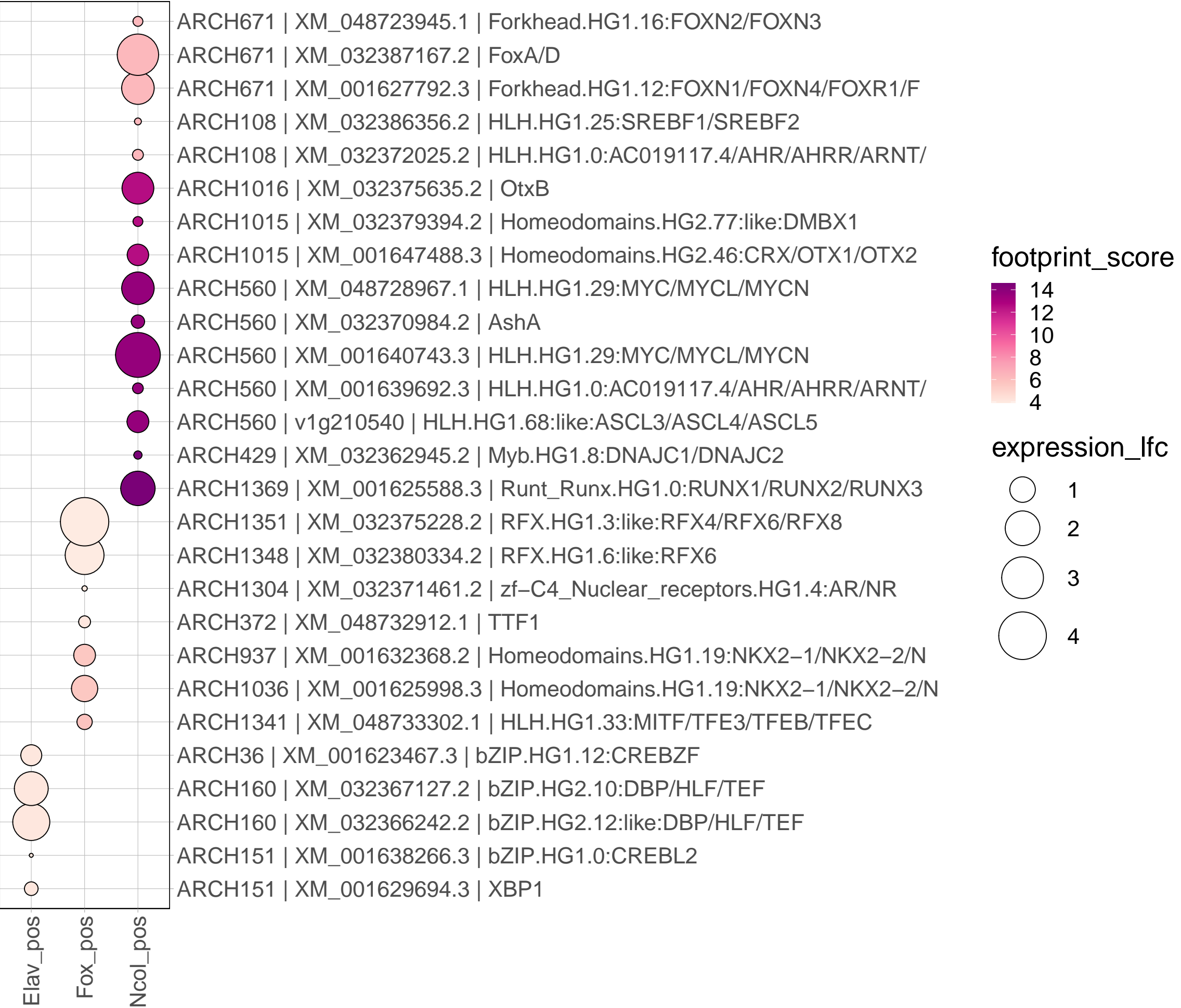
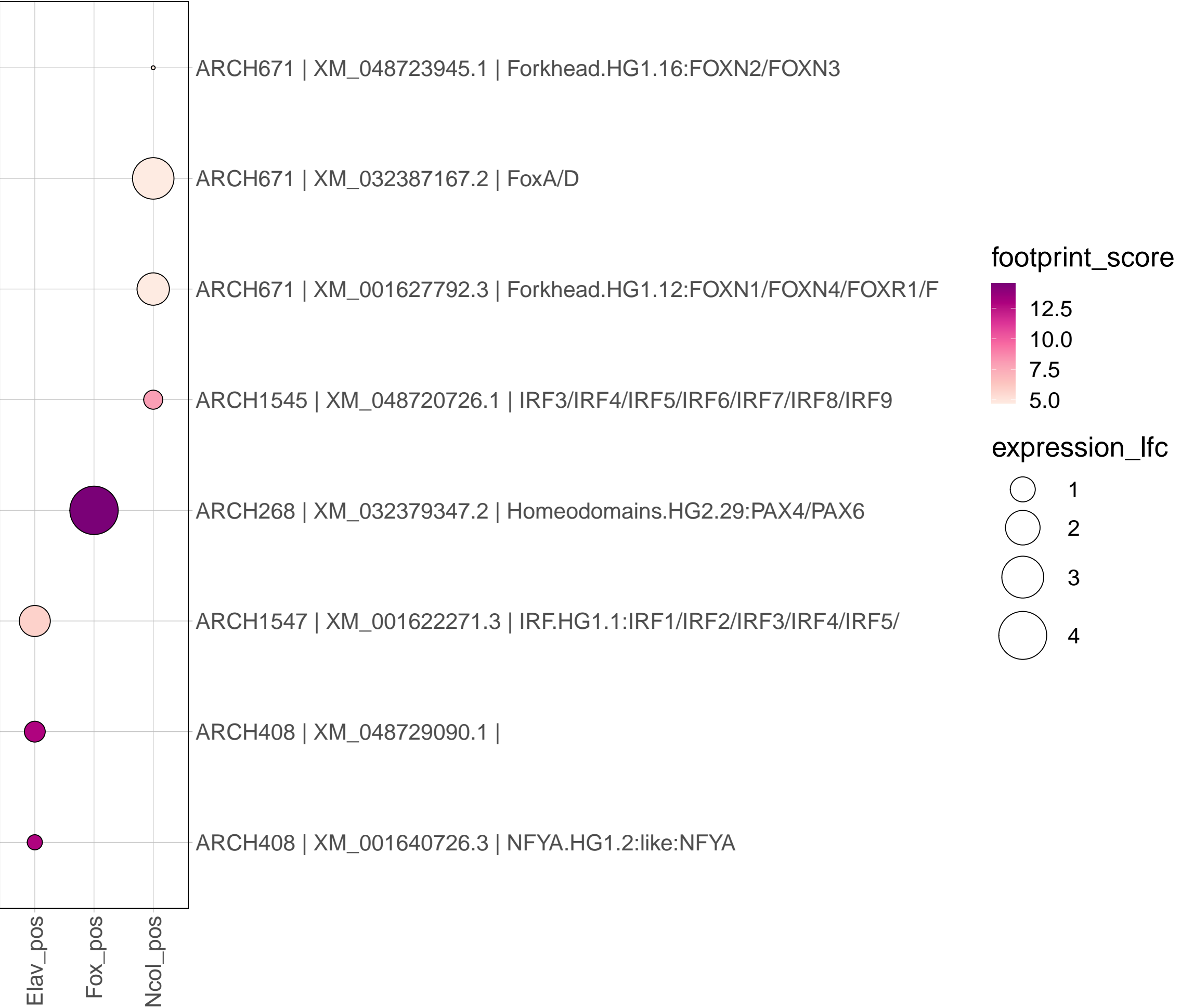


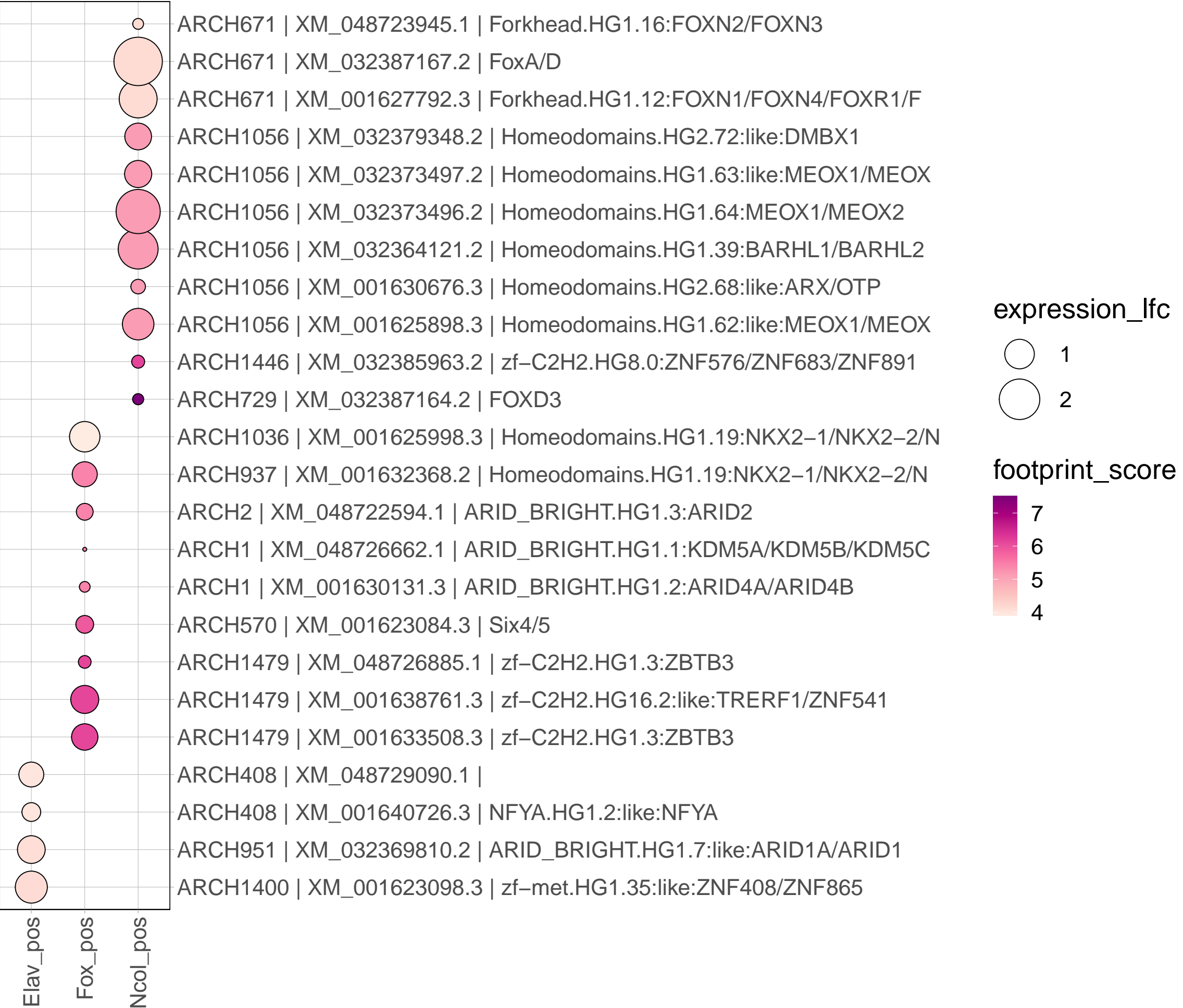
Nvec_vc1.1_XM_032385685.2
GPCR–rhodopsin:like:GPR176:likeclu:; 7tm_1/



