

Figure S7. Visualization of the mapping of the long-reads on the ESM015 and EP155 genome assemblies around the major detected synteny breaks.

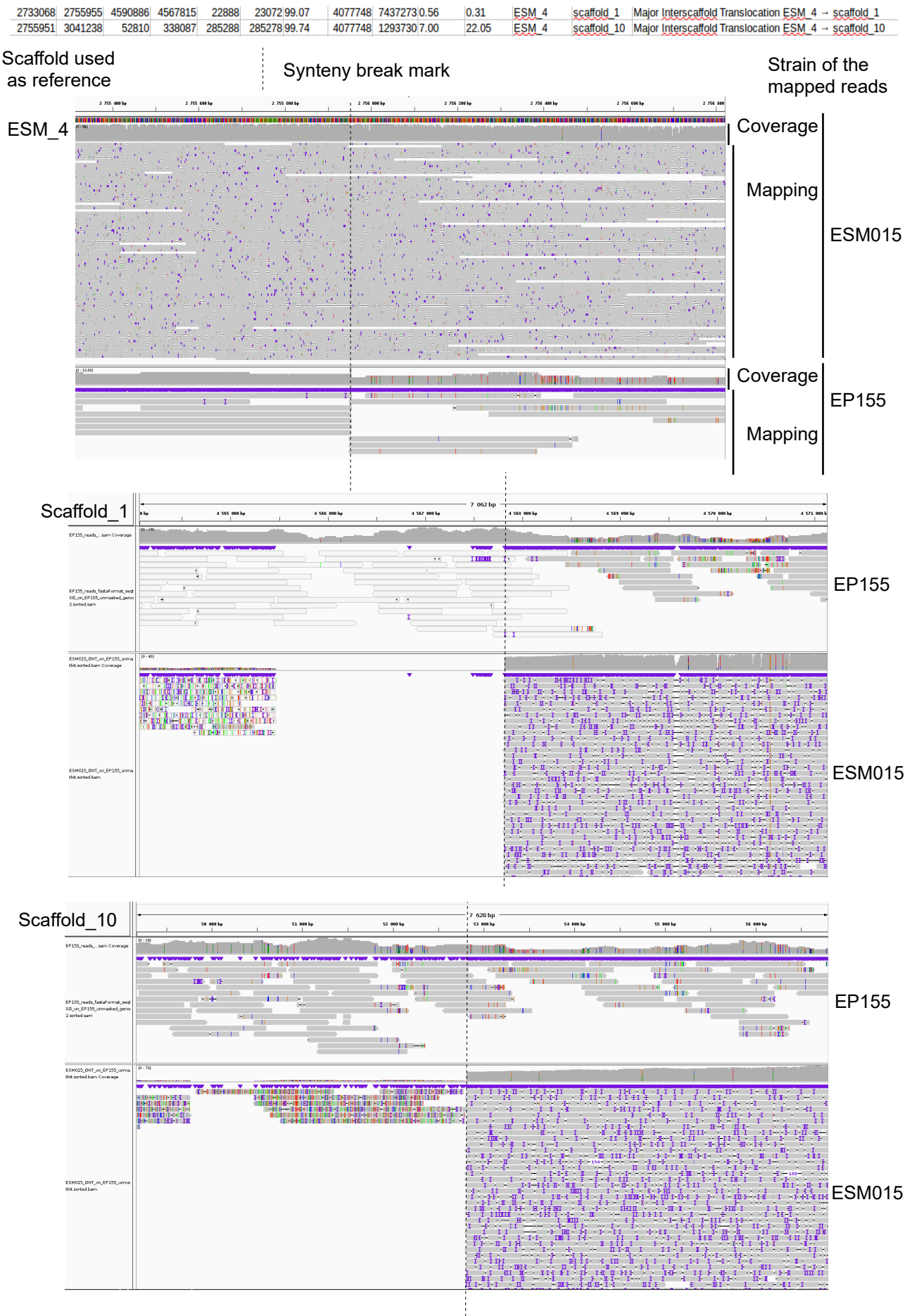


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2412868	2511629	884193	785452	98762	98742	99.86	5381768	3395974	1.84	2.91	ESM_2	scaffold_7	Major Interscaffold Translocation ESM_2 → scaffold_7
2511626	2522534	2543830	2554729	10909	10900	99.86	5381768	4337670	0.20	0.25	ESM_2	scaffold_5	Major Interscaffold Translocation ESM_2 → scaffold_5

