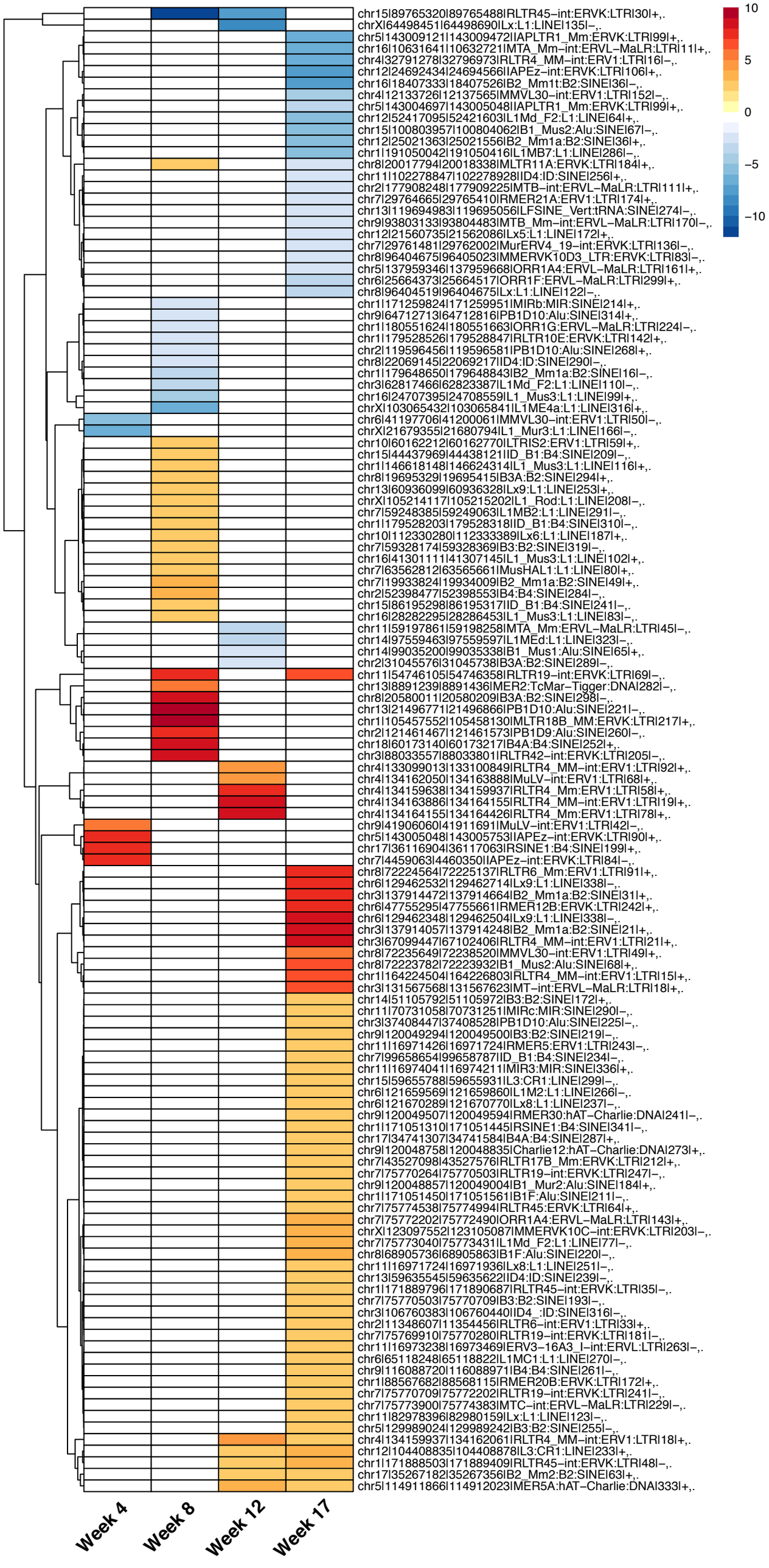
10
5
0
−5
−10
chr15|89765320|89765488|RLTR45−int:ERVK:LTR|30|+,.
chrX|64498451|64498690|Lx:L1:LINE|135|−,.
chr5|143009121|143009472|IAPLTR1_Mm:ERVK:LTR|99|+,.
chr16|10631641|10632721|MTA_Mm−int:ERVL−MaLR:LTR|11|+,.
chr4|32791278|32796973|RLTR4_MM−int:ERV1:LTR|16|−,.
chr12|24692434|24694566|IAPEz−int:ERVK:LTR|106|+,.
chr16|18407333|18407526|B2_Mm1t:B2:SINE|36|−,.
chr4|12133726|12137565|MMVL30−int:ERV1:LTR|152|−,.
chr5|143004697|143005048|IAPLTR1_Mm:ERVK:LTR|99|+,.
chr12|52417095|52421603|L1Md_F2:L1:LINE|64|+,.
chr15|100803957|100804062|B1_Mus2:Alu:SINE|67|−,.
