

Dataset stats

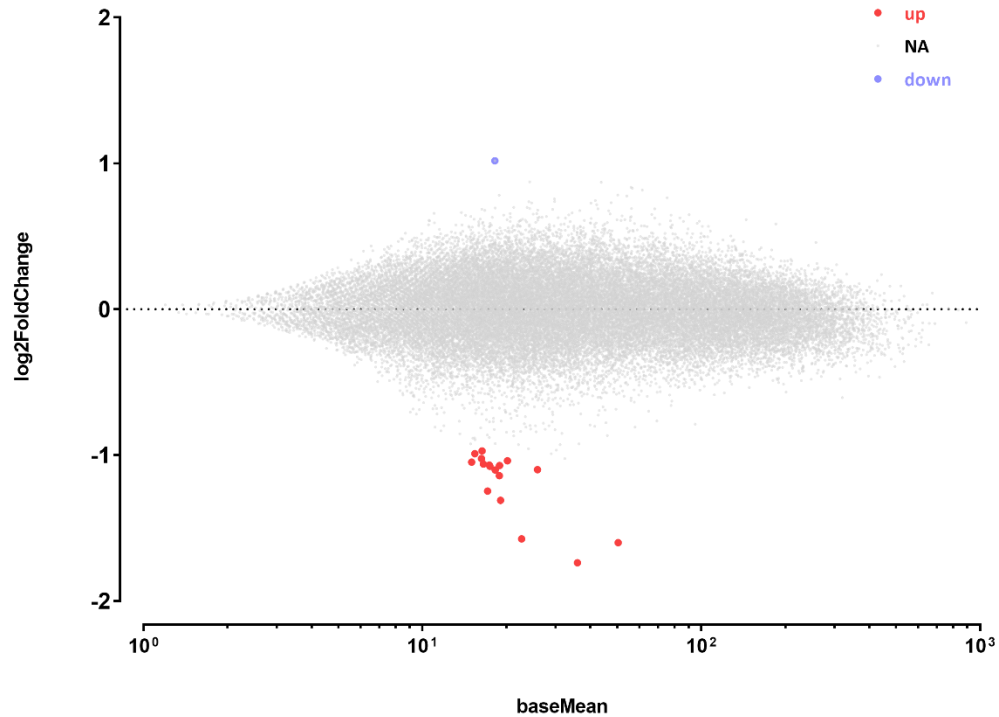
Species	Read Length	Library	Raw fragments	Unique non-chrM reads	Multi reads	Complexity	chrM reads	chrM fraction	Unique non-chrM reads after dedup	TSS ratio	MACS default peaks	FRiP (MACS)
<i>Homo sapiens</i> hg38-female	2x36	2019-07-16-MyoD-0hr-A1	15,148,615	13,042,232		0.83	3,524,400	0.21	11,485,906	16.22	30,398	295,942
<i>Homo sapiens</i> hg38-female	2x36	2019-07-16-MyoD-0hr-B1	14,317,815	11,532,368		0.81	3,465,618	0.23	9,999,578	18.72	29,955	331,050
<i>Homo sapiens</i> hg38-female	2x36	2019-07-16-MyoD-48hr-C1	10,334,712	8,120,012		0.82	2,456,018	0.23	7,053,030	18.75	28,630	329,840
<i>Homo sapiens</i> hg38-female	2x36	2019-07-16-MyoD-48hr-D1	8,729,615	6,715,900		0.82	1,831,020	0.21	5,802,154	19.49	25,151	340,275