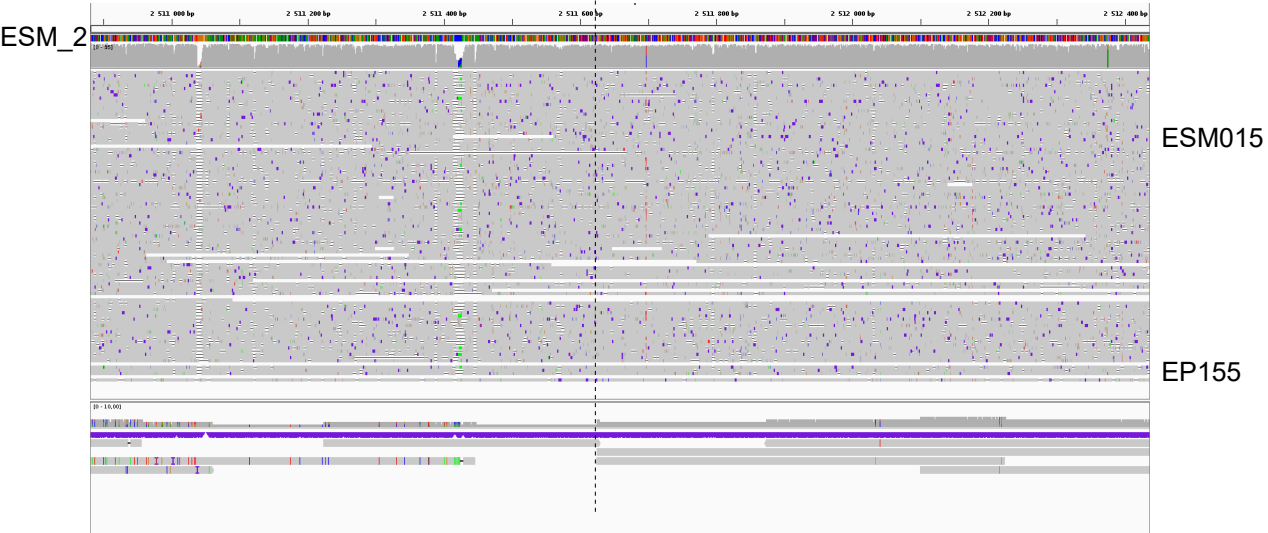


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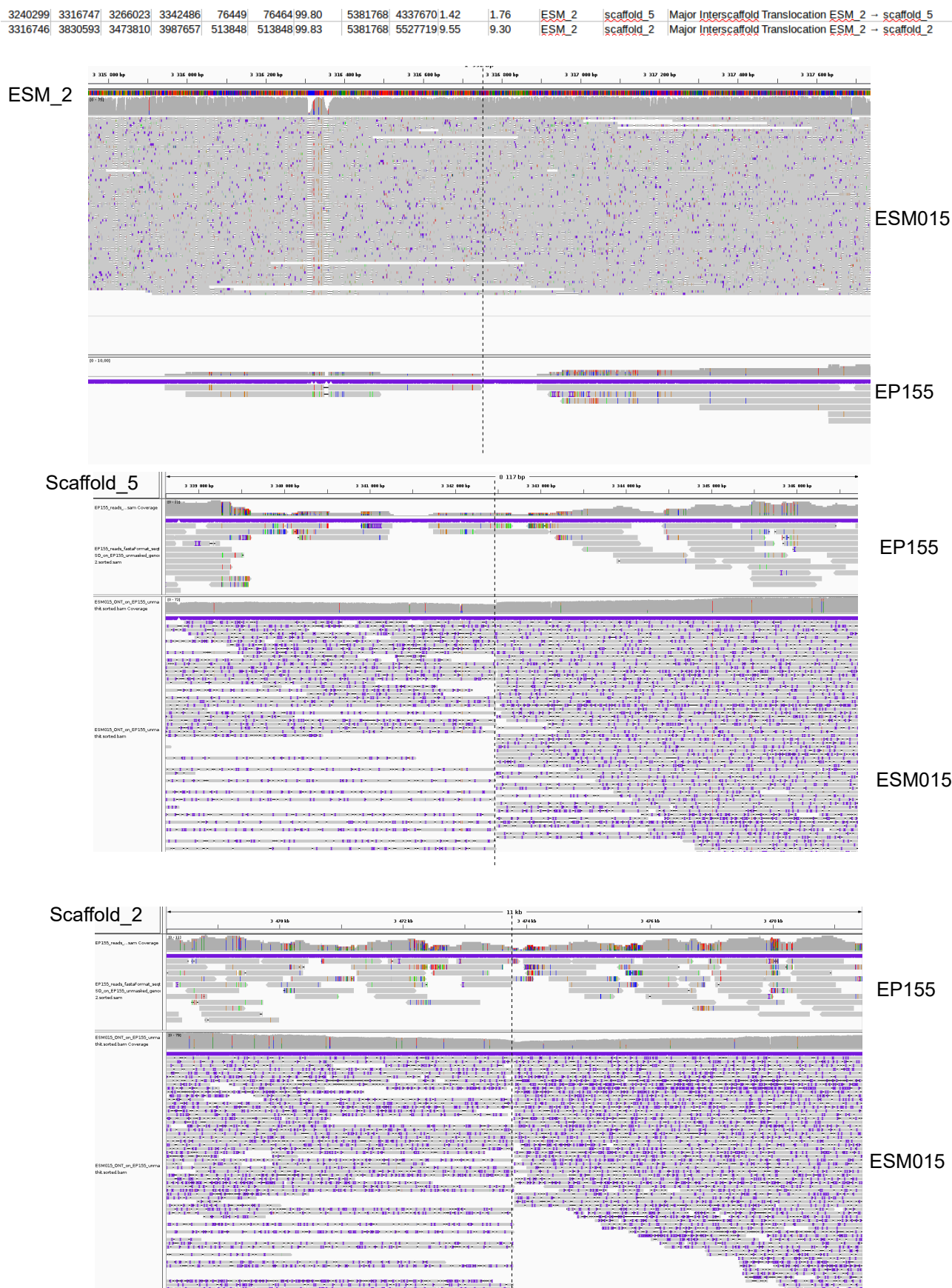