chr12|25021363|25021556|B2_Mm1a:B2:SINE|36|+,.
chr1|191050042|191050416|L1MB7:L1:LINE|286|−,.
chr8|20017794|20018338|MLTR11A:ERVK:LTR|184|+,.
chr11|102278847|102278928|ID4:ID:SINE|256|+,.
chr2|177908248|177909225|MTB−int:ERVL−MaLR:LTR|111|+,.
chr7|29764665|29765410|RMER21A:ERV1:LTR|174|+,.
chr13|119694983|119695056|LFSINE_Vert:tRNA:SINE|274|−,.
chr9|93803133|93804483|MTB_Mm−int:ERVL−MaLR:LTR|170|−,.
chr12|21560735|21562086|Lx5:L1:LINE|172|+,.
chr7|29761481|29762002|MurERV4_19−int:ERVK:LTR|136|−,.
chr8|96404675|96405023|MMERVK10D3_LTR:ERVK:LTR|83|−,.
chr5|137959346|137959668|ORR1A4:ERVL−MaLR:LTR|161|+,.
chr6|25664373|25664517|ORR1F:ERVL−MaLR:LTR|299|+,.
chr8|96404519|96404675|Lx:L1:LINE|122|−,.
chr1|171259824|171259951|MIRb:MIR:SINE|214|+,.
chr9|64712713|64712816|PB1D10:Alu:SINE|314|+,.
chr1|180551624|180551663|ORR1G:ERVL−MaLR:LTR|224|−,.
chr1|179528526|179528847|RLTR10E:ERVK:LTR|142|+,.
chr2|119596456|119596581|PB1D10:Alu:SINE|268|+,.
chr8|22069145|22069217|ID4:ID:SINE|290|−,.
chr1|179648650|179648843|B2_Mm1a:B2:SINE|16|−,.
chr3|62817466|62823387|L1Md_F2:L1:LINE|110|−,.
chr16|24707395|24708559|L1_Mus3:L1:LINE|99|+,.
chrX|103065432|103065841|L1ME4a:L1:LINE|316|+,.
chr6|41197706|41200061|MMVL30−int:ERV1:LTR|50|−,.
chrX|21679355|21680794|L1_Mur3:L1:LINE|166|−,.
chr10|60162212|60162770|LTRIS2:ERV1:LTR|59|+,.
chr15|44437969|44438121|ID_B1:B4:SINE|209|−,.
chr1|146618148|146624314|L1_Mus3:L1:LINE|116|+,.
chr8|19695329|19695415|B3A:B2:SINE|294|+,.
chr13|60936099|60936328|Lx9:L1:LINE|253|+,.
chrX|105214117|105215202|L1_Rod:L1:LINE|208|−,.
chr7|59248385|59249063|L1MB2:L1:LINE|291|−,.
chr1|179528203|179528318|ID_B1:B4:SINE|310|−,.
chr10|112330280|112333389|Lx6:L1:LINE|187|+,.
chr7|59328174|59328369|B3:B2:SINE|319|−,.
chr16|41301111|41307145|L1_Mus3:L1:LINE|102|+,.
chr7|63562812|63565661|MusHAL1:L1:LINE|80|+,.
chr7|19933824|19934009|B2_Mm1a:B2:SINE|49|+,.
chr2|52398477|52398553|B4:B4:SINE|284|−,.
chr15|86195298|86195317|ID_B1:B4:SINE|241|−,.
chr16|28282295|28286453|L1_Mus3:L1:LINE|83|−,.
chr11|59197861|59198258|MTA_Mm:ERVL−MaLR:LTR|45|−,.
chr14|97559463|97559597|L1MEd:L1:LINE|323|−,.
chr14|99035200|99035338|B1_Mus1:Alu:SINE|65|+,.
chr2|31045576|31045738|B3A:B2:SINE|289|−,.
chr11|54746105|54746358|RLTR19−int:ERVK:LTR|69|−,.
chr13|8891239|8891436|MER2:TcMar−Tigger:DNA|282|−,.
chr8|20580011|20580209|B3A:B2:SINE|298|−,.
chr13|21496771|21496866|PB1D10:Alu:SINE|221|−,.
chr1|105457552|105458130|MLTR18B_MM:ERVK:LTR|217|+,.
chr2|121461467|121461573|PB1D9:Alu:SINE|260|−,.
chr18|60173140|60173217|B4A:B4:SINE|252|+,.
chr3|88033557|88033801|RLTR42−int:ERVK:LTR|205|−,.
chr4|133099013|133100849|RLTR4_MM−int:ERV1:LTR|92|+,.
chr4|134162050|134163888|MuLV−int:ERV1:LTR|68|+,.
chr4|134159638|134159937|RLTR4_Mm:ERV1:LTR|58|+,.
chr4|134163886|134164155|RLTR4_MM−int:ERV1:LTR|19|+,.