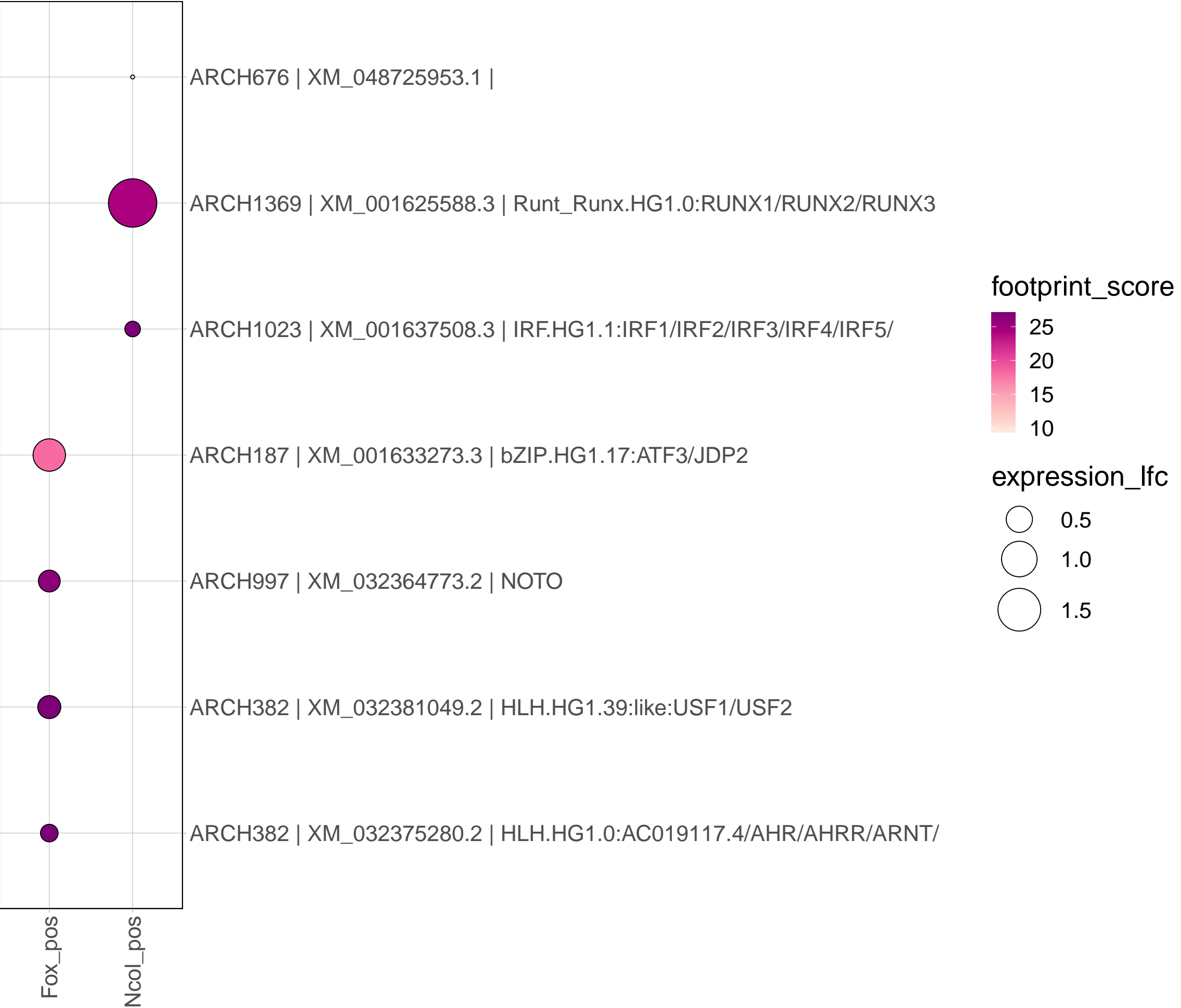
Nvec_vc1.1_XM_032367127.2
bZIP.HG2.10:DBP/HLF/TEF; DBP/HLF/TEF



Nvec_vc1.1_XM_048720228.1
Sushi, nidogen and EGF-like domain-; NIDO/

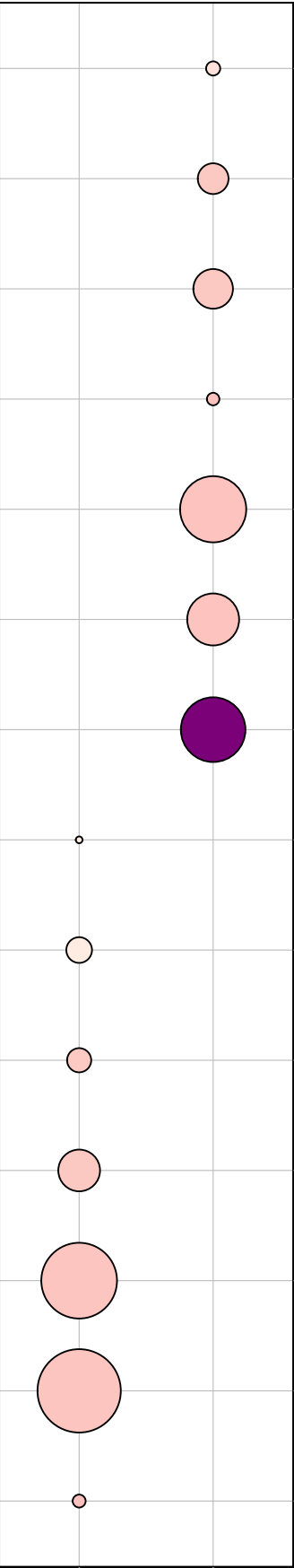


Nvec_vc1.1_XM_048734432.1
Folliculin–interacting protein 1; FNIP_N/FNIP_M/FNIP_C/



Nvec_vc1.1_XM_001627750.3

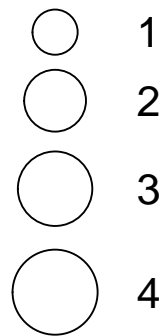
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Fox_pos