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1032112	1043922	3354296	3342486	11811	11811.99.99	2891089	4337670	0.41	0.27	ESM_7	scaffold_5	Major Interscaffold Translocation ESM_7 → scaffold_5
1043921	1118430	3473810	3399285	74510	74526.99.88	2891089	5527719	2.58	1.35	ESM_7	scaffold_2	Major Interscaffold Translocation ESM_7 → scaffold_2

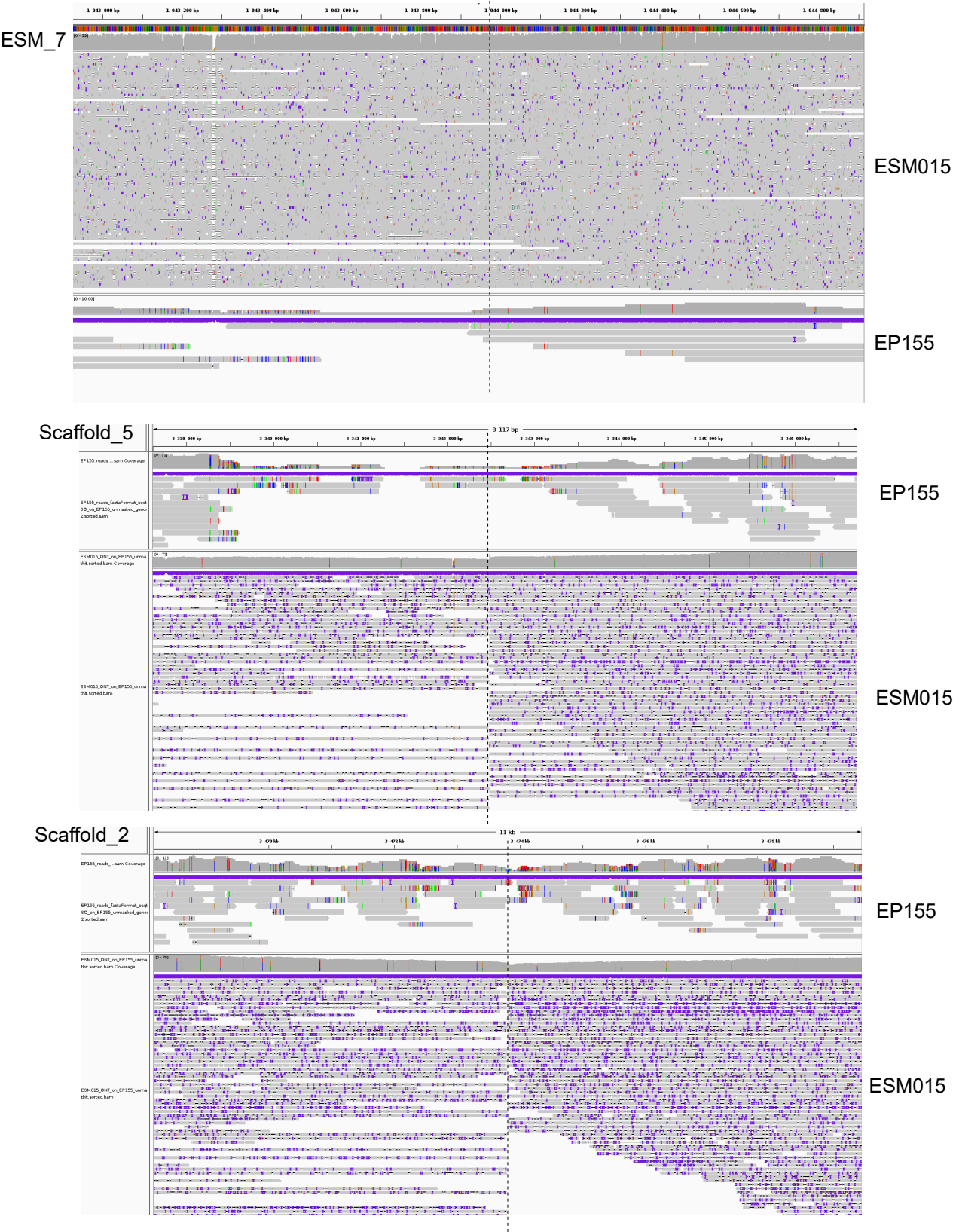




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1004321	1053110	736632	785455	48790	4882499.89	3530110	3395974	1.38	1.44	ESM_5	scaffold_7	Major Interscaffold Translocation ESM_5 → scaffold_7