Differential sites

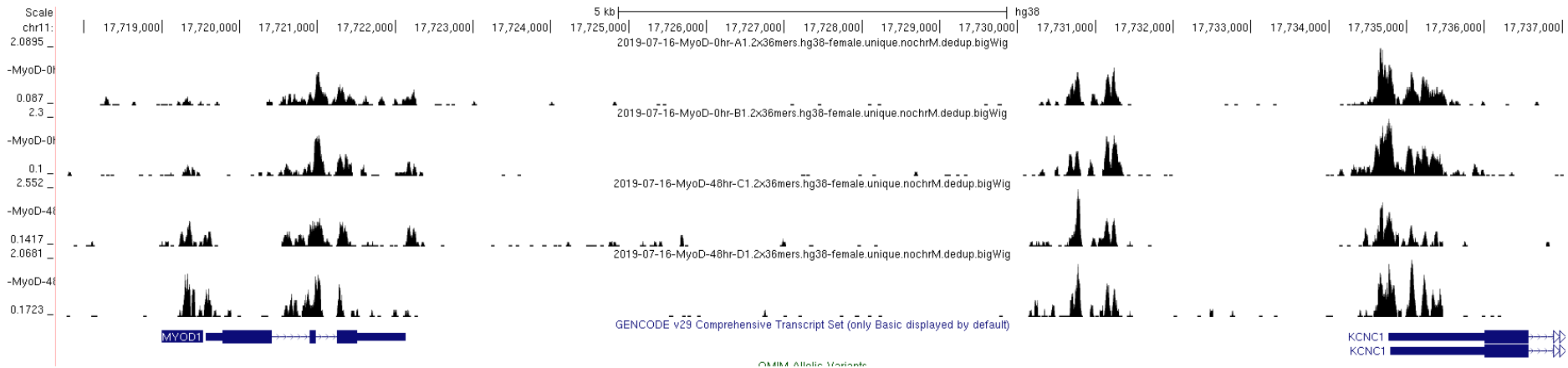
Number differential ATAC peaks

A	B	p-adj	
		up in B vs A	down in B vs A