Ncol_pos

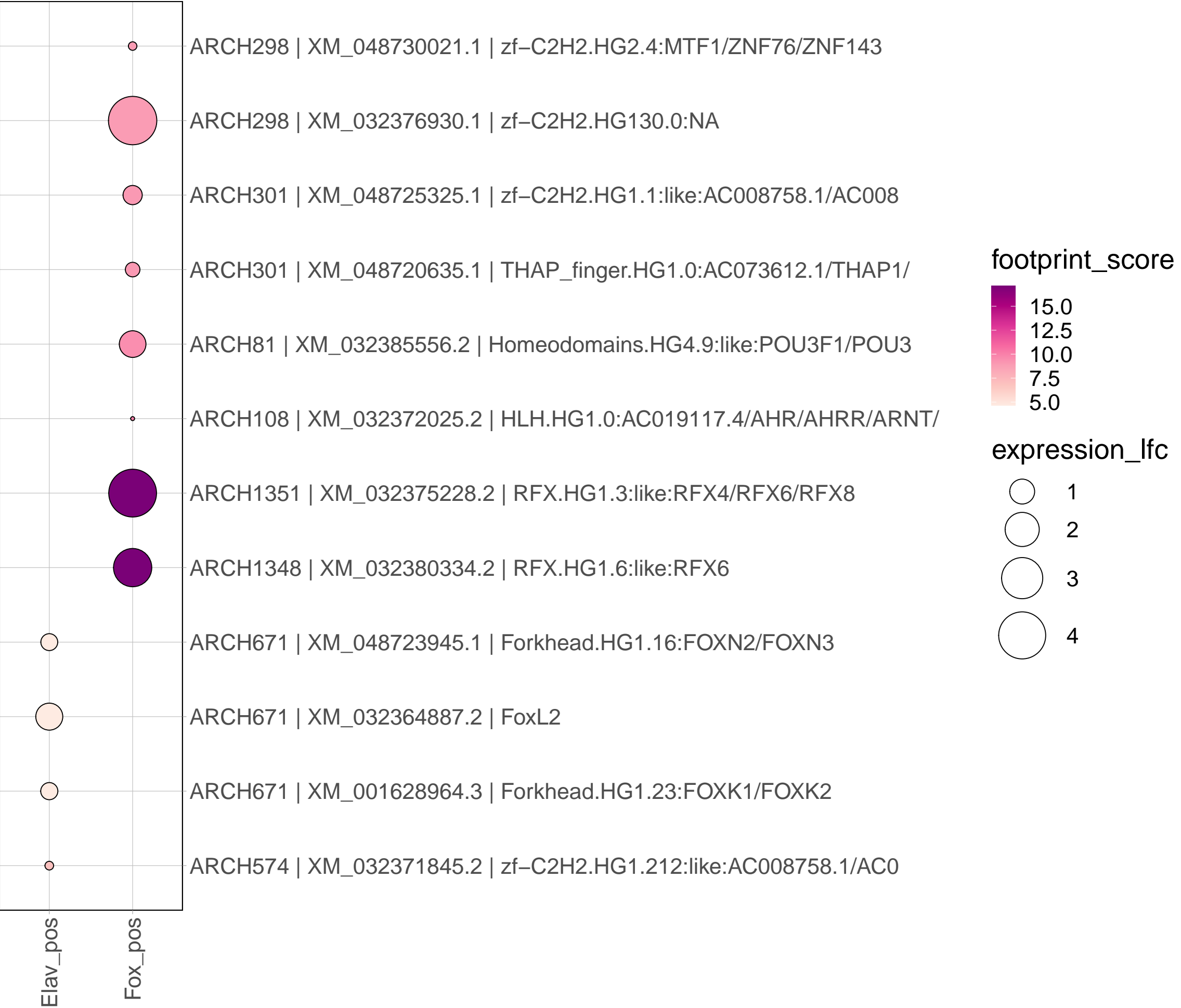
expression_lfc



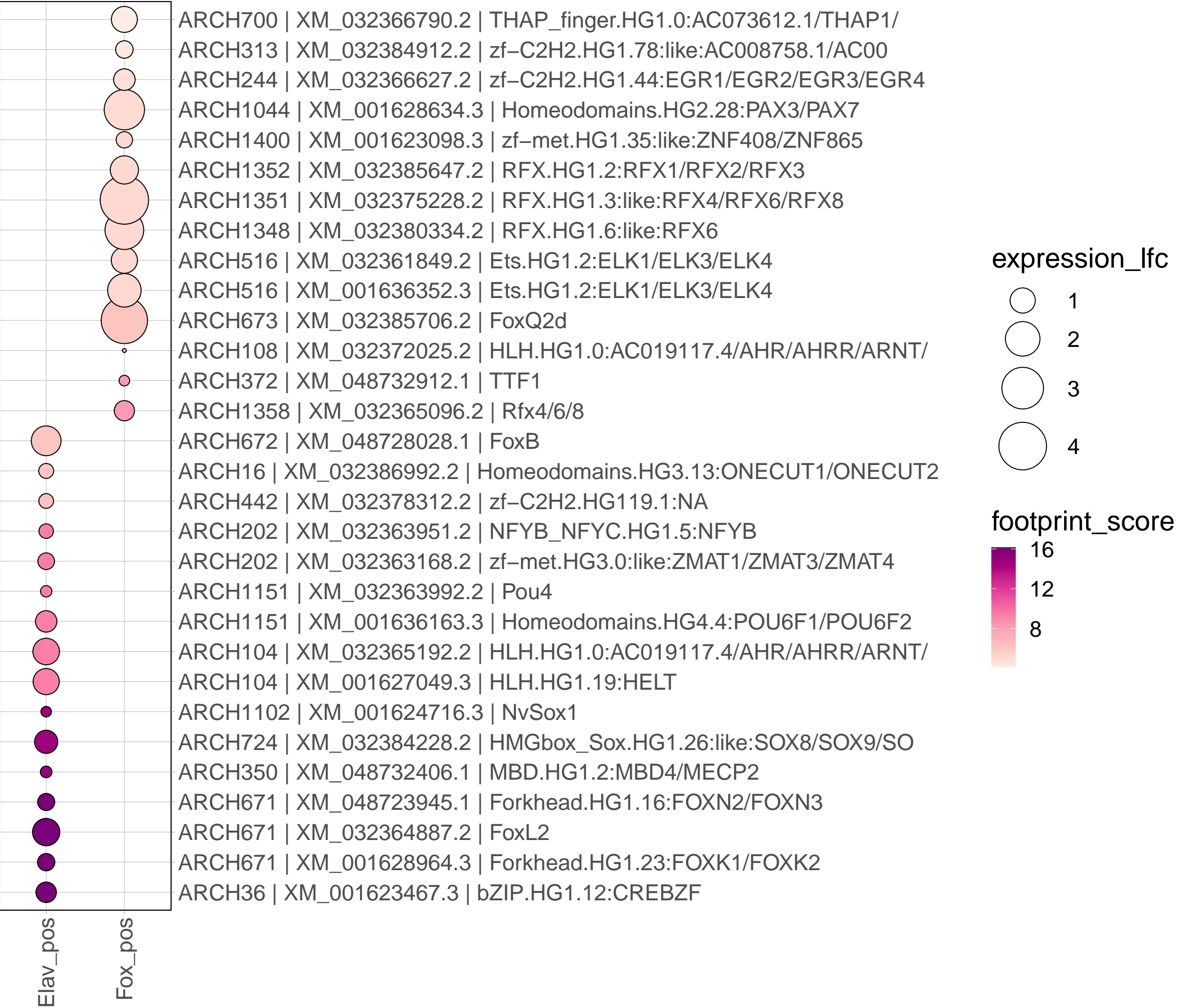
footprint_score



Nvec_vc1.1_XM_048722463.1
GPCR–rhodopsin:NA; 7tm_1/

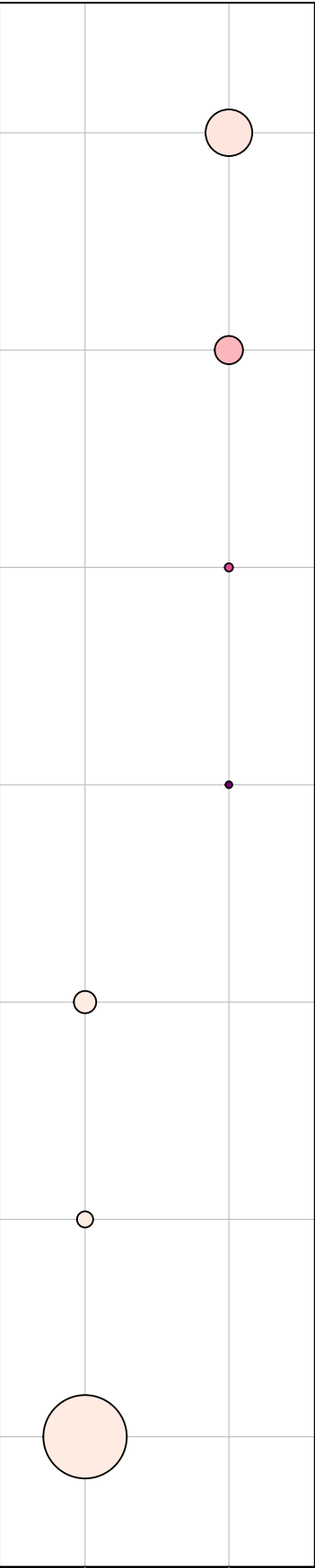


Nvec_vc1.1_XM_001627964.3
ion-NaKCa:KCNA1/KCNA10/KCNA2/KCNA3/; ion-NaKCa.HG1.6:KCNA1/KCNA



Nvec_vc1.1_XM_048729250.1

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Fox_pos

Ncol_pos

expression_lfc

2

4

6

footprint_score

15.0

12.5

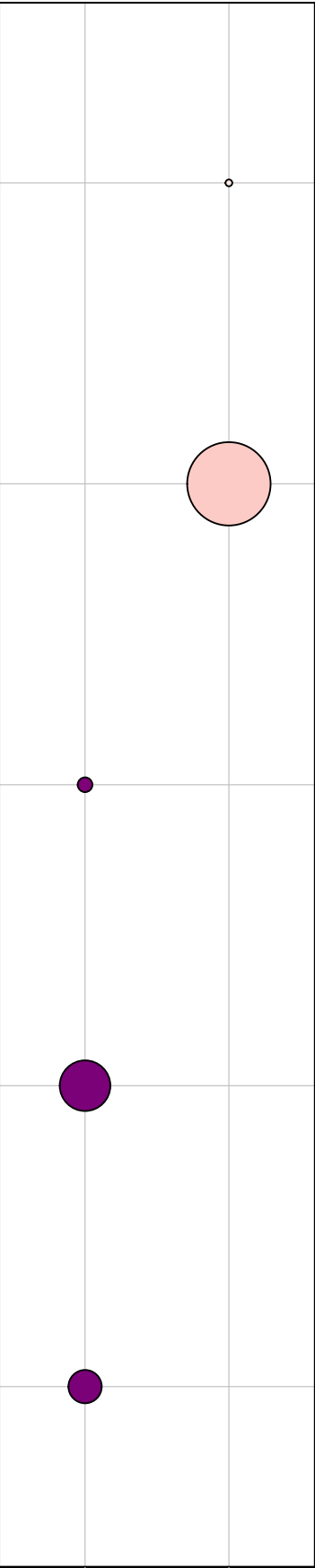
10.0

7.5

5.0

Nvec_vc1.1_XM_001641405.3

;
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Fox_pos

Ncol_pos

ARCH518 | XM_032385089.2 | Ets.HG1.7:GABPA

ARCH1425 | XM_001639701.3 | NvSox2

ARCH1104 | XM_032364225.2 | HMGbox_Sox.HG1.64:like:HMG20B

ARCH382 | XM_032381049.2 | HLH.HG1.39:like:USF1/USF2

ARCH382 | XM_032375280.2 | HLH.HG1.0:AC019117.4/AHR/AHRR/ARNT/

expression_lfc

0.25

0.50

0.75

footprint_score

14

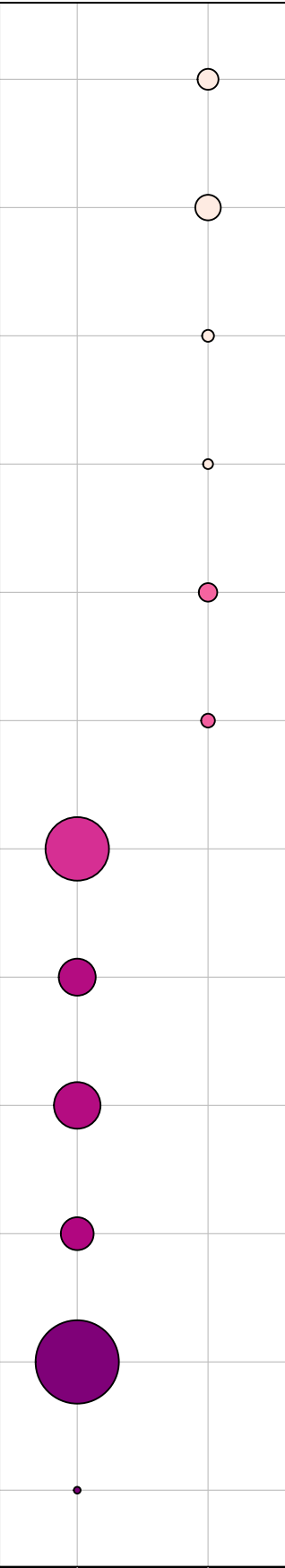
12

10

8

Nvec_vc1.1_XM_032375896.2

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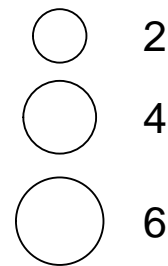
Fox_pos

Ncol_pos

footprint_score



expression_lfc



Nvec_vc1.1_XM_032376323.2

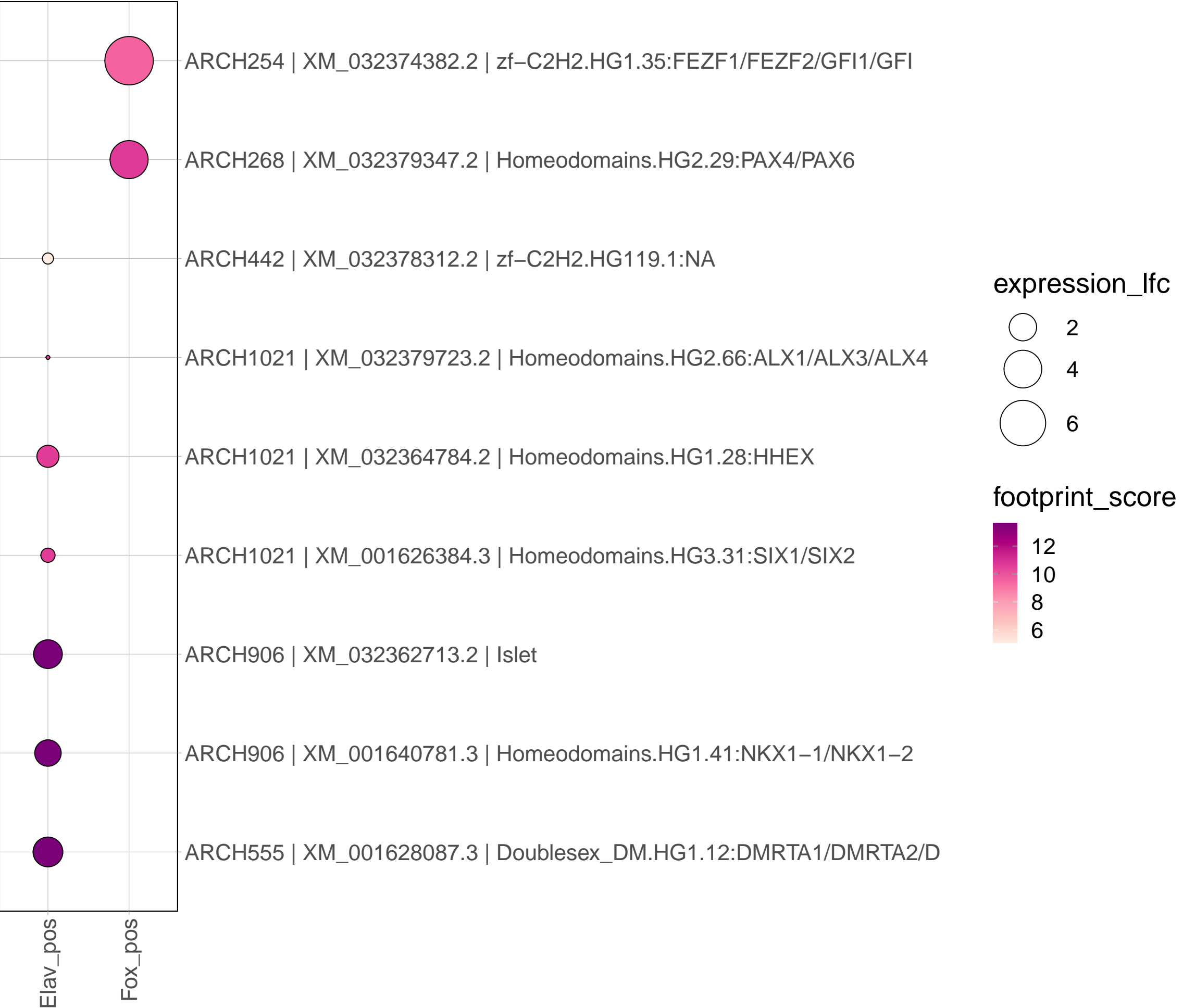
Pkinase:like:GUCY2C/GUCY2D/GUCY2F/N; Pkinase_Tyr/HNOBA/Guanylate_cy



