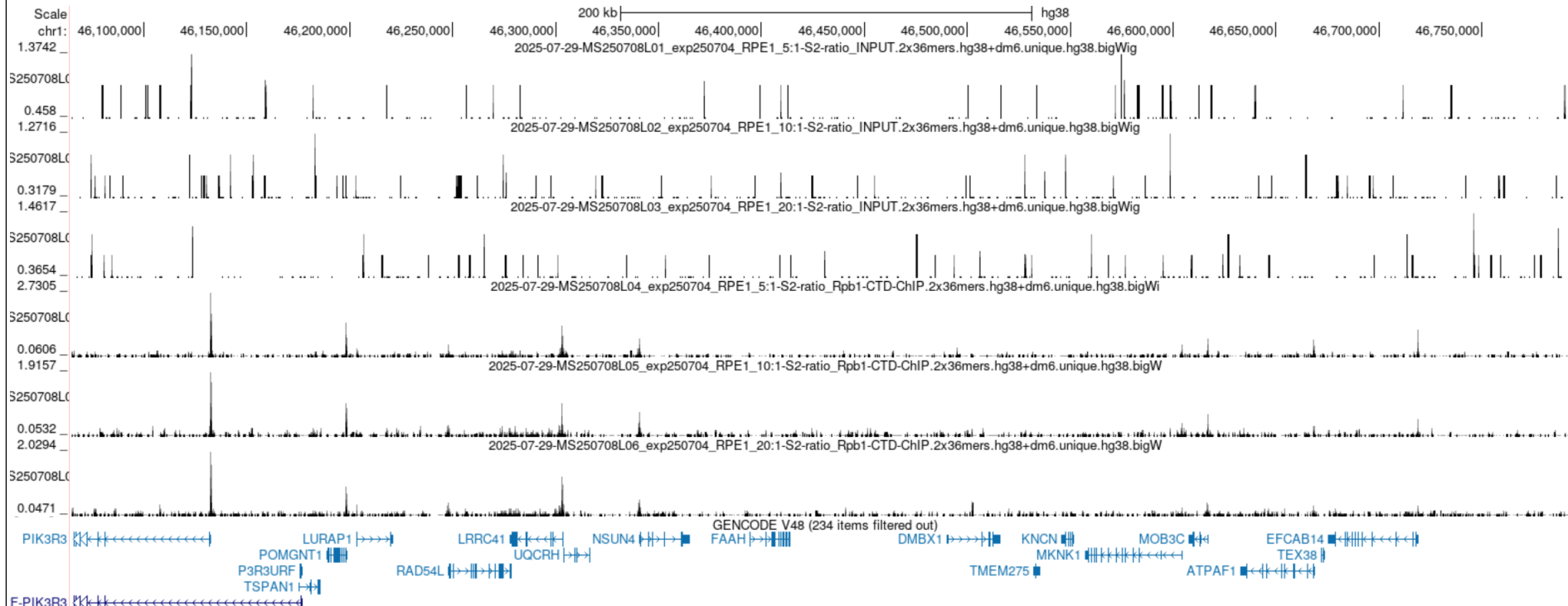


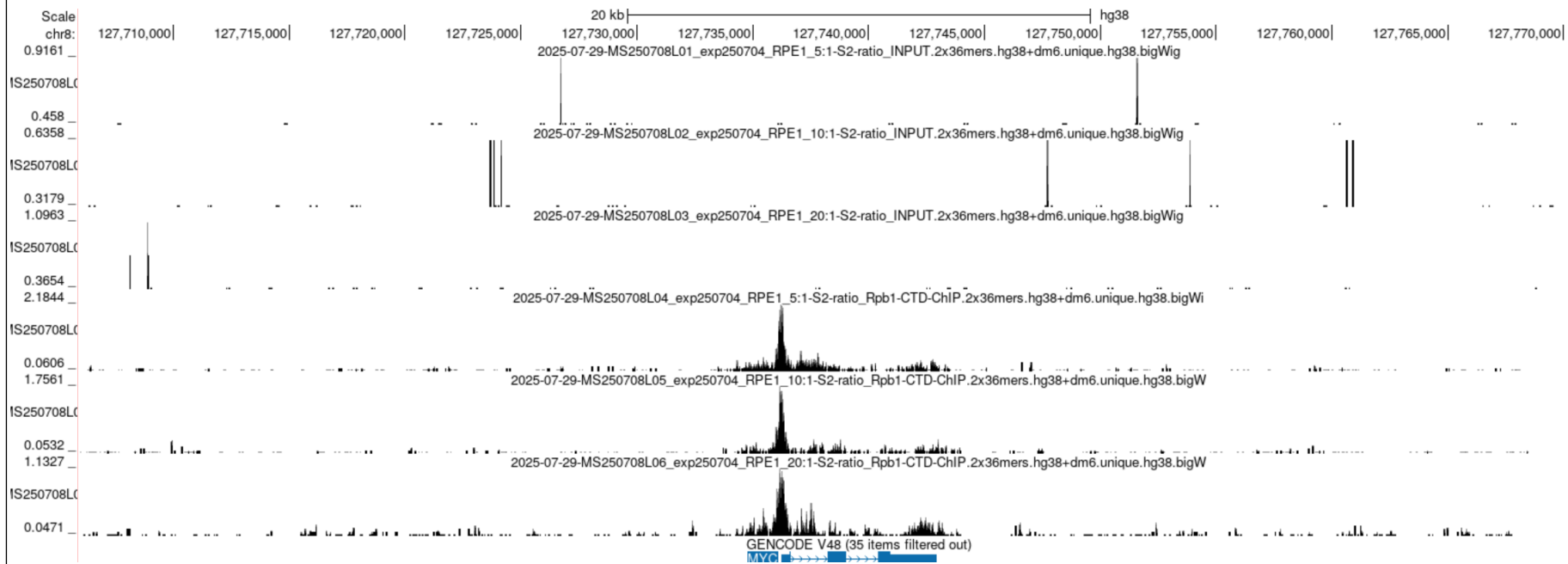
# MAPPING STATS

Library	Raw fragments	Species	2x36mers unique		
			Unique reads	Complexity	Fraction aligned
2025-07-29-MS250708L01_exp250704_RPE1_5:1-S2-ratio_INPUT	1,297,355	dm6	78,646	0.88	0.03
2025-07-29-MS250708L01_exp250704_RPE1_5:1-S2-ratio_INPUT	1,297,355	hg38	2,104,402	0.87	0.81
2025-07-29-MS250708L02_exp250704_RPE1_10:1-S2-ratio_INPUT	1,859,324	dm6	61,132	0.88	0.02
2025-07-29-MS250708L02_exp250704_RPE1_10:1-S2-ratio_INPUT	1,859,324	hg38	3,084,276	0.86	0.83
2025-07-29-MS250708L03_exp250704_RPE1_20:1-S2-ratio_INPUT	1,613,900	dm6	27,400	0.87	0.01
2025-07-29-MS250708L03_exp250704_RPE1_20:1-S2-ratio_INPUT	1,613,900	hg38	2,709,058	0.86	0.84
2025-07-29-MS250708L04_exp250704_RPE1_5:1-S2-ratio_Rpb1-CTD-ChIP	9,890,048	dm6	752,446	0.83	0.04
2025-07-29-MS250708L04_exp250704_RPE1_5:1-S2-ratio_Rpb1-CTD-ChIP	9,890,048	hg38	15,727,864	0.83	0.80
2025-07-29-MS250708L05_exp250704_RPE1_10:1-S2-ratio_Rpb1-CTD-ChIP	11,143,271	dm6	404,622	0.84	0.02
2025-07-29-MS250708L05_exp250704_RPE1_10:1-S2-ratio_Rpb1-CTD-ChIP	11,143,271	hg38	18,386,866	0.82	0.83
2025-07-29-MS250708L06_exp250704_RPE1_20:1-S2-ratio_Rpb1-CTD-ChIP	12,520,371	dm6	256,248	0.84	0.01
2025-07-29-MS250708L06_exp250704_RPE1_20:1-S2-ratio_Rpb1-CTD-ChIP	12,520,371	hg38	20,931,460	0.83	0.84