MyoD-0hr	MyoD-48hr	19	1

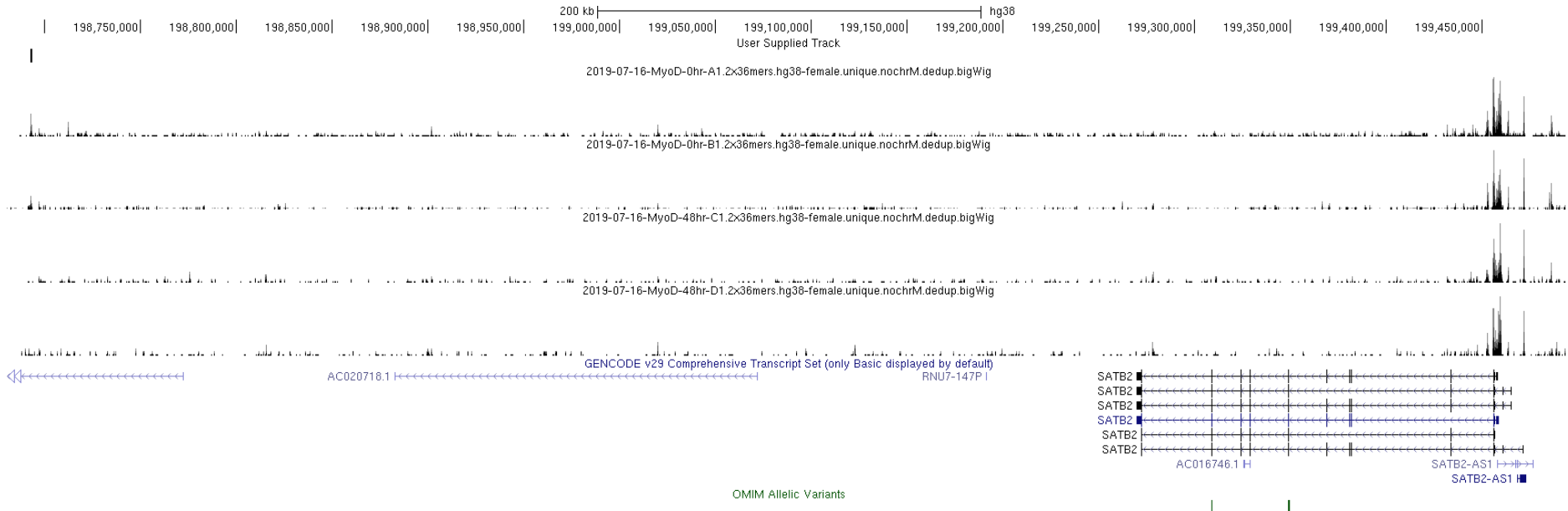
MyoD 0hr vs 48hr

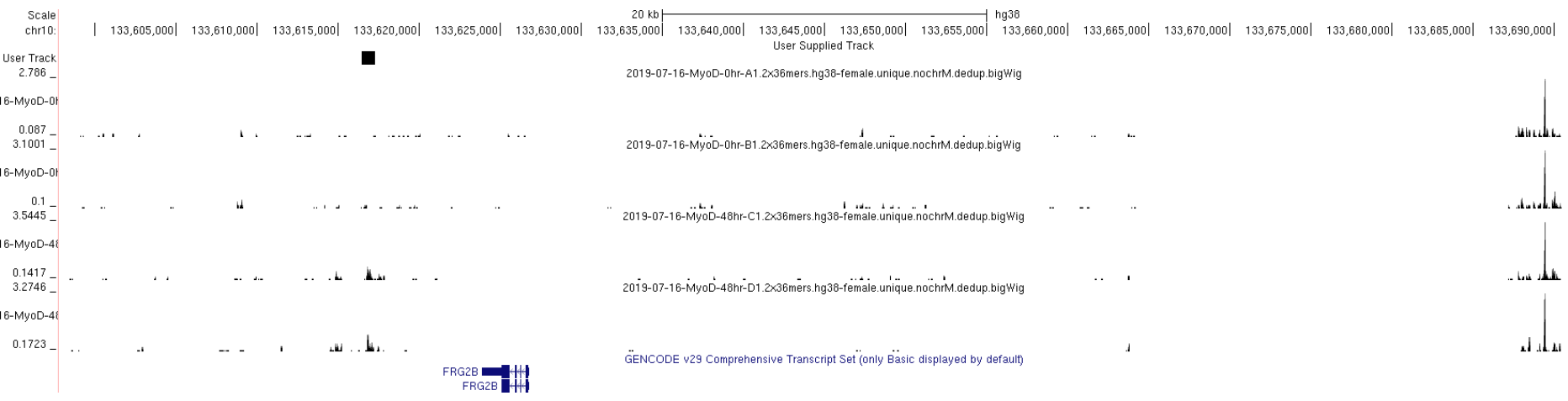