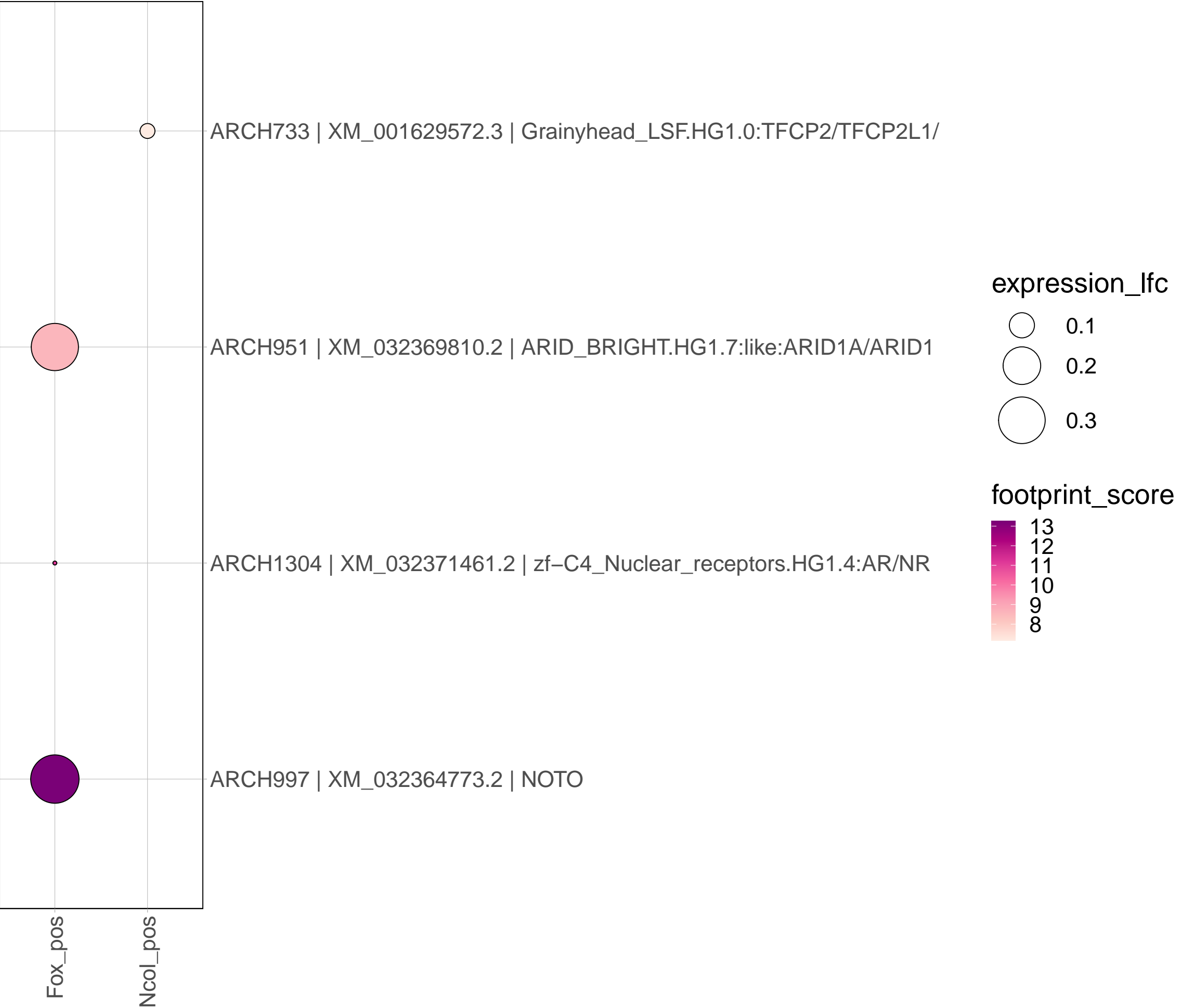
Nvec_vc1.1_XM_032381641.2
b(0,+)-type amino acid transporter ; AA_permease_2/



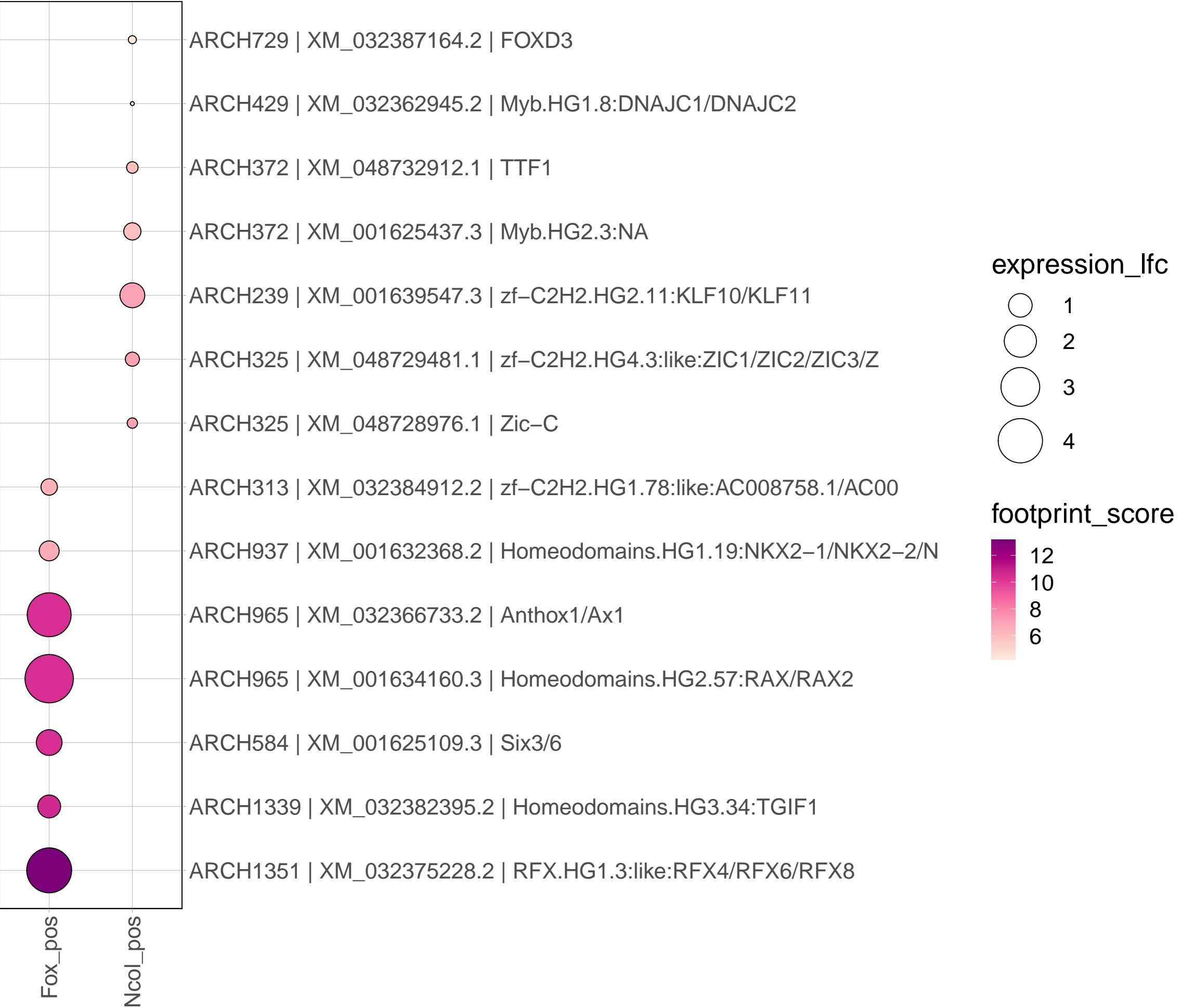
Nvec_vc1.1_XM_001632745.3
Protein NipSnap homolog 3A; NIPSNAP/NIPSNAP/



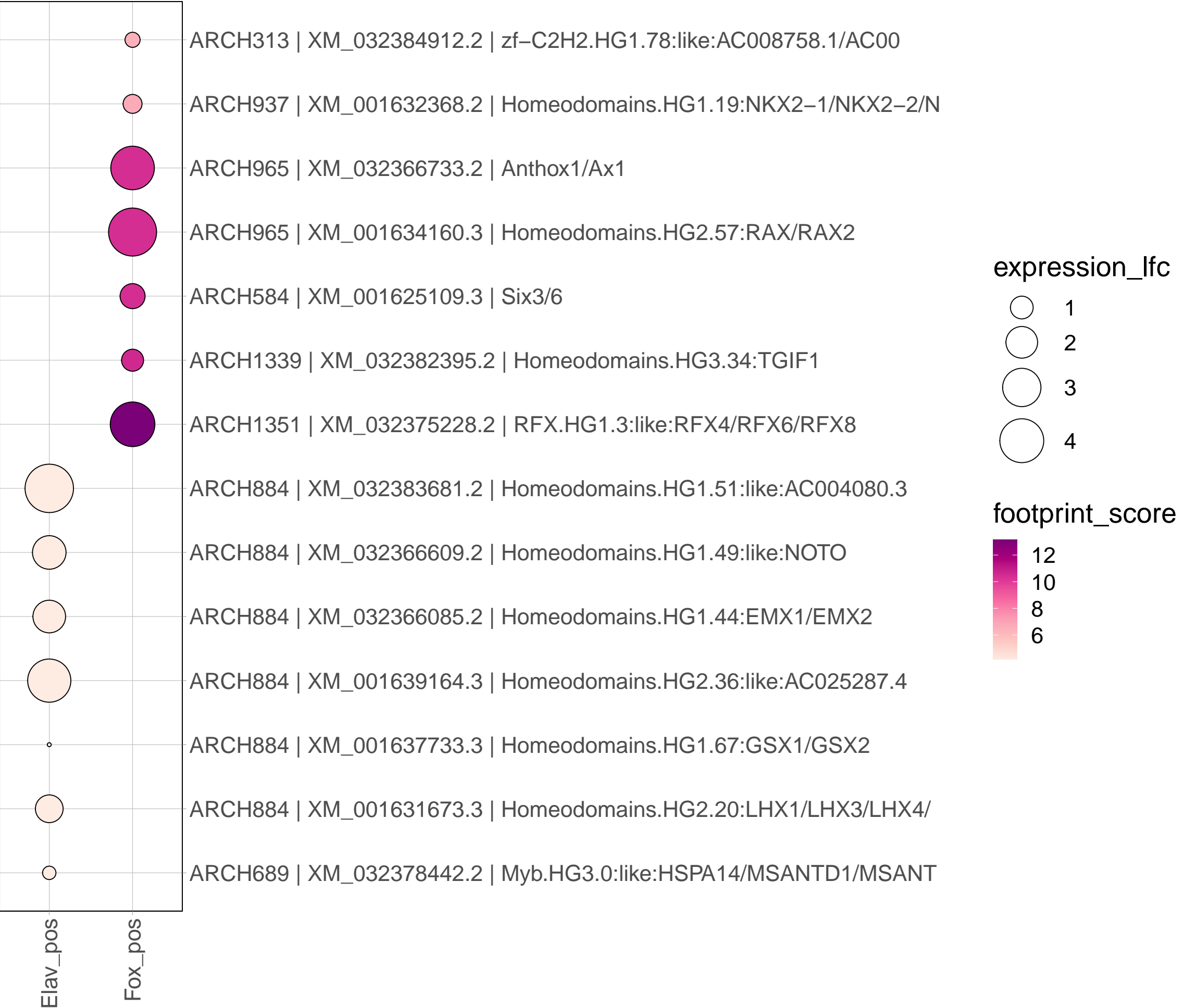
Nvec_vc1.1_XM_048724152.1
GDP–fucose protein O–fucosyltransfe; O–FucT/



