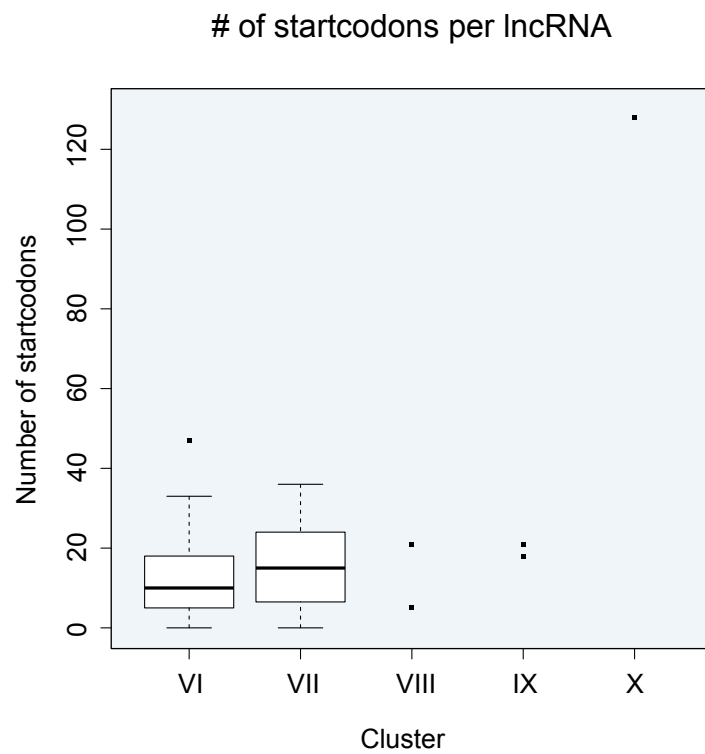




Additional file 8) Number of startcodons per lncRNA per cluster. For 5 ribosomal clusters, the number of potential ORFs is determined by counting the ATG translational start sites per transcript.