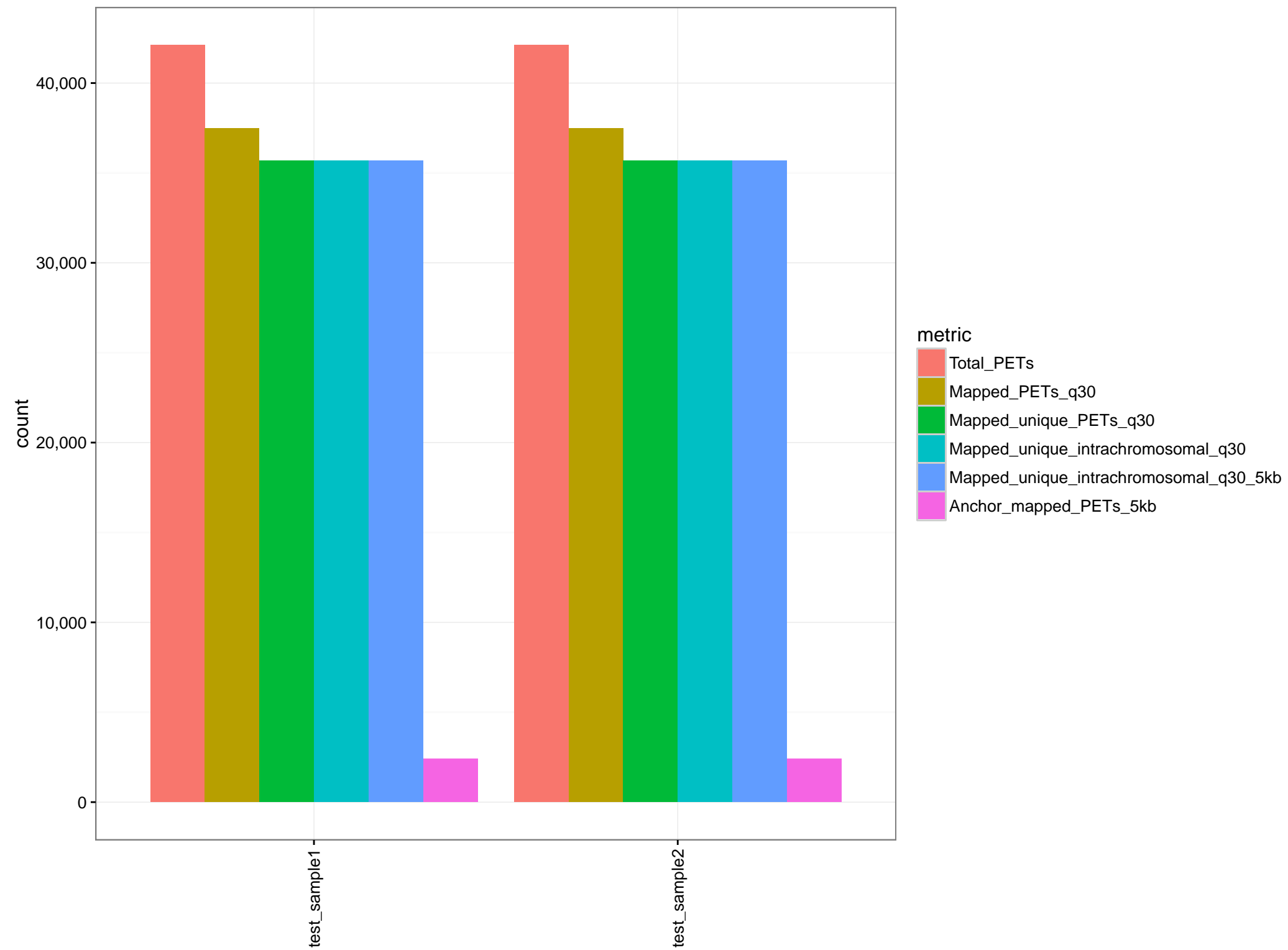


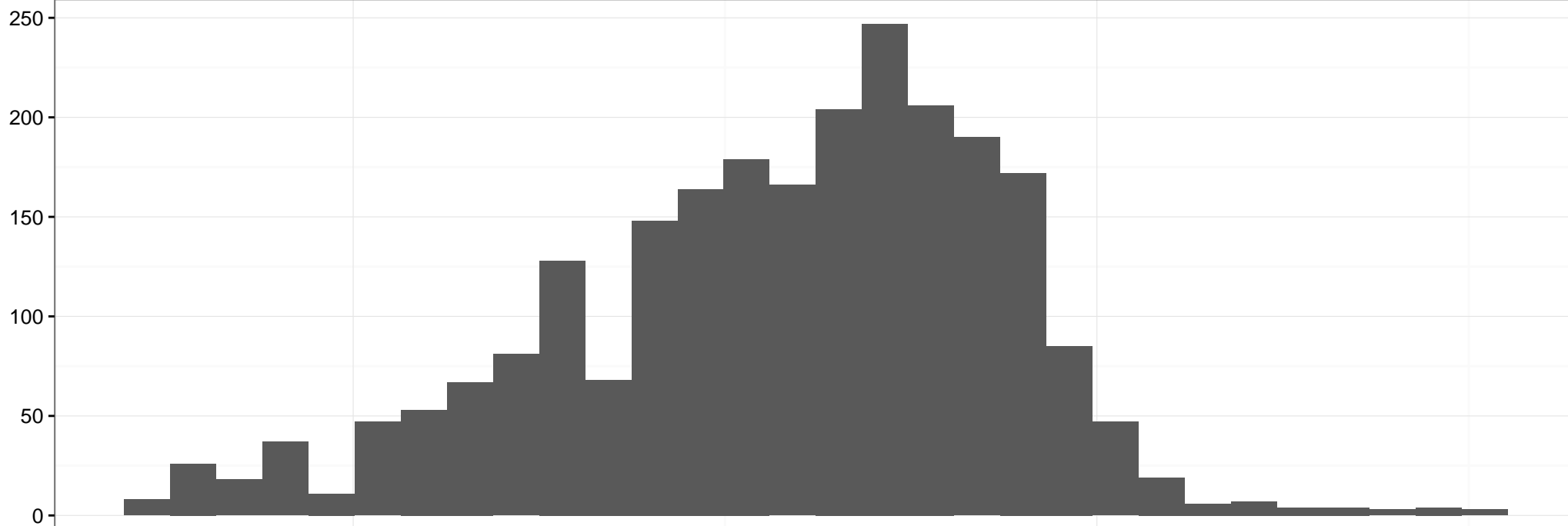
Counts



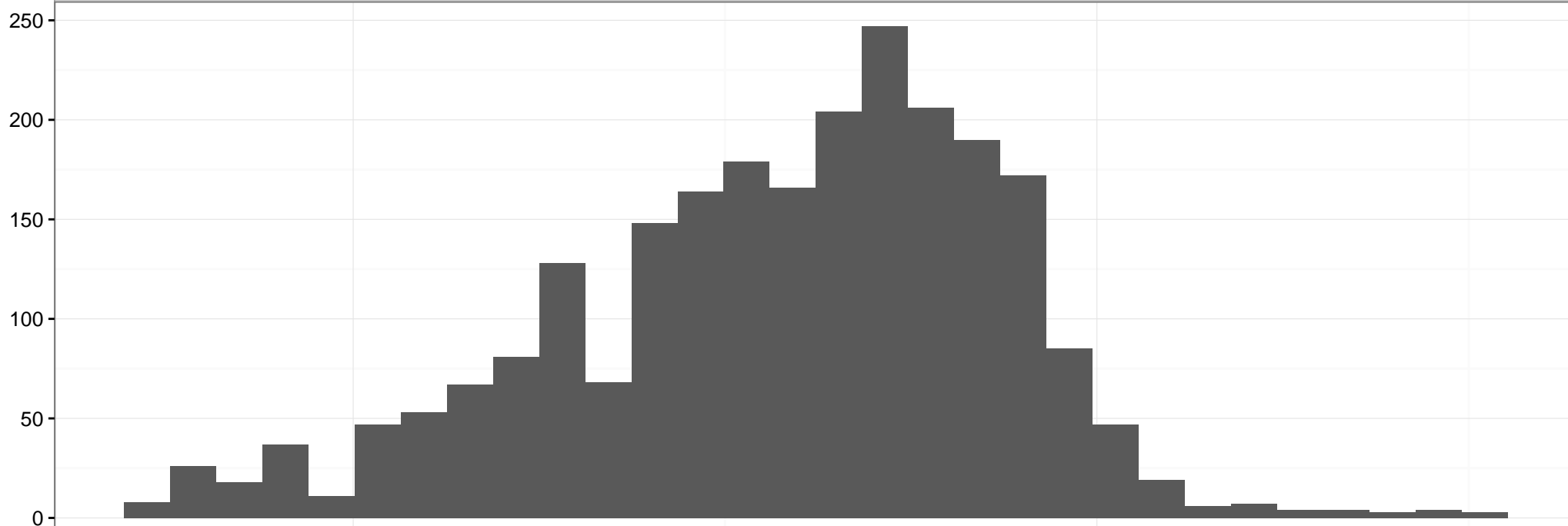
	test_sample1	test_sample2
<i>Total_PETs</i>	42,105	42,105
<i>Mapped_PETs_q30</i>	37,490	37,490
<i>Mapped_unique_PETs_q30</i>	35,674	35,674
<i>Mapped_unique_intrachromosomal_q30</i>	35,674	35,674
<i>Mapped_unique_intrachromosomal_q30_5kb</i>	35,674	35,674
<i>Anchor_mapped_PETs_5kb</i>	2,402	2,402

	test_sample1	test_sample2
<i>Total_PETs</i>	100.00	100.00
<i>Mapped_PETs_q30</i>	89.04	89.04
<i>Mapped_unique_PETs_q30</i>	84.73	84.73
<i>Mapped_unique_intrachromosomal_q30</i>	84.73	84.73
<i>Mapped_unique_intrachromosomal_q30_5kb</i>	84.73	84.73
<i>Anchor_mapped_PETs_5kb</i>	5.70	5.70

test_sample1



test_sample2



loop_length