1053107	1097962	2543833	2498955	44856	4487999.90	3530110	4337670	1.27	1.03	ESM_5	scaffold_5	Major Interscaffold Translocation ESM_5 → scaffold_5

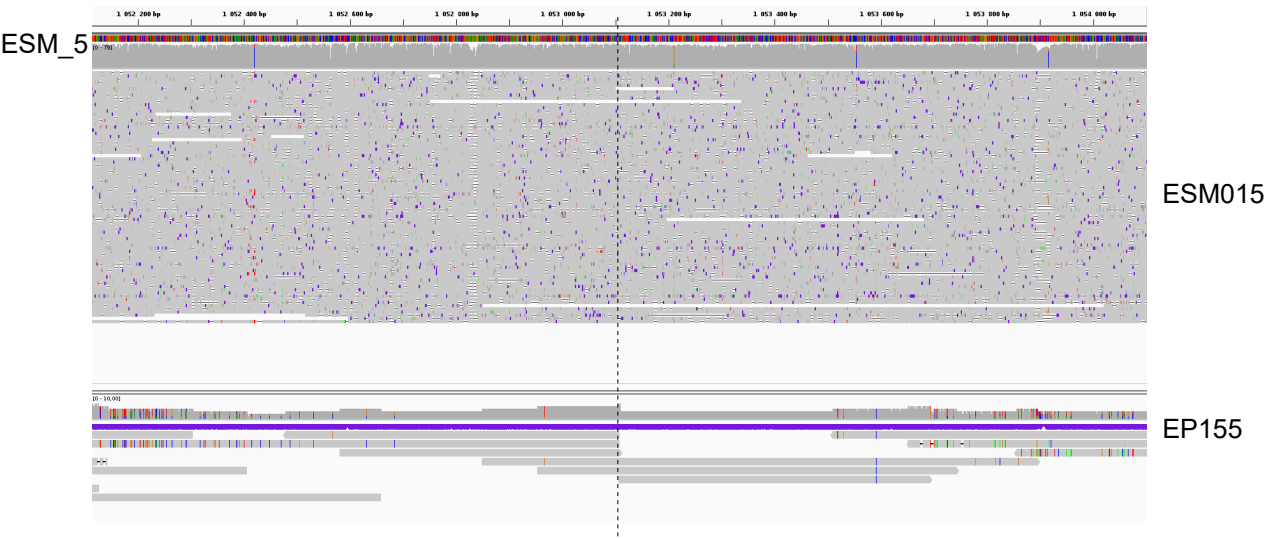


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1543121	1775927	2383767	2150857	232807	23291199.87	3437609	39507946.77	5.90	ESM_6	scaffold_6	Major Intrasc scaffold translocation ESM_6 → scaffold_6
1775920	1883961	3350	111362	108042	10801399.92	3437609	39507943.14	2.73	ESM_6	scaffold_6	Major Intrasc scaffold translocation ESM_6 → scaffold_6