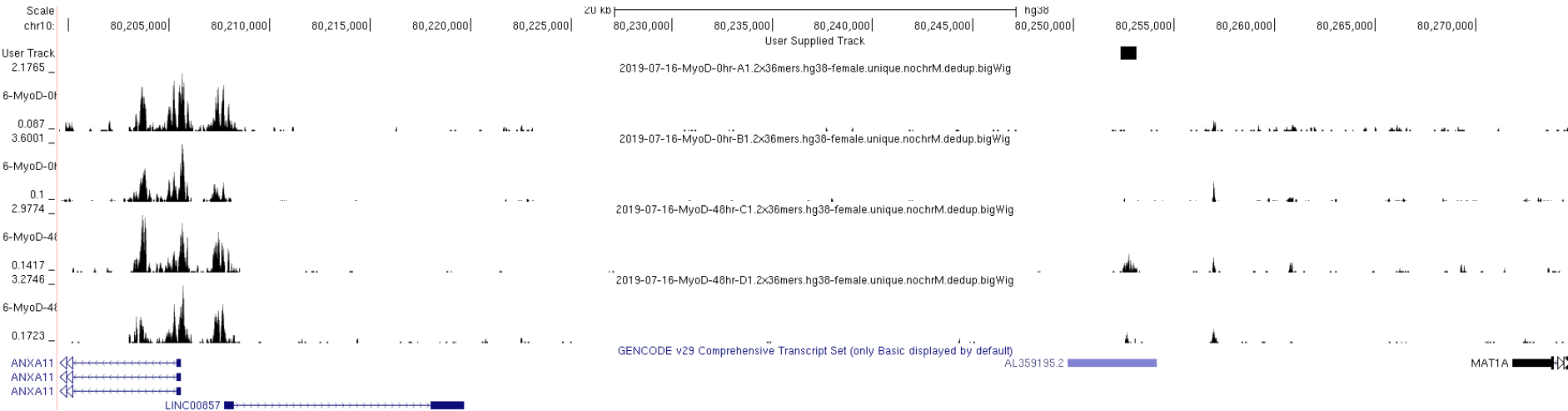
MYOD1

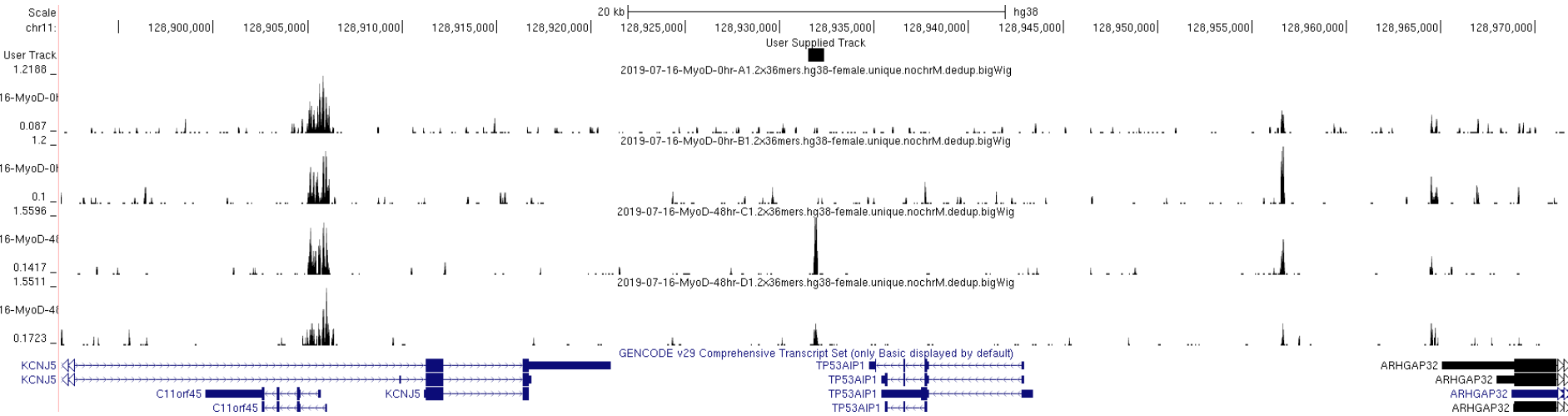


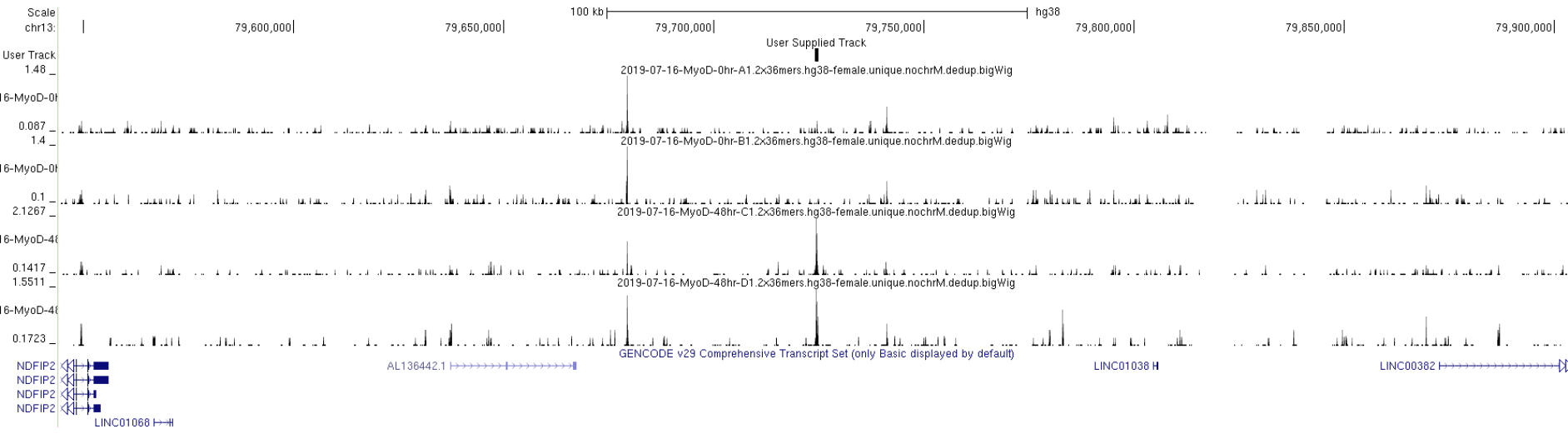
The other sites:

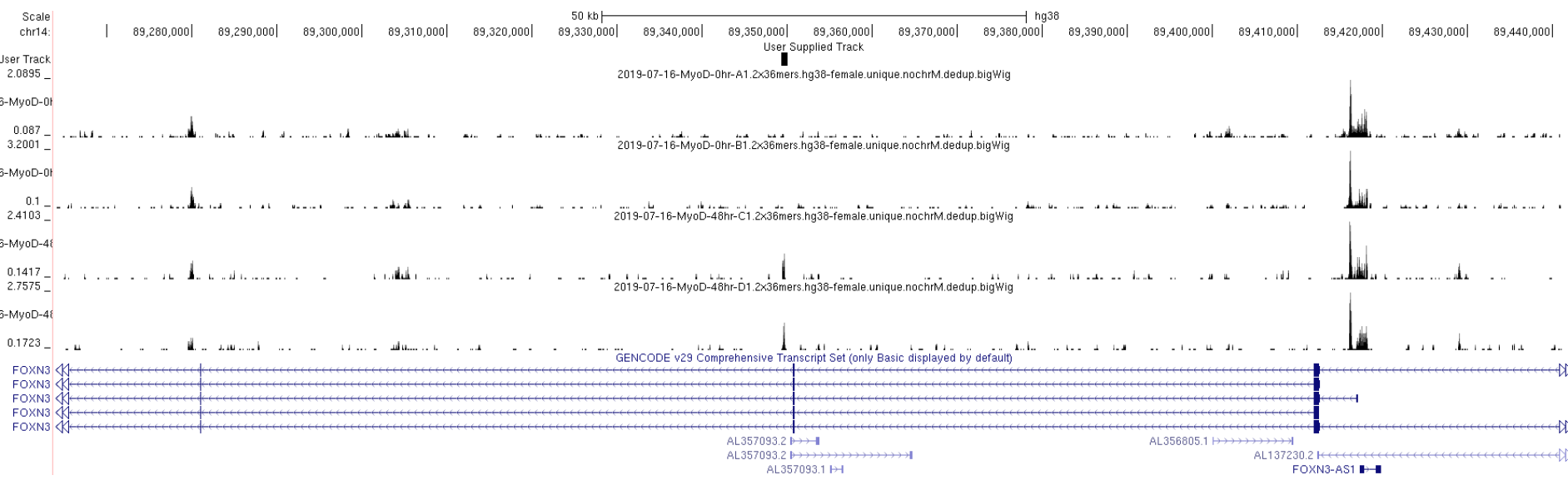


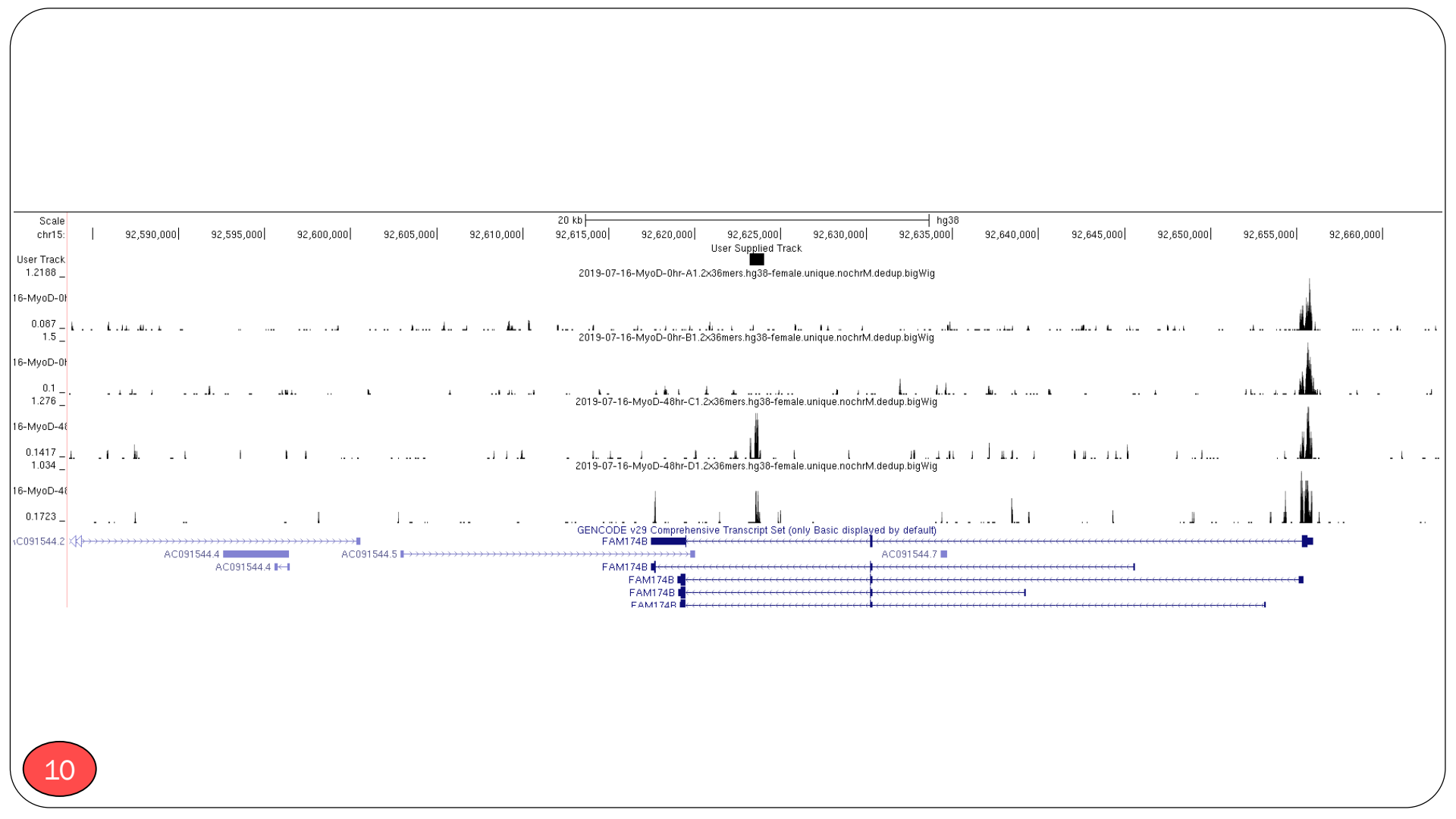


