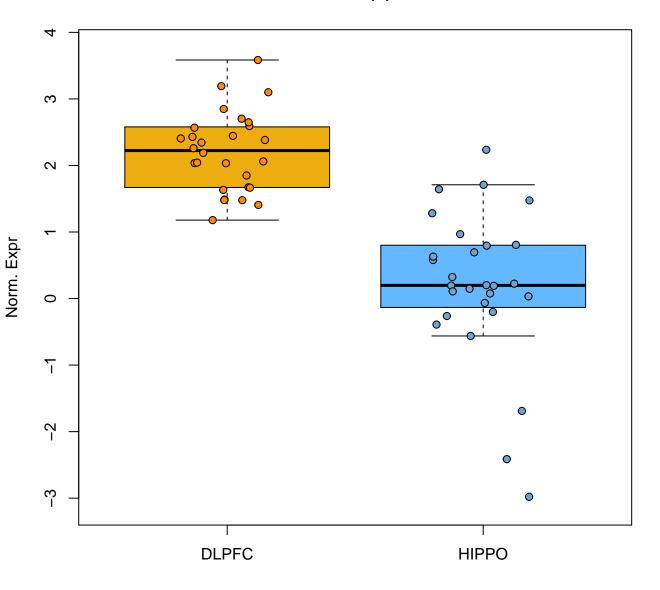
chr11:2166059-2166479(-) TH FDR 6.42e-06



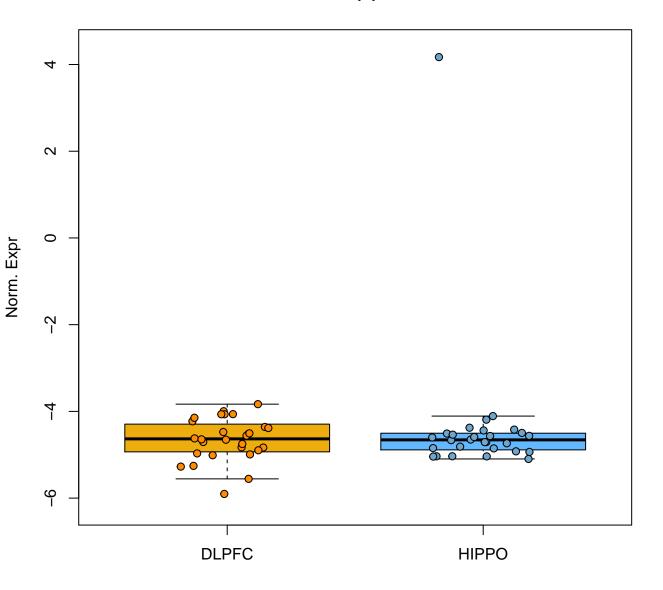
chr21:8401584-8445528(-) FDR 6.99e-06



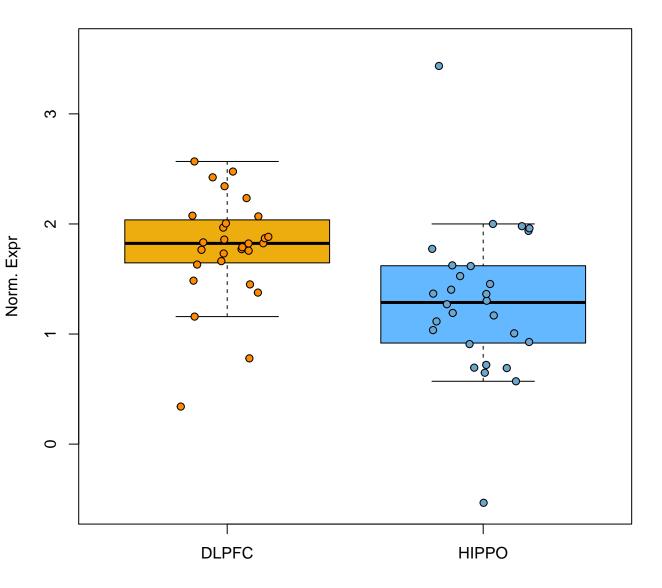
chr9:116976786-117008091(-) ASTN2 FDR 7.12e-06



