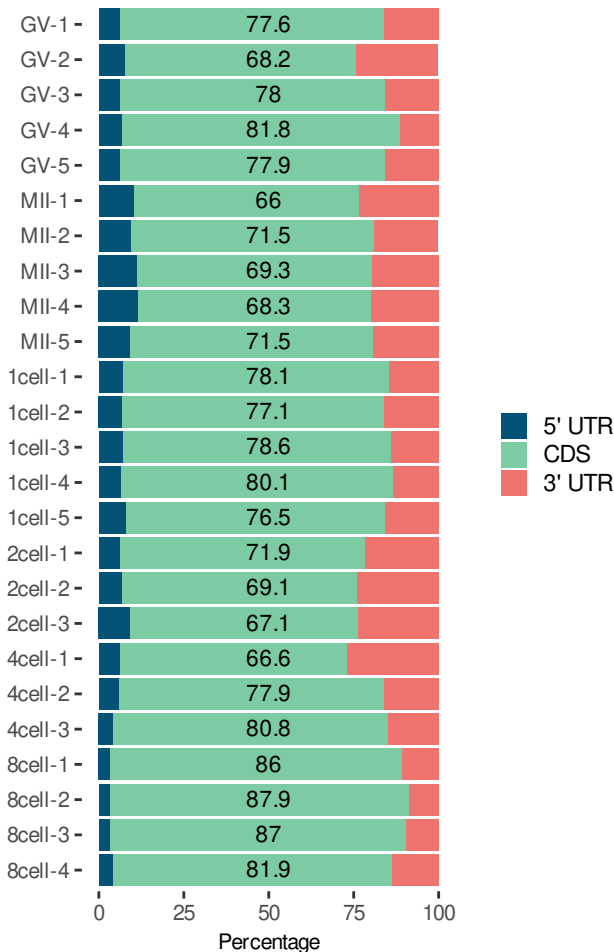Distribution of RPFs by transcript regions

Experiment	5' UTR	CDS	3' UTR
GV-1		77.6	
GV-2		68.2	
GV-3		78	
GV-4		81.8	
GV-5		77.9	
MII-1		66	
MII-2		71.5	
MII-3		69.3	
MII-4		68.3	
MII-5		71.5	
1cell-1		78.1	
1cell-2		77.1	
1cell-3		78.6	
1cell-4		80.1	
1cell-5		76.5	
2cell-1		71.9	
2cell-2		69.1	
2cell-3		67.1	
4cell-1		66.6	
4cell-2		77.9	
4cell-3		80.8	
8cell-1		86	
8cell-2		87.9	
8cell-3		87	
8cell-4		81.9	