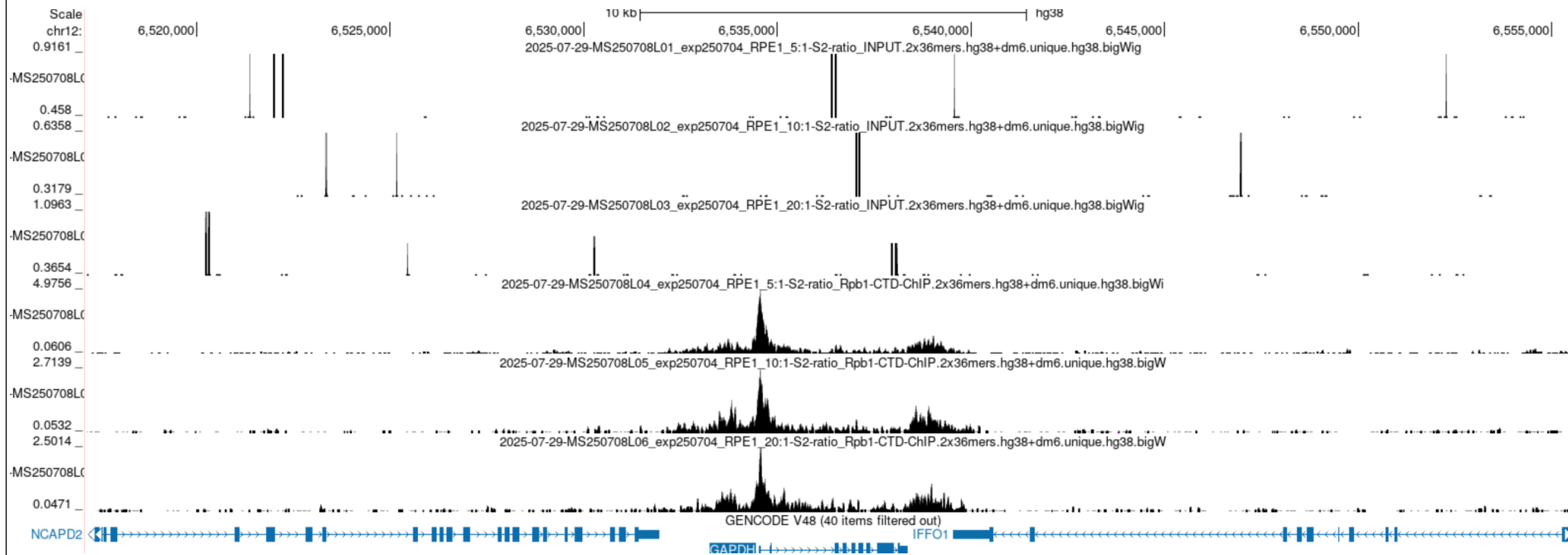
# BROWSER SNAPSHOTS, HUMAN



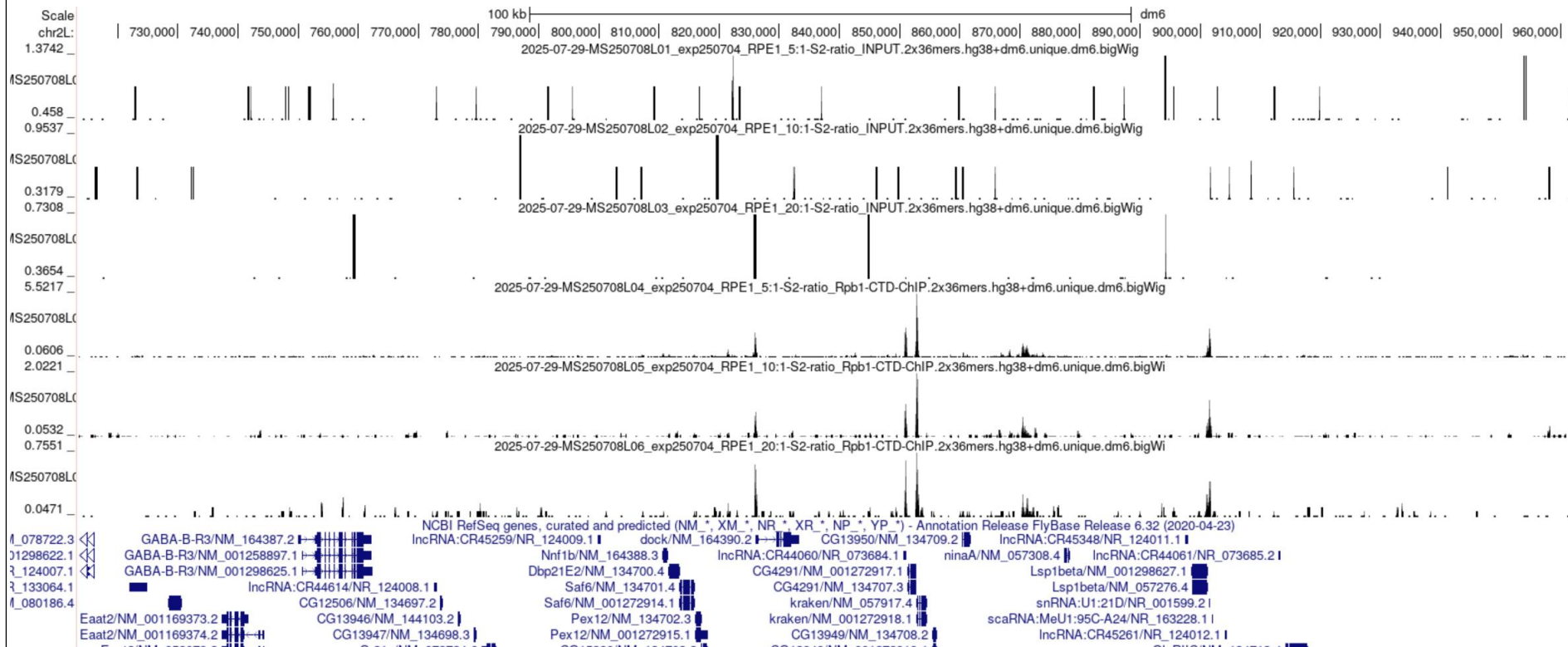