Nvec_vc1.1_XM_048731865.1
GPCR–rhodopsin:like:GPR176:likeclu:; 7tm_1/

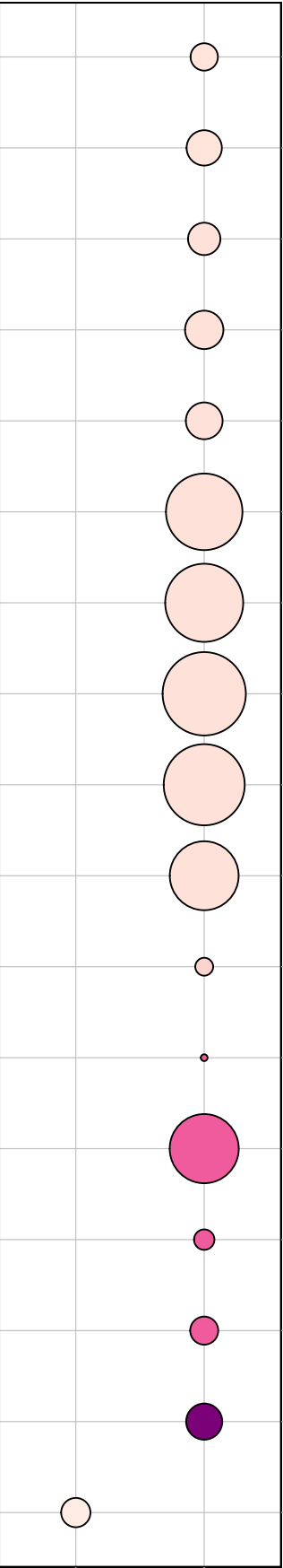


Nvec_vc1.1_XM_048731890.1
GPCR–rhodopsin:like:GPR176:likeclu:; 7tm_1/



Nvec_NVE13556

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ARCH1341 | XM_048733302.1 | HLH.HG1.33:MITF/TFE3/TFEB/TFEC

ARCH1341 | XM_032366030.2 | HLH.HG1.5:like:CCDC169-SOHLH2/SOHLH

ARCH160 | XM_032367127.2 | bZIP.HG2.10:DBP/HLF/TEF

ARCH152 | XM_032376926.2 | bZIP.HG3.2:like:MAF/MAFA/MAFB/NRL

ARCH152 | XM_001631543.3 | bZIP.HG1.15:CREB3/CREB3L1/CREB3L2/C

ARCH146 | XM_048730283.1 | bZIP.HG1.17:ATF3/JDP2

ARCH146 | XM_048730161.1 | Cnido-Fos1

ARCH146 | XM_032385896.2 | Cnido-Jun

ARCH146 | XM_032374029.2 | bZIP.HG1.6:JUN/JUNB/JUND

ARCH146 | XM_001631093.3 | bZIP.HG1.17:ATF3/JDP2

ARCH1023 | XM_001637508.3 | IRF.HG1.1:IRF1/IRF2/IRF3/IRF4/IRF5/

ARCH676 | XM_048725953.1 |

ARCH636 | XM_001631763.3 | PaxA

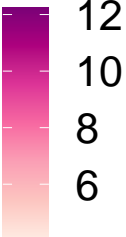
ARCH372 | XM_048732912.1 | TTF1

ARCH372 | XM_001625437.3 | Myb.HG2.3:NA

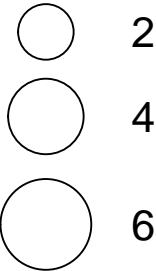
ARCH1425 | XM_001639701.3 | NvSox2

ARCH387 | XM_001625700.3 | NRF1.HG1.2:NRF1

footprint_score

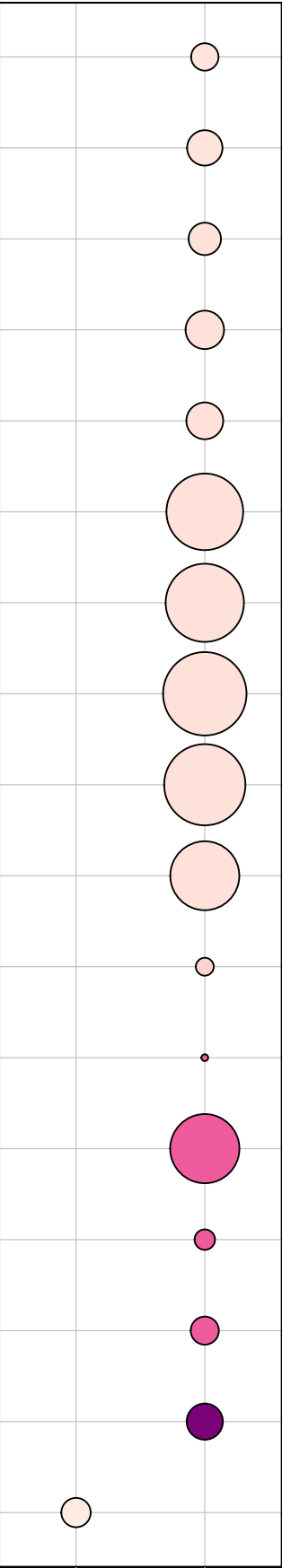


expression_lfc



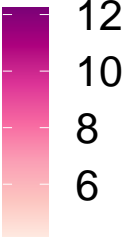
Nvec_vc1.1_XM_032387399.2

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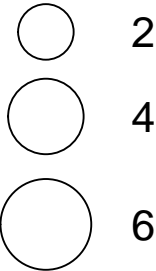


ARCH1341 XM_048733302.1 HLH.HG1.33:MITF/TFE3/TFEB/TFEC
ARCH1341 XM_032366030.2 HLH.HG1.5:like:CCDC169-SOHLH2/SOHLH
ARCH160 XM_032367127.2 bZIP.HG2.10:DBP/HLF/TEF
ARCH152 XM_032376926.2 bZIP.HG3.2:like:MAF/MAFA/MAFB/NRL
ARCH152 XM_001631543.3 bZIP.HG1.15:CREB3/CREB3L1/CREB3L2/C
ARCH146 XM_048730283.1 bZIP.HG1.17:ATF3/JDP2
ARCH146 XM_048730161.1 Cnido-Fos1
ARCH146 XM_032385896.2 Cnido-Jun
ARCH146 XM_032374029.2 bZIP.HG1.6:JUN/JUNB/JUND
ARCH146 XM_001631093.3 bZIP.HG1.17:ATF3/JDP2
ARCH1023 XM_001637508.3 IRF.HG1.1:IRF1/IRF2/IRF3/IRF4/IRF5/
ARCH676 XM_048725953.1
ARCH636 XM_001631763.3 PaxA
ARCH372 XM_048732912.1 TTF1
ARCH372 XM_001625437.3 Myb.HG2.3:NA
ARCH1425 XM_001639701.3 NvSox2
ARCH387 XM_001625700.3 NRF1.HG1.2:NRF1