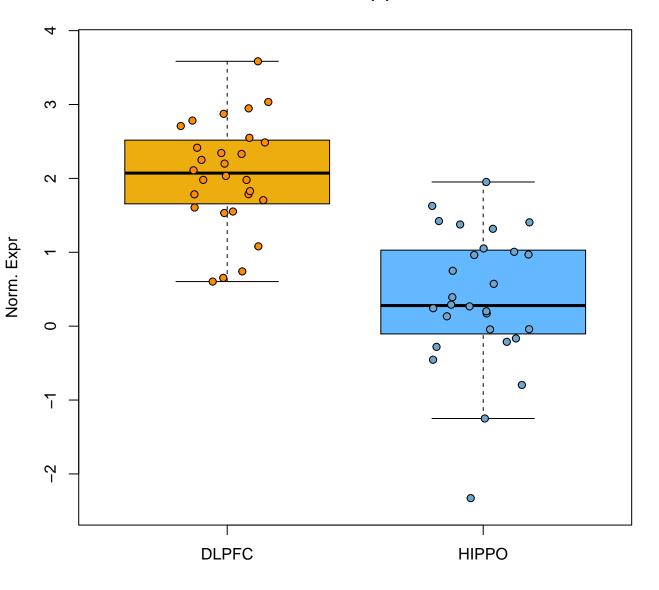
chrM:2167-2244(+) FDR 7.4e-06



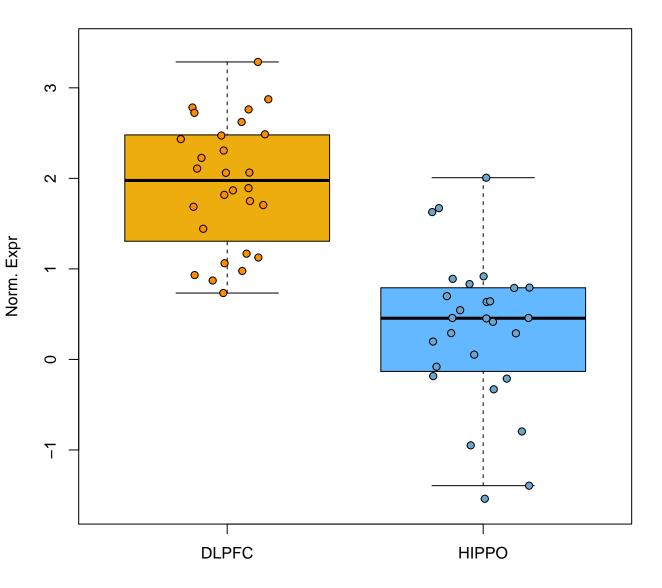
chr13:95599425-95606002(-) DZIP1 FDR 9.15e-06