# BROWSER SNAPSHOTS, HUMAN



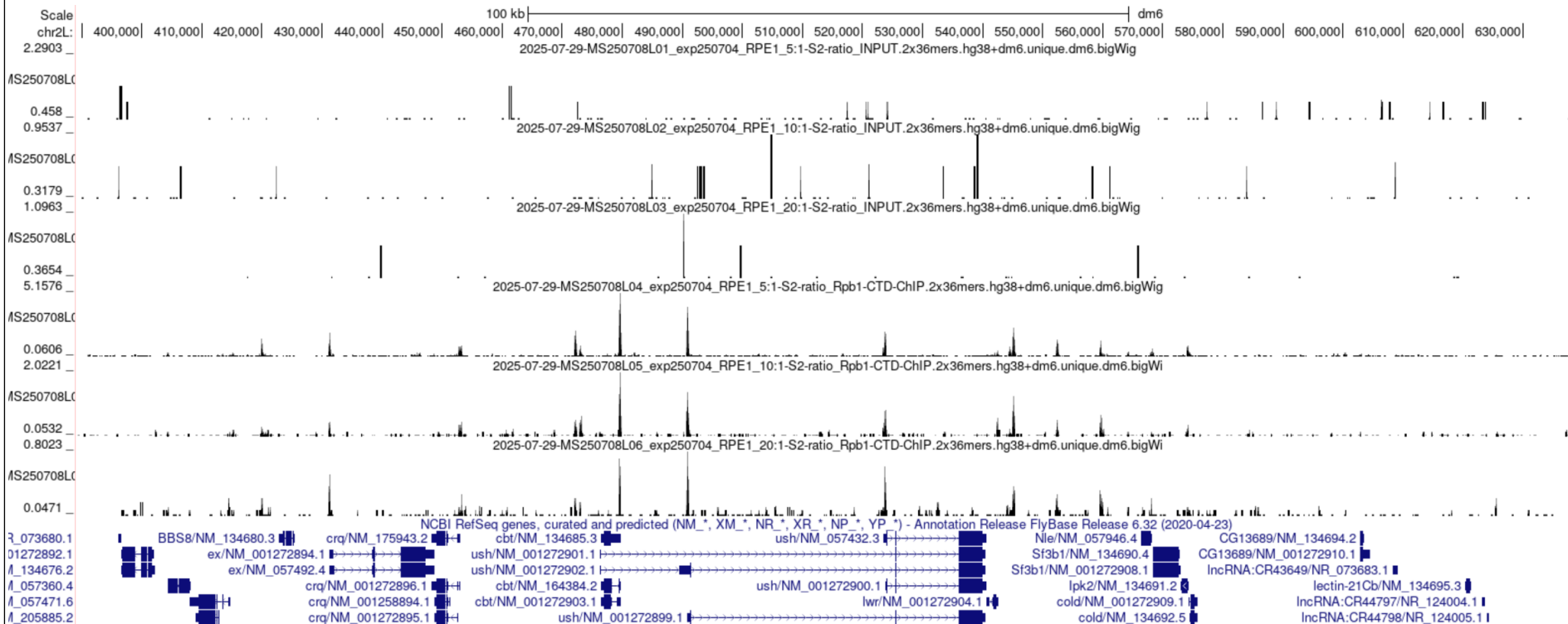
# BROWSER SNAPSHOTS, HUMAN



