

Annotation Tracks for Genome Browser

Su(Hw)1: *Adh* region sites that match the consensus with a p value of -12 or better and that map to fragments with enrichment of at least 1.7 fold in ChIP from at least one of the chromatin sources. 60 sites

Su(Hw)2: *Adh* region sites that match the consensus with a p value between -12 and -15 and that map to fragments with enrichment of at least 1.7 fold in ChIP from at least one of the chromatin sources. 18 sites.

Su(Hw)3: *Adh* region sites that match the consensus with a p value of -15 or better and that map to fragments with enrichment of at least 1.7 fold in ChIP from at least one of the chromatin sources. 42 sites.

Su(Hw)4: *Adh* region sites that match the consensus with a p value of -15 or better and that map to fragments whose ChIP enrichment does not pass the 1.7 fold threshold. 37 sites.

----- cut below and paste for UCSC -----

browser position chr2L:13500000-16500000

track name=Su(Hw)1 description="SuHw Patser p value ≤ -12 and enrichment ≥ 1.7 fold" color=255,0,0

chr2L	13584993	13585012
chr2L	13607659	13607678
chr2L	13614204	13614223
chr2L	13629839	13629858
chr2L	13632623	13632642
chr2L	13638889	13638908
chr2L	13641670	13641689
chr2L	13667151	13667170
chr2L	13677997	13678016
chr2L	13714589	13714608
chr2L	13824516	13824535
chr2L	13894059	13894078
chr2L	13942758	13942777
chr2L	13969073	13969092
chr2L	13973053	13973072
chr2L	13988200	13988219
chr2L	14014963	14014982
chr2L	14020712	14020731
chr2L	14041051	14041070
chr2L	14082802	14082821
chr2L	14089608	14089627
chr2L	14134128	14134147
chr2L	14134237	14134256
chr2L	14166557	14166576
chr2L	14175641	14175660
chr2L	14179505	14179524

chr2L	14179577	14179596
chr2L	14234119	14234138
chr2L	14234544	14234563
chr2L	14555301	14555320
chr2L	14601019	14601038
chr2L	14688545	14688564
chr2L	14807926	14807945
chr2L	14864561	14864580
chr2L	14919464	14919483
chr2L	15052715	15052734
chr2L	15053529	15053548
chr2L	15075047	15075066
chr2L	15137627	15137646
chr2L	15159546	15159565
chr2L	15555313	15555332
chr2L	15591842	15591861
chr2L	15631187	15631206
chr2L	15634144	15634163
chr2L	15634525	15634544
chr2L	15660497	15660516
chr2L	15667741	15667760
chr2L	15668122	15668141
chr2L	15695494	15695513
chr2L	15717602	15717621
chr2L	15774643	15774662
chr2L	15811071	15811090
chr2L	15898198	15898217
chr2L	15910924	15910943
chr2L	15946636	15946655
chr2L	16116870	16116889
chr2L	16349908	16349927
chr2L	16352558	16352577
chr2L	16355644	16355663
chr2L	16402841	16402860

track name=Su(Hw)2 description=" SuHw -15< Patser p value <= -12 and enrichment >= 1.7 fold " color=0,0,255

chr2L	13973053	13973072
chr2L	14134237	14134256
chr2L	14175641	14175660
chr2L	14179505	14179524
chr2L	14179577	14179596
chr2L	14601019	14601038
chr2L	15052715	15052734
chr2L	15591842	15591861
chr2L	15631187	15631206
chr2L	15634144	15634163
chr2L	15634525	15634544
chr2L	15667741	15667760
chr2L	15668122	15668141
chr2L	15774643	15774662

chr2L	15811071	15811090
chr2L	15946636	15946655
chr2L	16349908	16349927
chr2L	16402841	16402860

track name=Su(Hw)3 description=" SuHw Patser p value <= -15 and enrichment >= 1.7 fold " color=255,0,0

chr2L	13584993	13585012
chr2L	13607659	13607678
chr2L	13614204	13614223
chr2L	13629839	13629858
chr2L	13632623	13632642
chr2L	13638889	13638908
chr2L	13641670	13641689
chr2L	13667151	13667170
chr2L	13677997	13678016
chr2L	13714589	13714608
chr2L	13824516	13824535
chr2L	13894059	13894078
chr2L	13942758	13942777
chr2L	13969073	13969092
chr2L	13988200	13988219
chr2L	14014963	14014982
chr2L	14020712	14020731
chr2L	14041051	14041070
chr2L	14082802	14082821
chr2L	14089608	14089627
chr2L	14134128	14134147
chr2L	14166557	14166576
chr2L	14234119	14234138
chr2L	14234544	14234563
chr2L	14555301	14555320
chr2L	14688545	14688564
chr2L	14807926	14807945
chr2L	14864561	14864580
chr2L	14919464	14919483
chr2L	15053529	15053548
chr2L	15075047	15075066
chr2L	15137627	15137646
chr2L	15159546	15159565
chr2L	15555313	15555332
chr2L	15660497	15660516
chr2L	15695494	15695513
chr2L	15717602	15717621
chr2L	15898198	15898217
chr2L	15910924	15910943
chr2L	16116870	16116889
chr2L	16352558	16352577
chr2L	16355644	16355663

track name=Su(Hw)4 description=" SuHw Patser p value < -15 and enrichment < 1.7 fold " color=128,128,128

chr2L	13576244	13576263
chr2L	13632623	13632642
chr2L	13641670	13641689
chr2L	13749739	13749758
chr2L	13969073	13969092
chr2L	14057173	14057192
chr2L	14106974	14106993
chr2L	14110653	14110672
chr2L	14352281	14352300
chr2L	14688545	14688564
chr2L	14788814	14788833
chr2L	14882942	14882961
chr2L	14939553	14939572
chr2L	14991308	14991327
chr2L	14998235	14998254
chr2L	15056641	15056660
chr2L	15137627	15137646
chr2L	15159546	15159565
chr2L	15212509	15212528
chr2L	15213622	15213641
chr2L	15281950	15281969
chr2L	15370457	15370476
chr2L	15543249	15543268
chr2L	15615023	15615042
chr2L	15658912	15658931
chr2L	15693909	15693928
chr2L	15836263	15836282
chr2L	15872750	15872769
chr2L	15932483	15932502
chr2L	15952587	15952606
chr2L	15961036	15961055
chr2L	16013343	16013362
chr2L	16042590	16042609
chr2L	16056445	16056464
chr2L	16129490	16129509
chr2L	16327530	16327549
chr2L	16365484	16365503