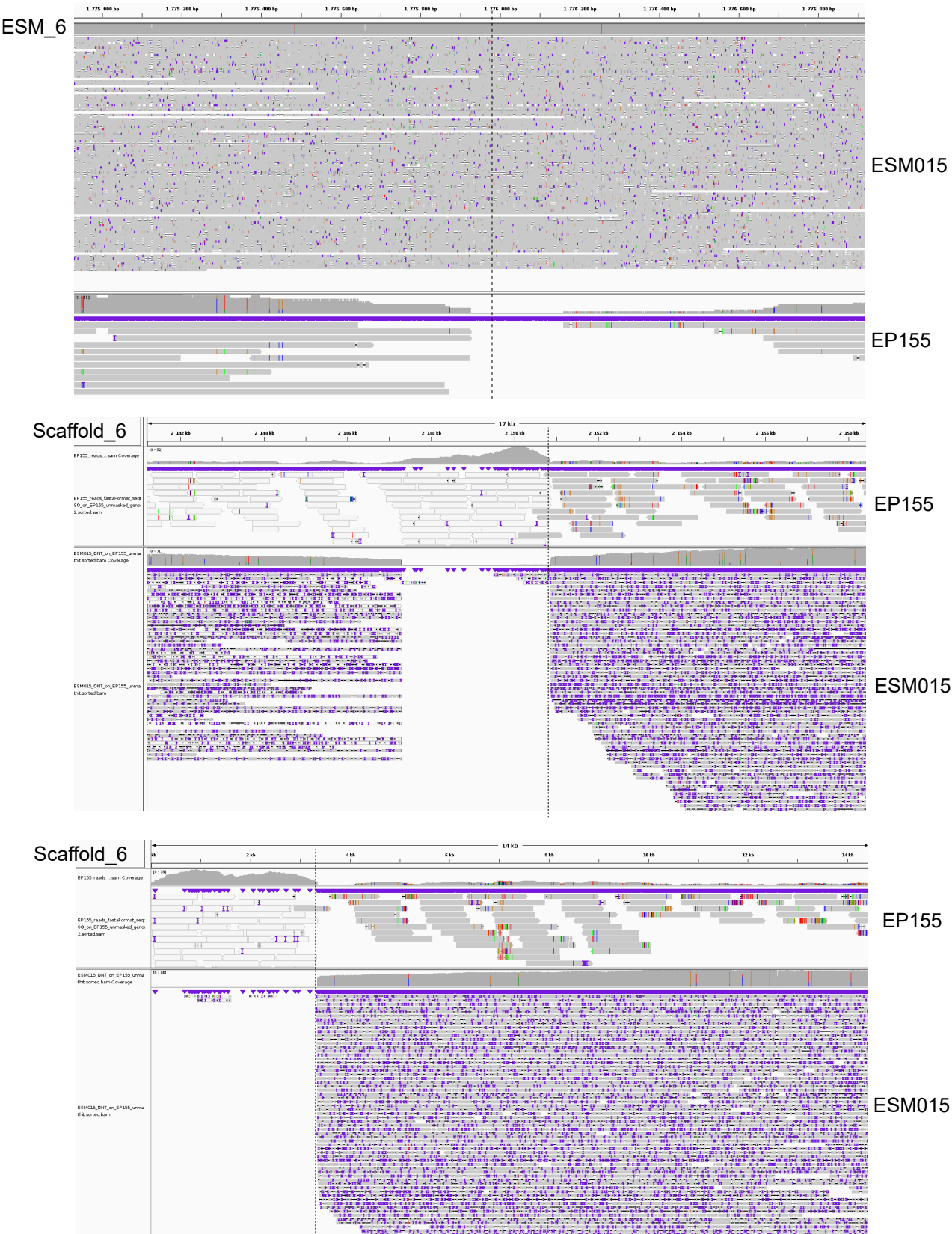


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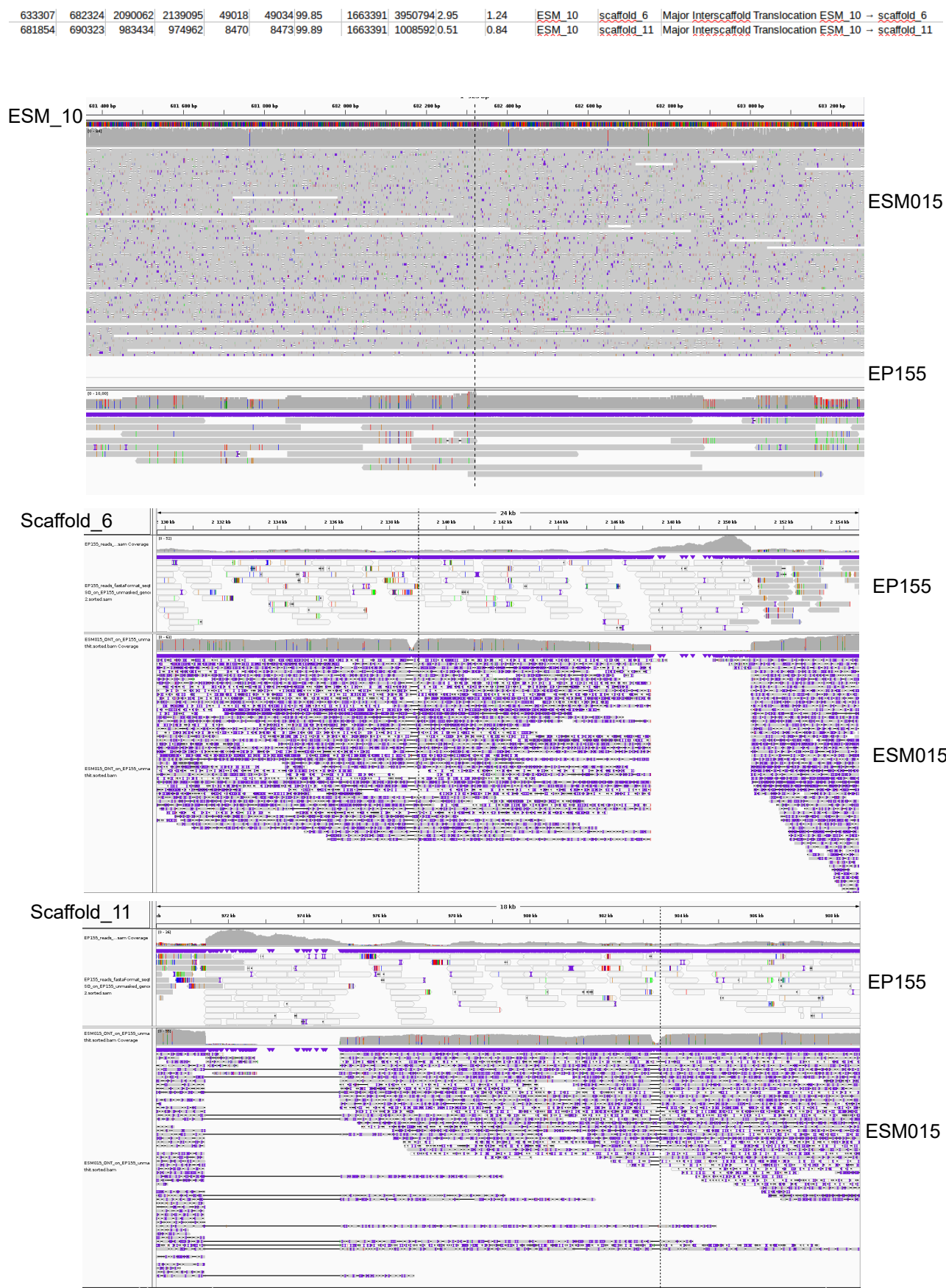




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75387	93013	1423790	1406164	17627	17627	97.48	599387	5527719	2.94	0.32	ESM_11	scaffold_2	Major Intrascaffold Translocation ESM_11 → scaffold_2
115383	179603	591138	655400	64221	64263	98.97	599387	5527719	10.71	1.16	ESM_11	scaffold_2	Major Intrascaffold Translocation ESM_11 → scaffold_2