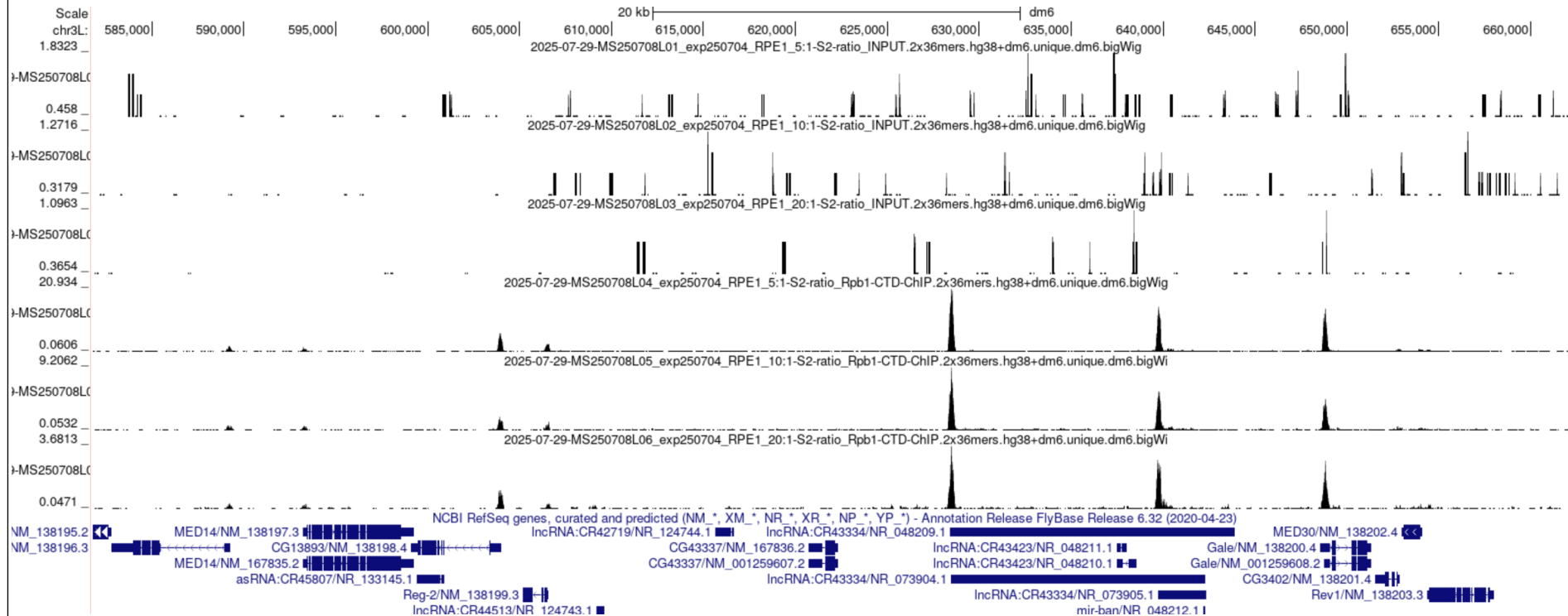
# BROWSER SNAPSHOTS, DROSOPHILA



# BROWSER SNAPSHOTS, DROSOPHILA