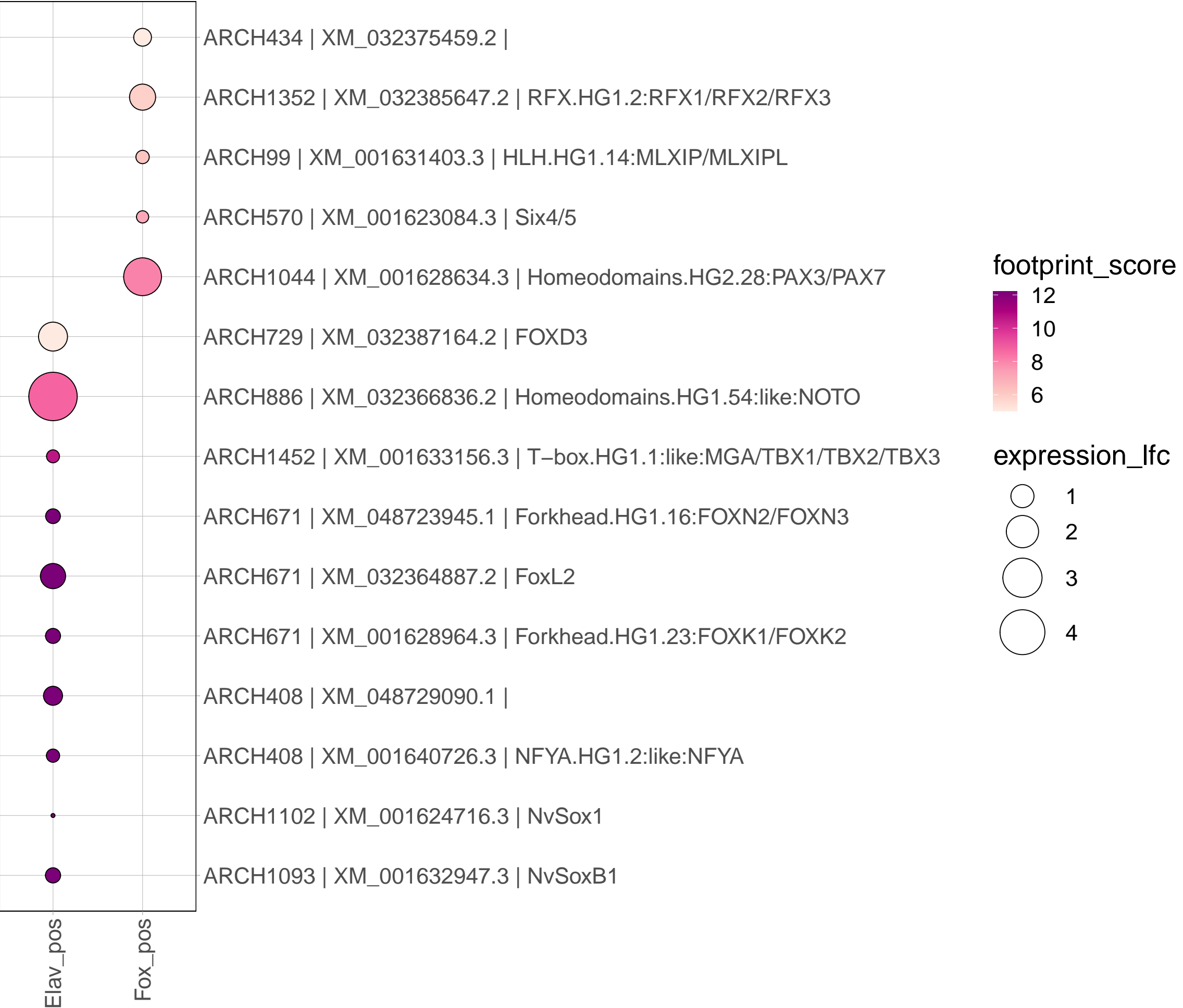
footprint_score



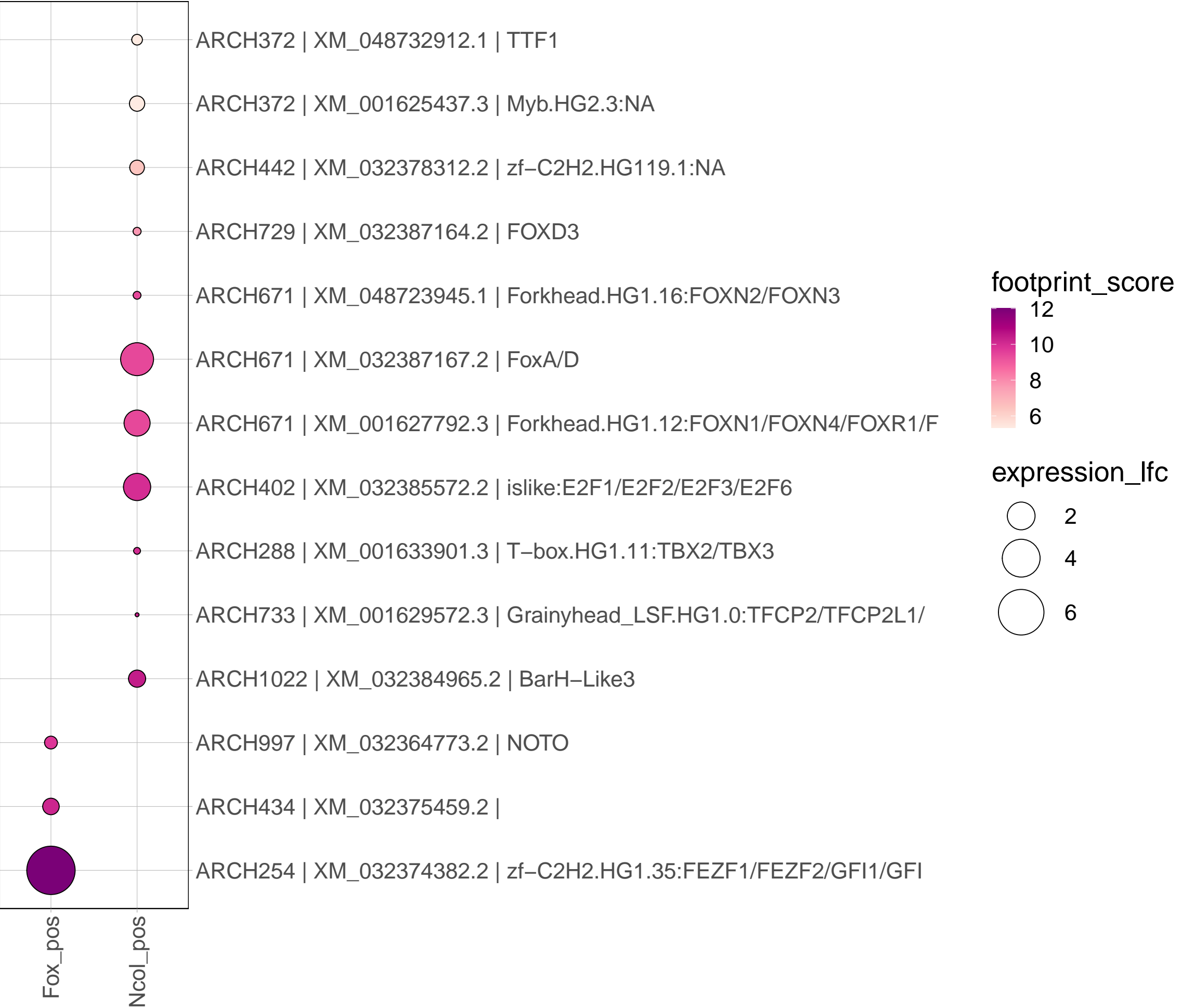
expression_lfc



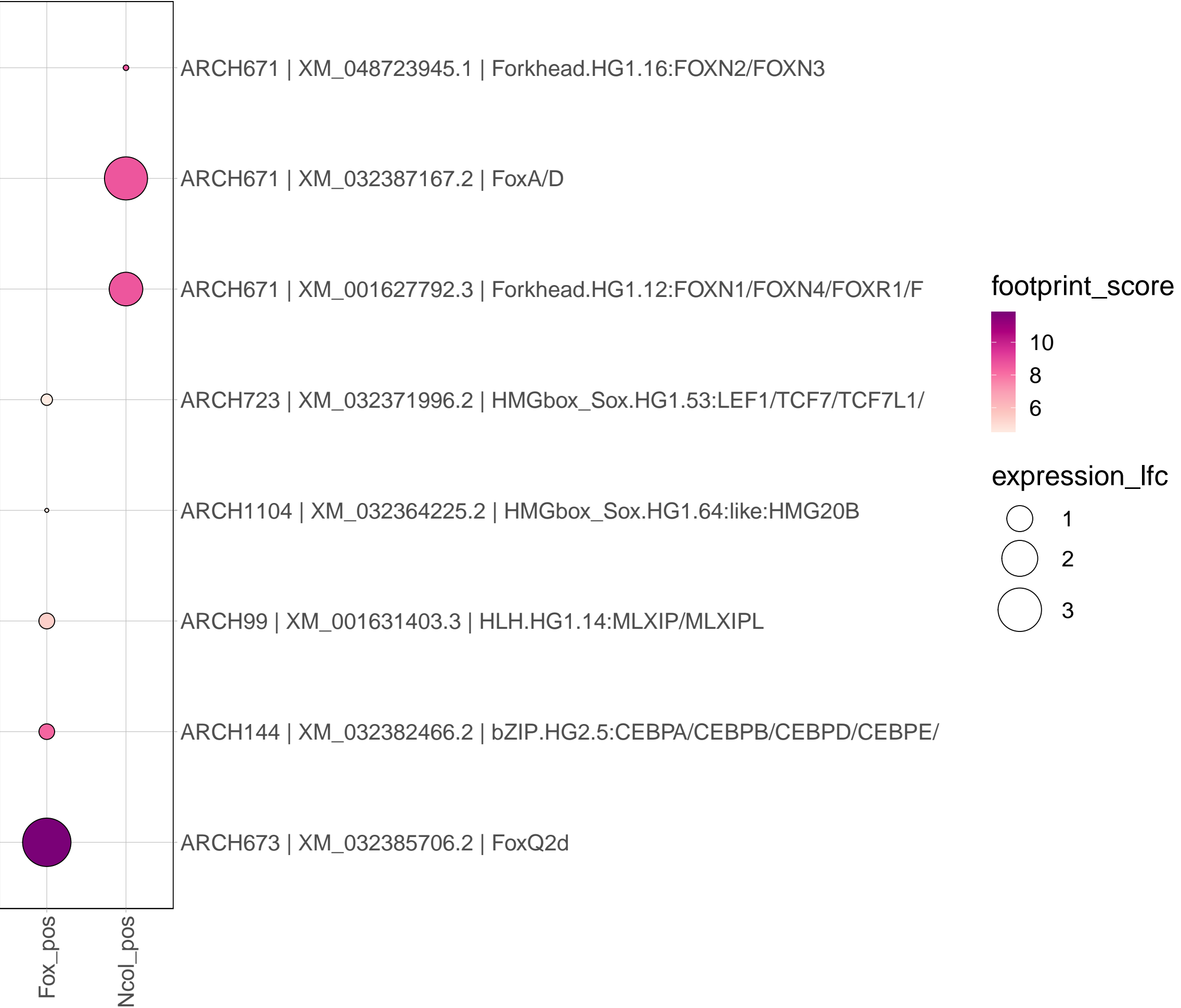
Nvec_vc1.1_XM_048729747.1
Kinesin-like protein KIF16B; Kinesin/FHA/



Nvec_vc1.1_XM_032382956.2
Galectin; VHL/

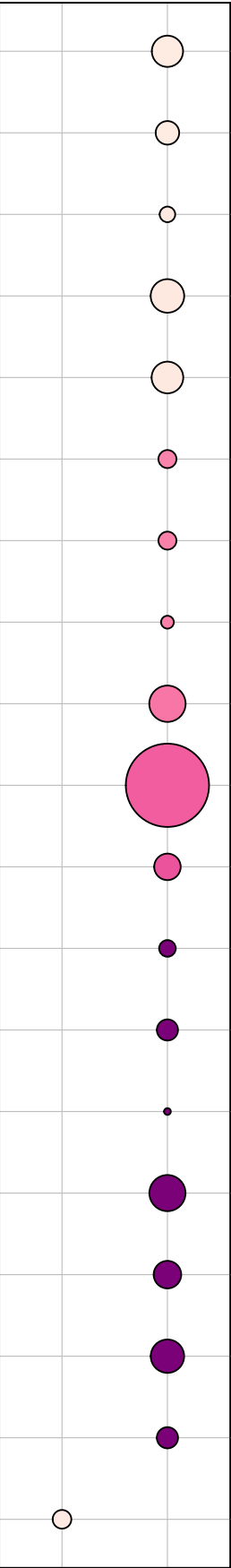


Nvec_vc1.1_XM_001627245.3
cDNA FLJ56852, highly similar to Hu; I_LWEQ/



Nvec_vc1.1_XM_001640356.3

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ARCH404 XM_032367573.2 SAND.HG1.1:like:GMEB1/GMEB2
ARCH1457 XM_001635374.3 Homeodomains.HG3.7:PBX1/PBX2/PBX3/P
ARCH1479 XM_048726885.1 zf-C2H2.HG1.3:ZBTB3
ARCH1479 XM_001638761.3 zf-C2H2.HG16.2:like:TRERF1/ZNF541
ARCH1479 XM_001633508.3 zf-C2H2.HG1.3:ZBTB3
ARCH239 XM_001639547.3 zf-C2H2.HG2.11:KLF10/KLF11
ARCH723 XM_032371996.2 HMGbox_Sox.HG1.53:LEF1/TCF7/TCF7L1/
ARCH1104 XM_032364225.2 HMGbox_Sox.HG1.64:like:HMG20B
ARCH202 XM_032363168.2 zf-met.HG3.0:like:ZMAT1/ZMAT3/ZMAT4
ARCH254 XM_032374382.2 zf-C2H2.HG1.35:FEZF1/FEZF2/GFI1/GFI
ARCH238 XM_032387218.2 zf-C2H2.HG57.1:like:ZBTB48
ARCH1591 XM_048722339.1 HLH.HG1.0:AC019117.4/AHR/AHRR/ARNT/
ARCH1591 XM_001635172.3 HLH.HG1.38:MXD1/MXD3/MXD4/MXI1
ARCH1571 XM_001623131.3 NFYB_NFYC.HG3.0:TAF11/TAF11L3/TAF11
ARCH1435 XM_001628041.3 HMGbox_Sox.HG1.24:HMGB1/HMGB2/HMGB3
ARCH1435 XM_001627423.3 HMGbox_Sox.HG1.10:BBX
ARCH1434 XM_032372264.2 NvSoxC
ARCH115 XM_001628279.3 HLH.HG1.76:ATOH8
ARCH94 XM_032384172.2 zf-C2H2.HG1.40:GZF1/ZBTB24

