

Genomic map showing the distribution of tRNA and LTR elements across a genomic region. The x-axis represents the Base position (0 to 250,000). A dashed horizontal line separates the top and bottom halves of the map. Black bars represent tRNA and LTR elements, with their names and sizes labeled above them. Green text labels identify specific elements: tRNA, LTR, and R063. A red dot is located at the bottom left of the map.

Element Name	Base Position (approx.)	Size (bp)	Type
1613*	10,000	100	tRNA
1100*	15,000	100	tRNA
1822	20,000	100	tRNA
2125*	25,000	100	tRNA
1259*	55,000	100	tRNA
2287	90,000	100	tRNA
1744*	120,000	100	tRNA
1938*	125,000	100	tRNA
2018	140,000	100	tRNA
2256*	155,000	100	tRNA
1616	165,000	100	tRNA
1302	170,000	100	tRNA
2112	175,000	100	tRNA
1278	190,000	100	tRNA
2296	245,000	100	tRNA
2189	10,000	100	tRNA
1885	25,000	100	tRNA
1398*	35,000	100	tRNA
2470	40,000	100	tRNA
1819	100,000	100	tRNA
2277*	110,000	100	tRNA
1791	125,000	100	tRNA
1237	135,000	100	tRNA
1851	145,000	100	tRNA
1351*	140,000	100	tRNA
2130*	165,000	100	tRNA
1806*	170,000	100	tRNA
1283*	245,000	100	tRNA
2489*	180,000	100	tRNA
2125*	25,000	100	tRNA
1938*	125,000	100	tRNA
1616	165,000	100	tRNA
1302	170,000	100	tRNA
2112	175,000	100	tRNA
1278	190,000	100	tRNA
2296	245,000	100	tRNA
2189	10,000	100	tRNA
1885	25,000	100	tRNA
1398*	35,000	100	tRNA
2470	40,000	100	tRNA
1819	100,000	100	tRNA
2277*	110,000	100	tRNA
1791	125,000	100	tRNA
1237	135,000	100	tRNA
1851	145,000	100	tRNA
1351*	140,000	100	tRNA
2130*	165,000	100	tRNA
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1283*	245,000	100	tRNA
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1885	25,000	100	tRNA
1398*	35,000	100	tRNA
2470	40,000	100	tRNA
1819	100,000	100	tRNA
2277*	110,000	100	