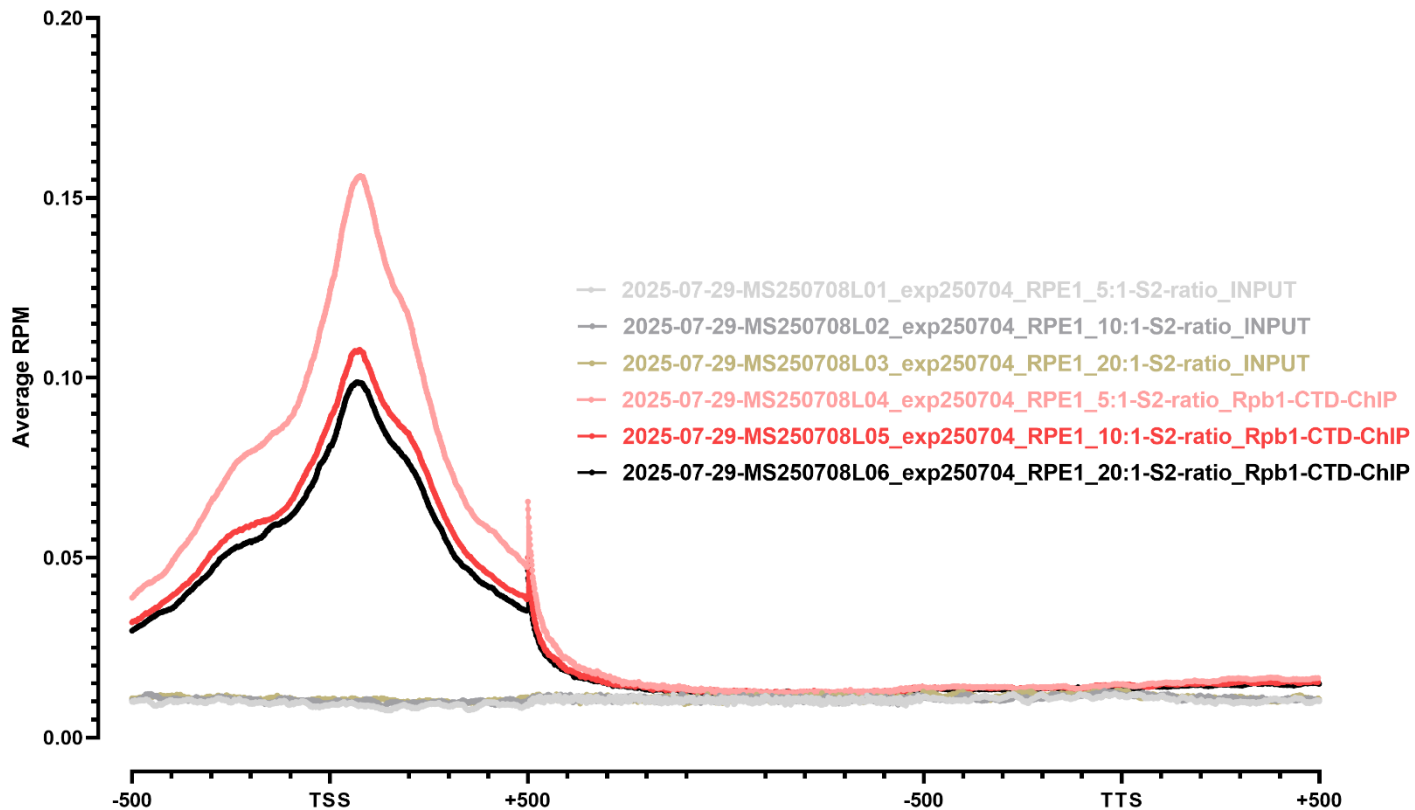


# BROWSER SNAPSHOTS, DROSOPHILA



# METAGENE PROFILES

hg38



# METAGENE PROFILES

dm6