footprint_score



10
8
6

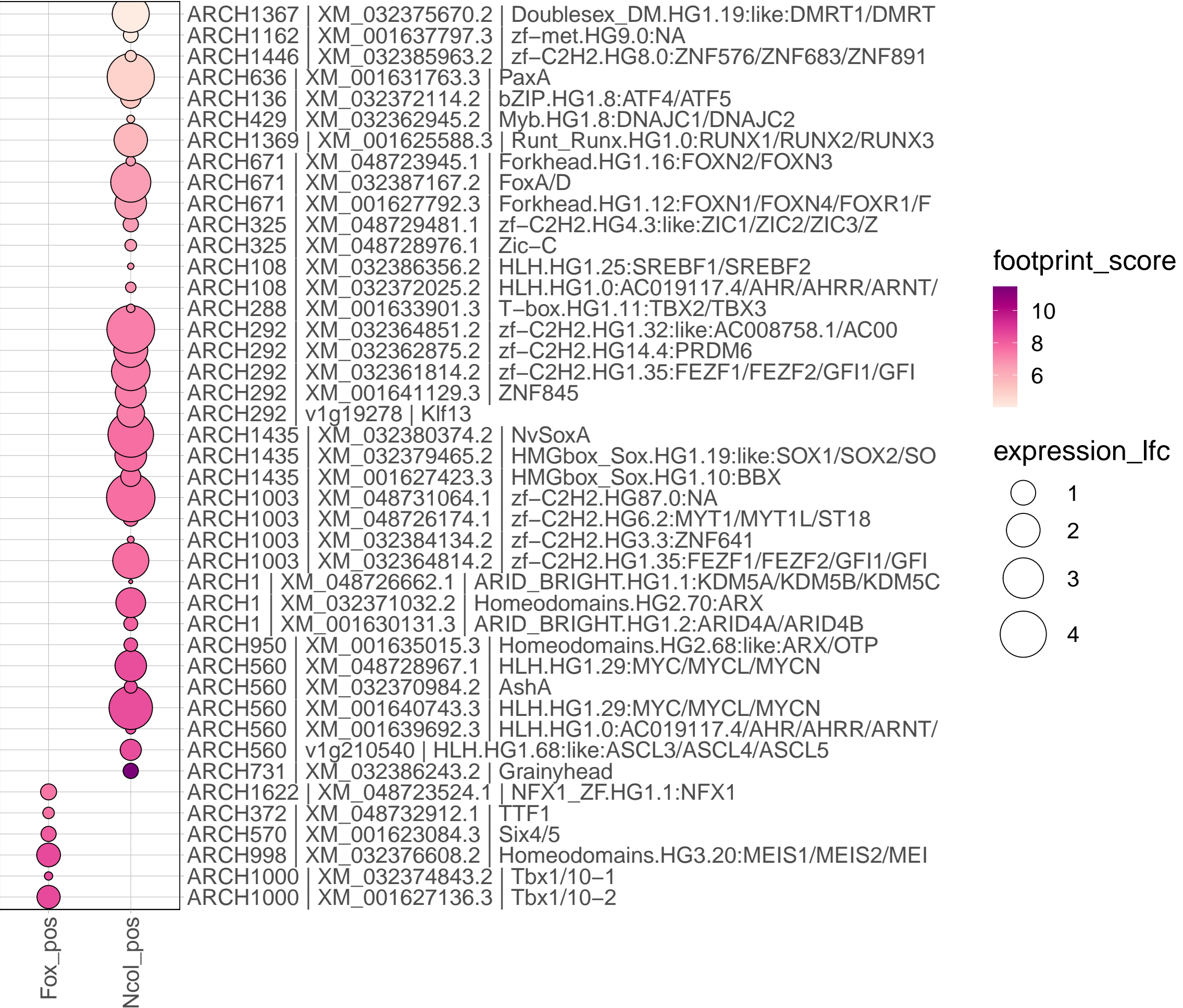
expression_lfc



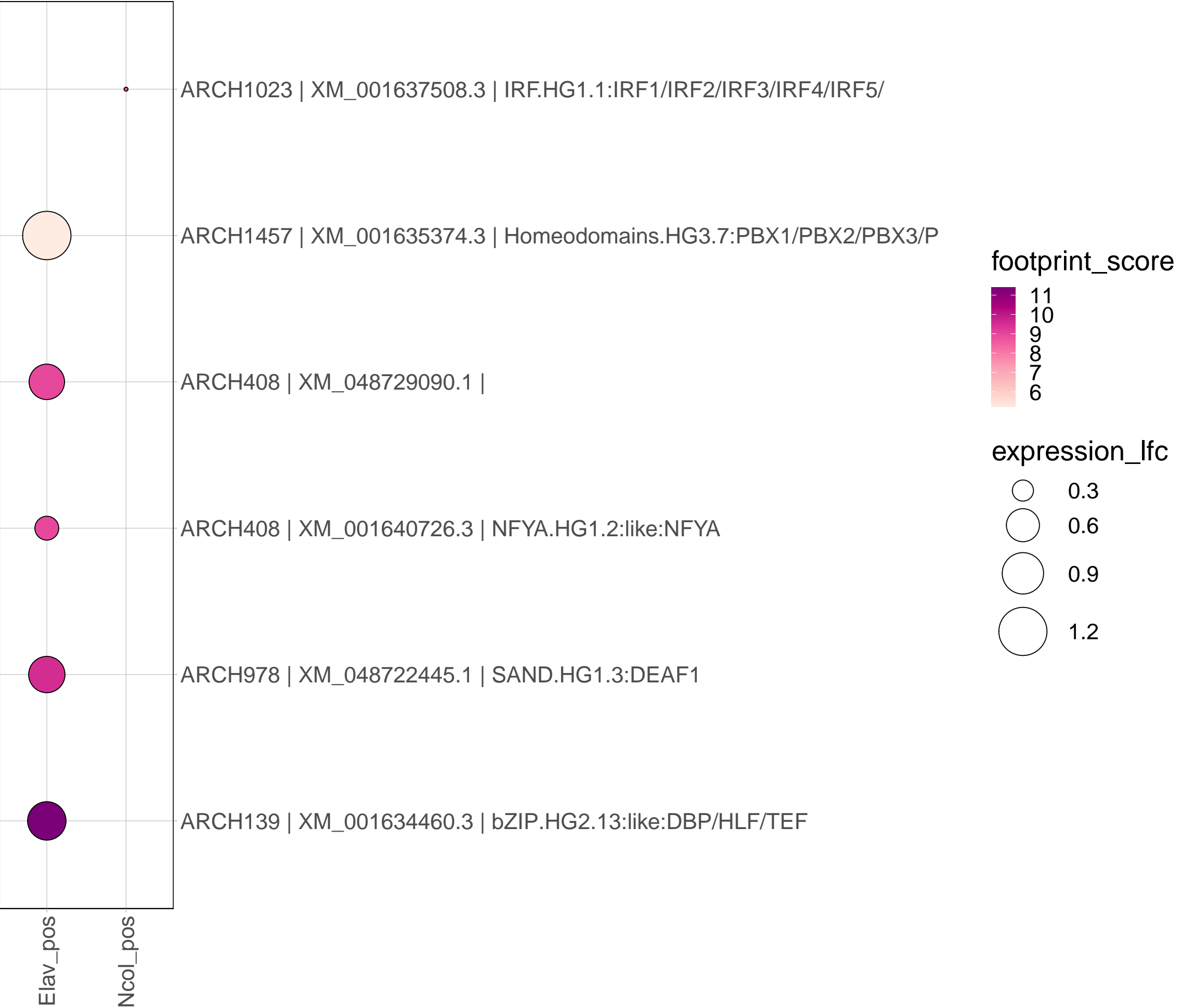
2
4
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Elav_pos
Fox_pos

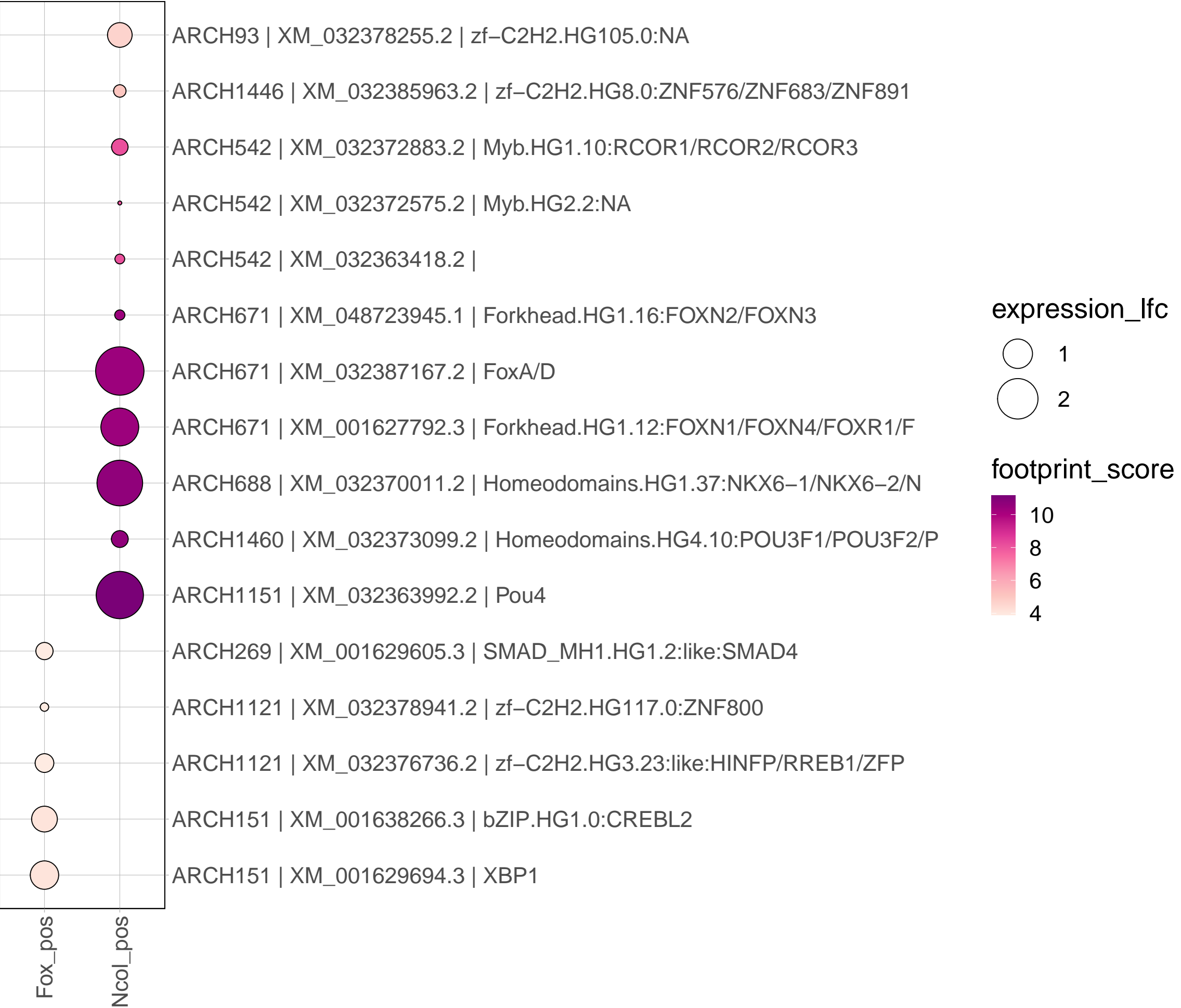
Nvec_vc1.1_XM_032377937.2
ion-neuro-gated:like:GABRA1/GABRA2/; Neur_chan_LBD/Neur_chan_memb/



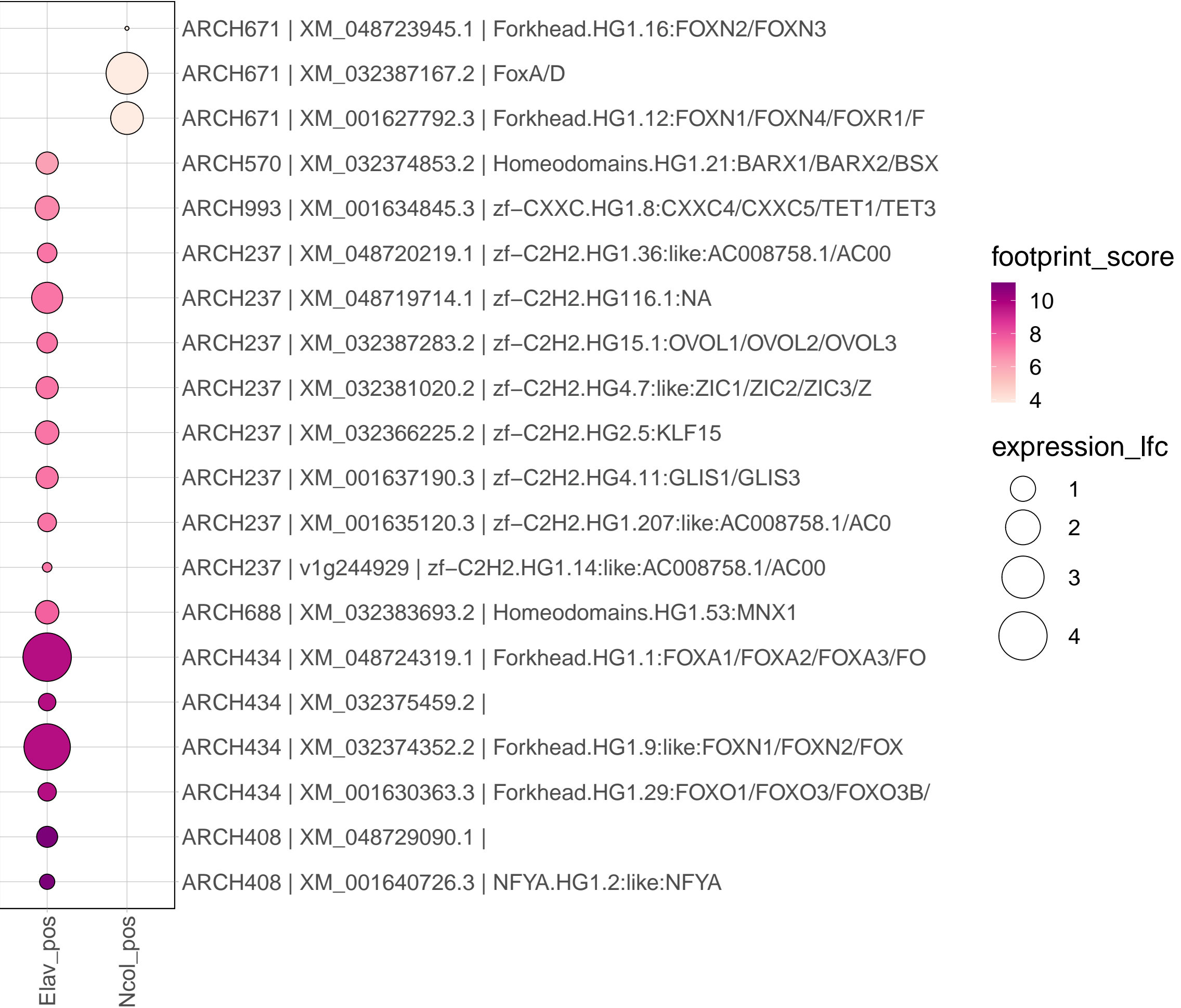
Nvec_vc1.1_XM_001636964.3
cDNA FLJ77769, highly similar to Ho; NAD_kinase/