chr4|134164155|134164426|RLTR4_Mm:ERV1:LTR|78|+,.
chr9|41906060|41911691|MuLV−int:ERV1:LTR|42|−,.
chr5|143005048|143005753|IAPEz−int:ERVK:LTR|90|+,.
chr17|36116904|36117063|RSINE1:B4:SINE|199|+,.
chr7|4459063|4460350|IAPEz−int:ERVK:LTR|84|−,.
chr8|72224564|72225137|RLTR6_Mm:ERV1:LTR|91|+,.
chr6|129462532|129462714|Lx9:L1:LINE|338|−,.
chr3|137914472|137914664|B2_Mm1a:B2:SINE|31|+,.
chr6|47755295|47755661|RMER12B:ERVK:LTR|242|+,.
chr6|129462348|129462504|Lx9:L1:LINE|338|−,.
chr3|137914057|137914248|B2_Mm1a:B2:SINE|21|+,.
chr3|67099447|67102406|RLTR4_MM−int:ERV1:LTR|21|+,.
chr8|72235649|72238520|MMVL30−int:ERV1:LTR|49|+,.
chr8|72223782|72223932|B1_Mus2:Alu:SINE|68|+,.
chr1|164224504|164226803|RLTR4_MM−int:ERV1:LTR|15|+,.
chr3|131567568|131567623|MT−int:ERVL−MaLR:LTR|18|+,.
chr14|51105792|51105972|B3:B2:SINE|172|+,.
chr11|70731058|70731251|MIRc:MIR:SINE|290|−,.
chr3|37408447|37408528|PB1D10:Alu:SINE|225|−,.
chr9|120049294|120049500|B3:B2:SINE|219|−,.
chr11|16971426|16971724|RMER5:ERV1:LTR|243|−,.
chr7|99658654|99658787|ID_B1:B4:SINE|234|−,.
chr11|16974041|16974211|MIR3:MIR:SINE|336|+,.
chr15|59655788|59655931|L3:CR1:LINE|299|−,.
chr6|121659569|121659860|L1M2:L1:LINE|266|−,.
chr6|121670289|121670770|Lx8:L1:LINE|237|−,.
chr9|120049507|120049594|RMER30:hAT−Charlie:DNA|241|−,.
chr1|171051310|171051445|RSINE1:B4:SINE|341|−,.
chr17|34741307|34741584|B4A:B4:SINE|287|+,.
chr9|120048758|120048835|Charlie12:hAT−Charlie:DNA|273|+,.
chr7|43527098|43527576|RLTR17B_Mm:ERVK:LTR|212|+,.
chr7|75770264|75770503|RLTR19−int:ERVK:LTR|247|−,.
chr9|120048857|120049004|B1_Mur2:Alu:SINE|184|+,.
chr1|171051450|171051561|B1F:Alu:SINE|211|−,.
chr7|75774538|75774994|RLTR45:ERVK:LTR|64|+,.
chr7|75772202|75772490|ORR1A4:ERVL−MaLR:LTR|143|+,.
chrX|123097552|123105087|MMERVK10C−int:ERVK:LTR|203|−,.
chr7|75773040|75773431|L1Md_F2:L1:LINE|77|−,.
chr8|68905736|68905863|B1F:Alu:SINE|220|−,.
chr11|16971724|16971936|Lx8:L1:LINE|251|−,.
chr13|59635545|59635622|ID4:ID:SINE|239|−,.
chr1|171889796|171890687|RLTR45−int:ERVK:LTR|35|−,.
chr7|75770503|75770709|B3:B2:SINE|193|−,.
chr3|106760383|106760440|ID4_:ID:SINE|316|−,.
chr2|11348607|11354456|RLTR6−int:ERV1:LTR|33|+,.
chr7|75769910|75770280|RLTR19−int:ERVK:LTR|181|−,.
chr11|16973238|16973469|ERV3−16A3_I−int:ERVL:LTR|263|−,.
chr6|65118248|65118822|L1MC1:L1:LINE|270|−,.
chr9|116088720|116088971|B4:B4:SINE|261|−,.
chr1|88567682|88568115|RMER20B:ERVK:LTR|172|+,.
chr7|75770709|75772202|RLTR19−int:ERVK:LTR|241|−,.
chr7|75773900|75774383|MTC−int:ERVL−MaLR:LTR|229|−,.
chr11|82978396|82980159|Lx:L1:LINE|123|−,.
chr5|129989024|129989242|B3:B2:SINE|255|−,.
chr4|134159937|134162061|RLTR4_MM−int:ERV1:LTR|18|+,.
chr12|104408835|104408878|L3:CR1:LINE|233|+,.
chr1|171888503|171889409|RLTR45−int:ERVK:LTR|48|−,.
chr17|35267182|35267356|B2_Mm2:B2:SINE|63|+,.
chr5|114911866|114912023|MER5A:hAT−Charlie:DNA|333|+,.
Week 4
Week 8
Week 12
Week 17