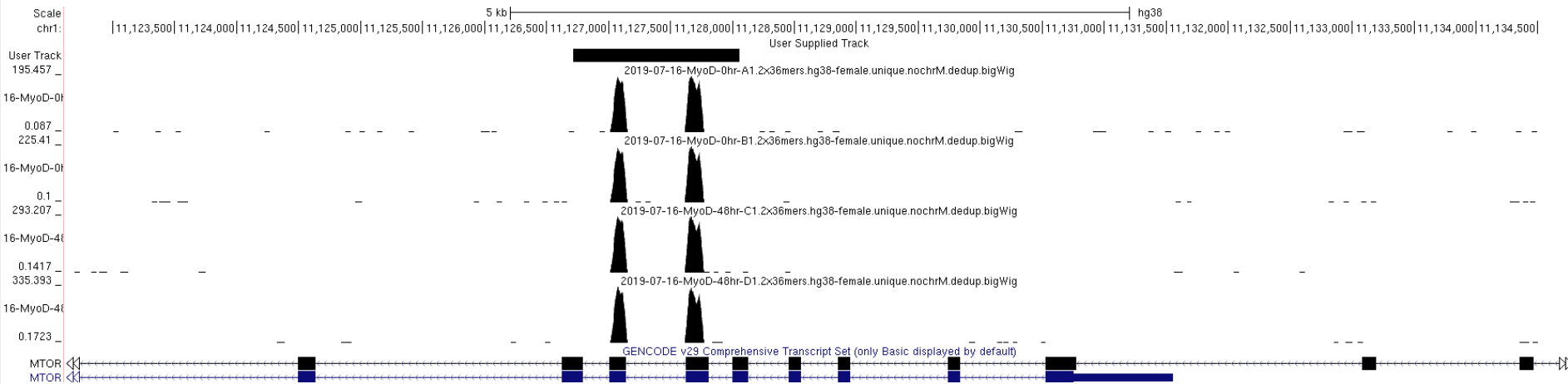












Clear artifact. Does anyone have plasmids with MTOR around the lab?