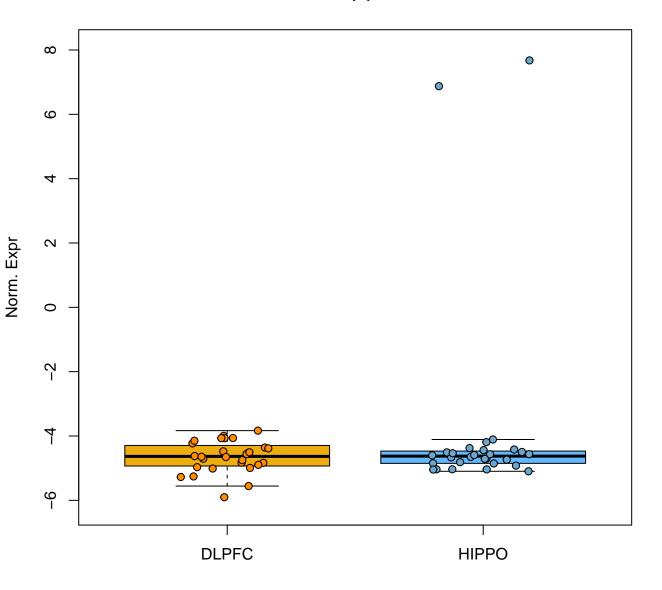
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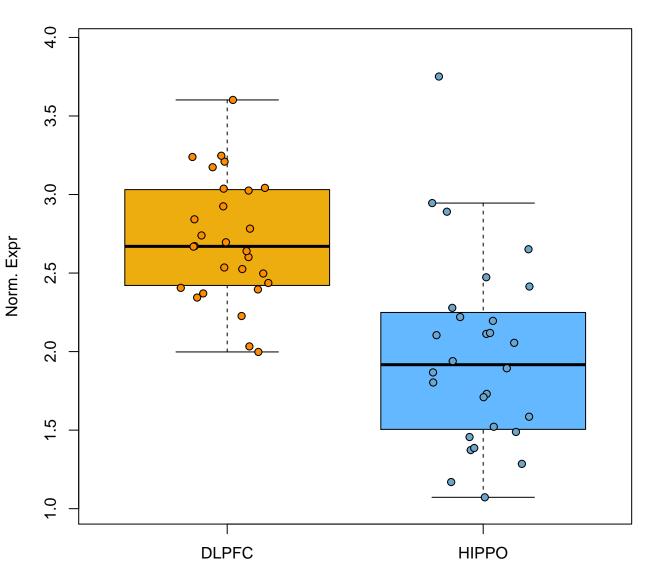
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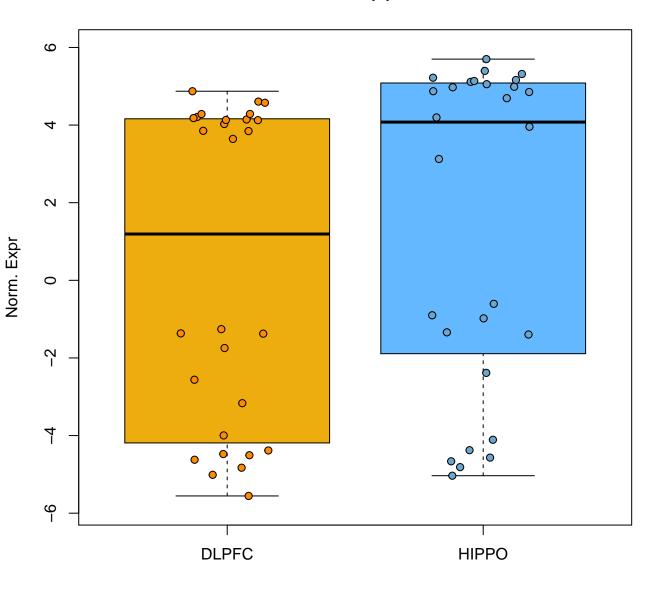
chrM:2160-2354(+) FDR 1.01e-05



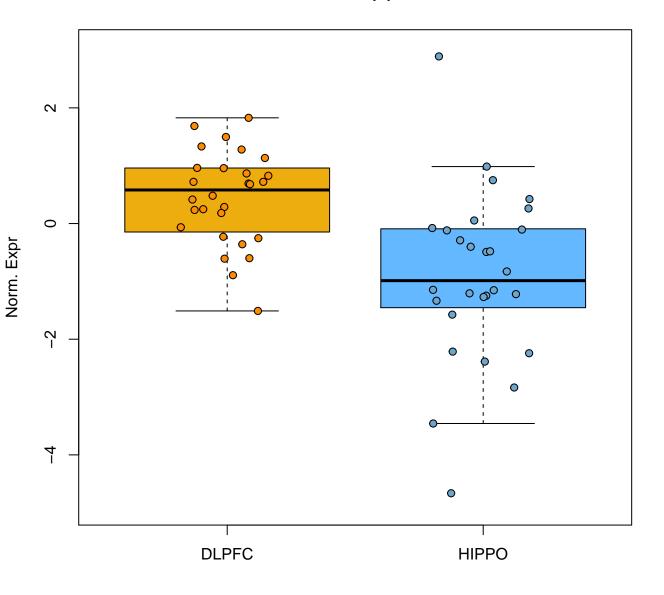
chr7:38432213-38436271(-) AMPH FDR 1.37e-05



chrX:73837504-73841381(-) XIST FDR 1.38e-05



chr8:60266050-60279688(-) CA8 FDR 1.43e-05



chr10:24589303-24591224(-) ARHGAP21 FDR 1.47e-05

