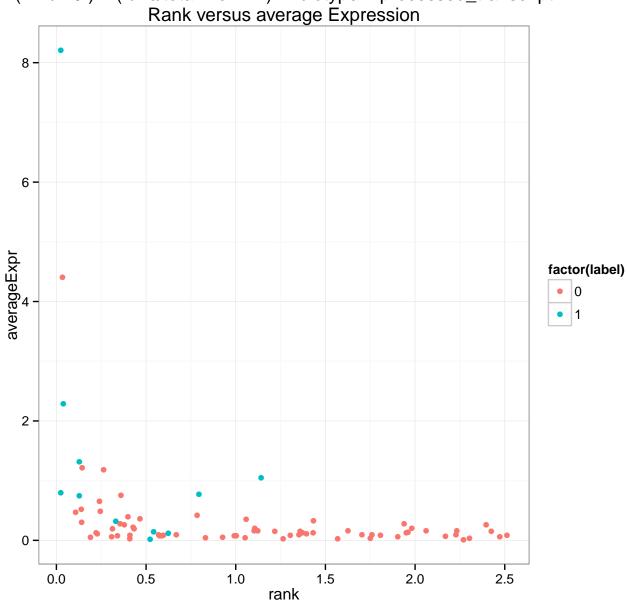
lncRNA group = IDRlessthan0_01 ::: data cols = Inpa (11 / 79) = (func/total lncRNA) ::: biotype = processed_transcript



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