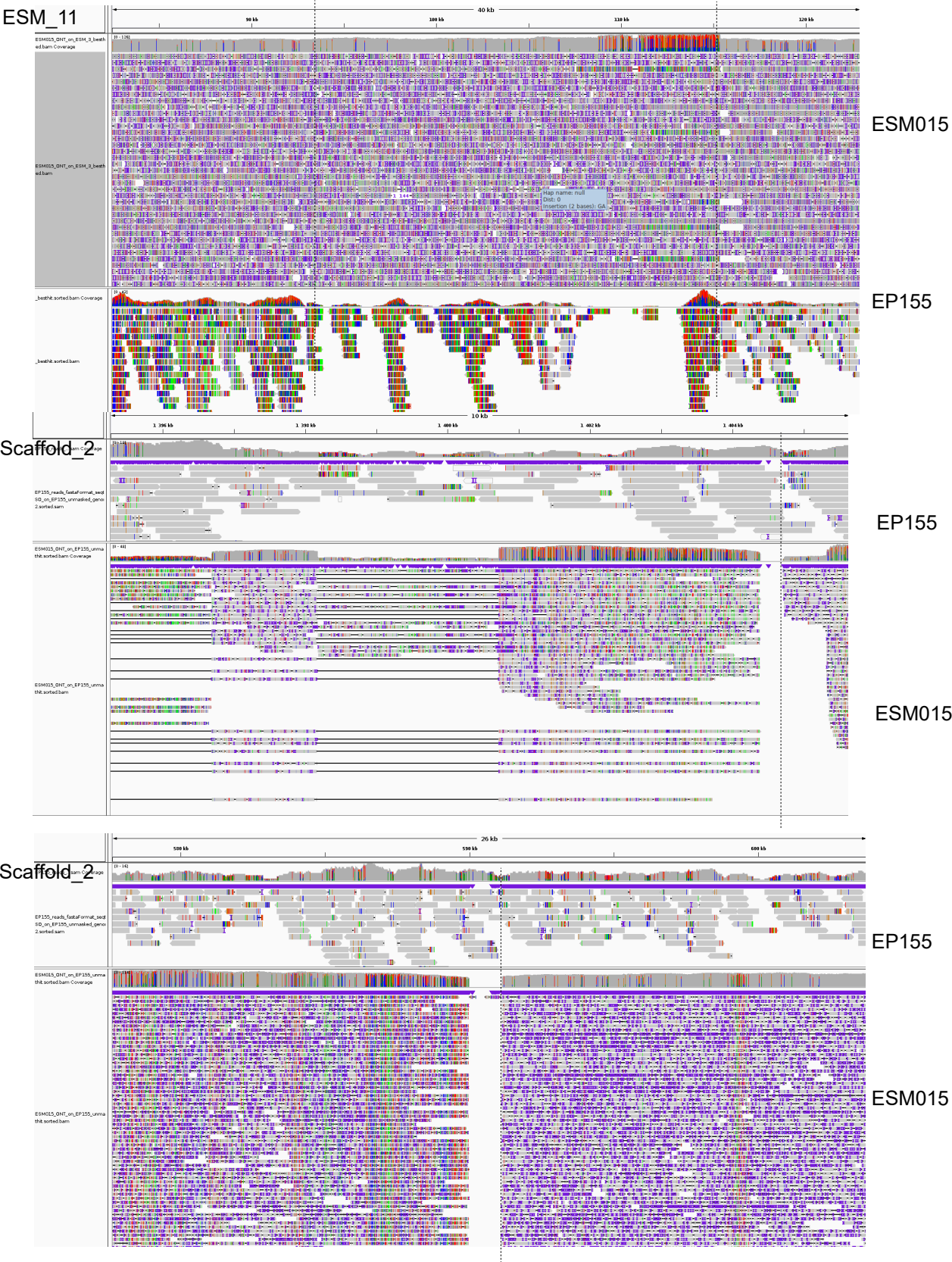




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187183	202667	574535	590007	15485	15473.97.07	500935	5527719.3.09	0.28	ESM_12	scaffold_2	Major Intrascaffold Translocation ESM_12 → scaffold_2
211902	234003	1377750	1355648	22102	22103.99.81	500935	5527719.4.41	0.40	ESM_12	scaffold_2	Major Intrascaffold Translocation ESM_12 → scaffold_2