Nvec_vc1.1_XM_001623818.3
ion-NaKCa:like:KCNA1/KCNA2/KCNA3/KC; BTB_2/Ion_trans/

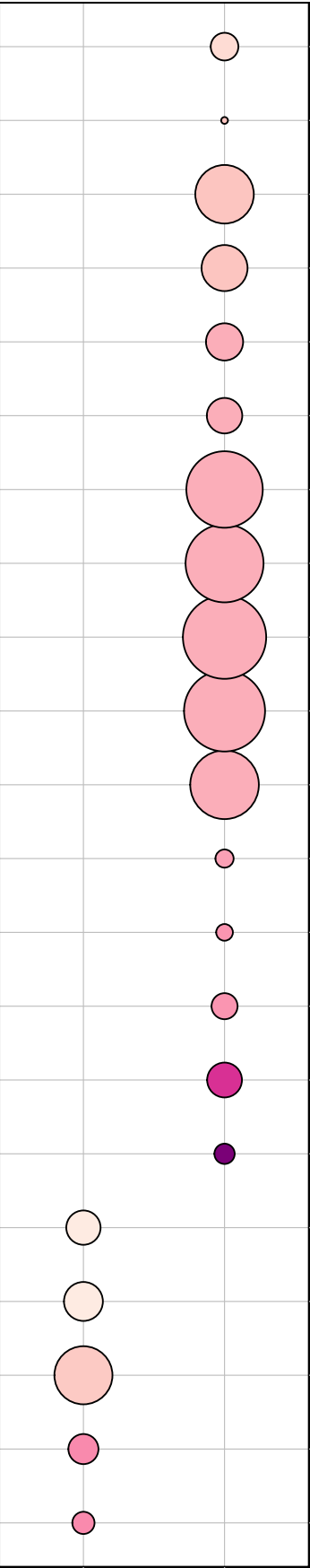


Nvec_vc1.1_XM_001637022.3
GPCR-rhodopsin:NA; 7tm_1/



Nvec_vc1.1_XM_048731921.1

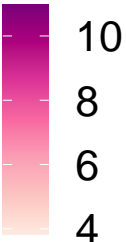
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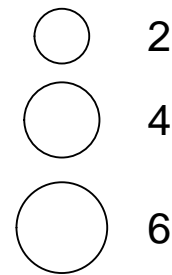
Elav_pos

Ncol_pos

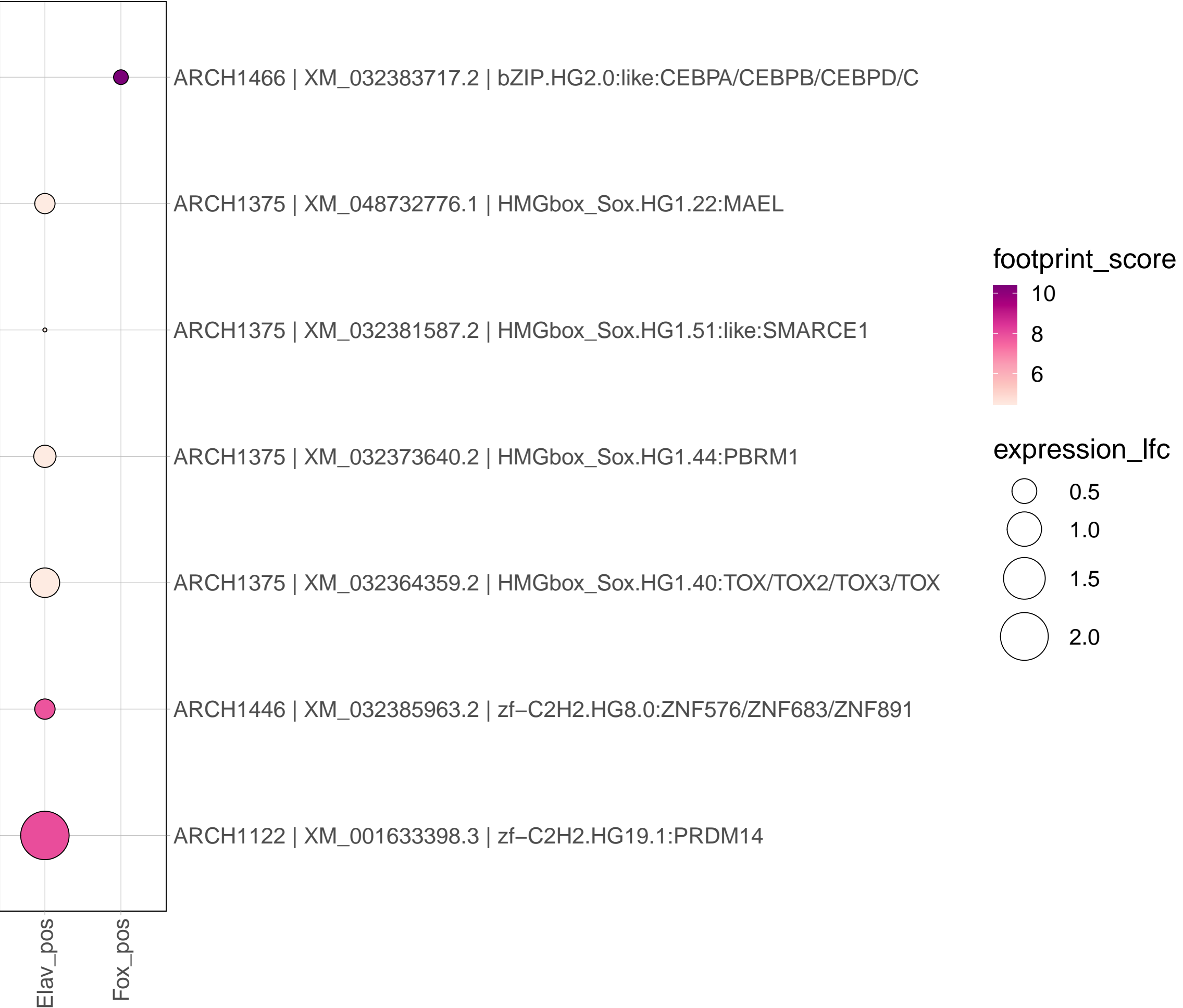
footprint_score



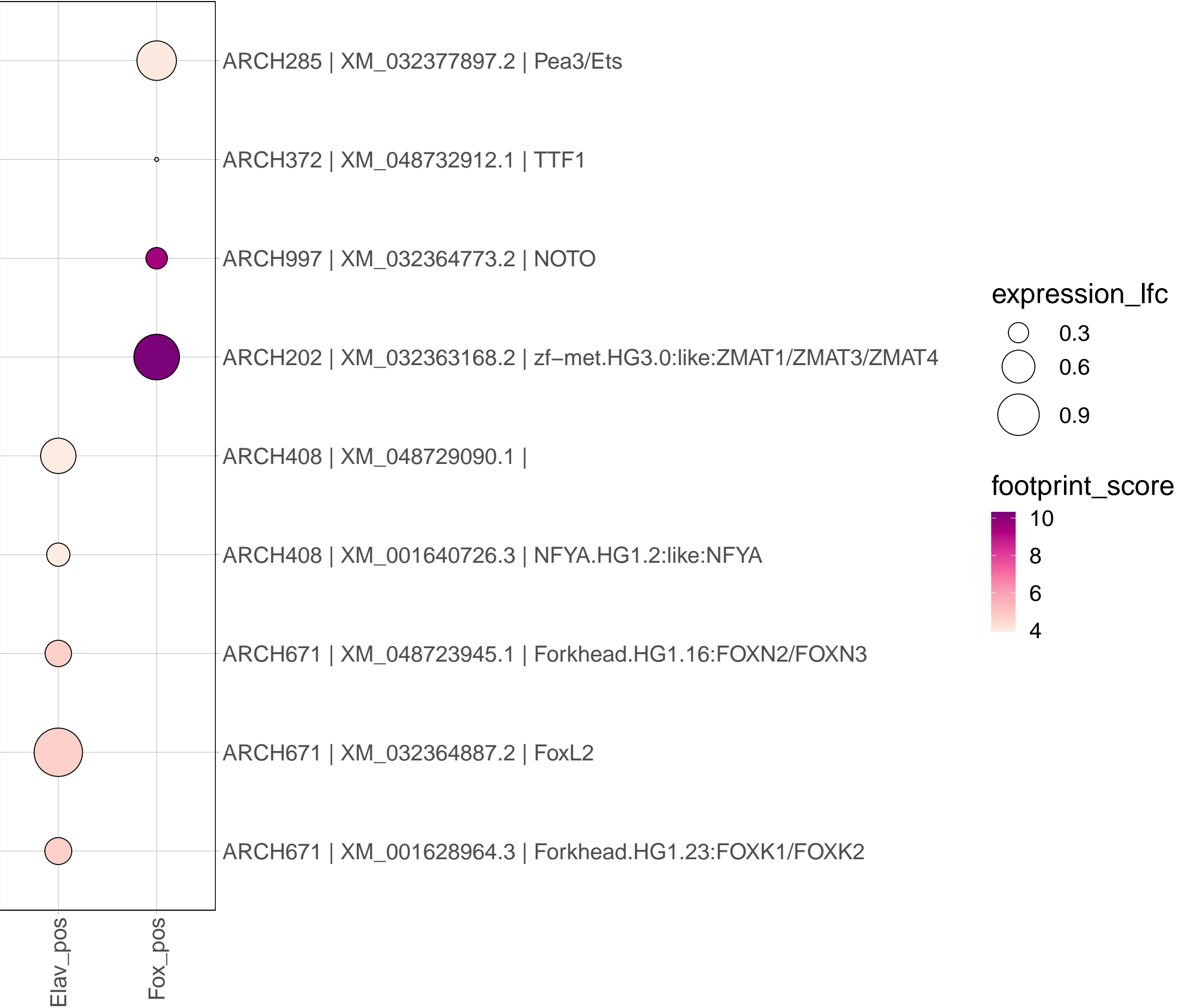
expression_lfc



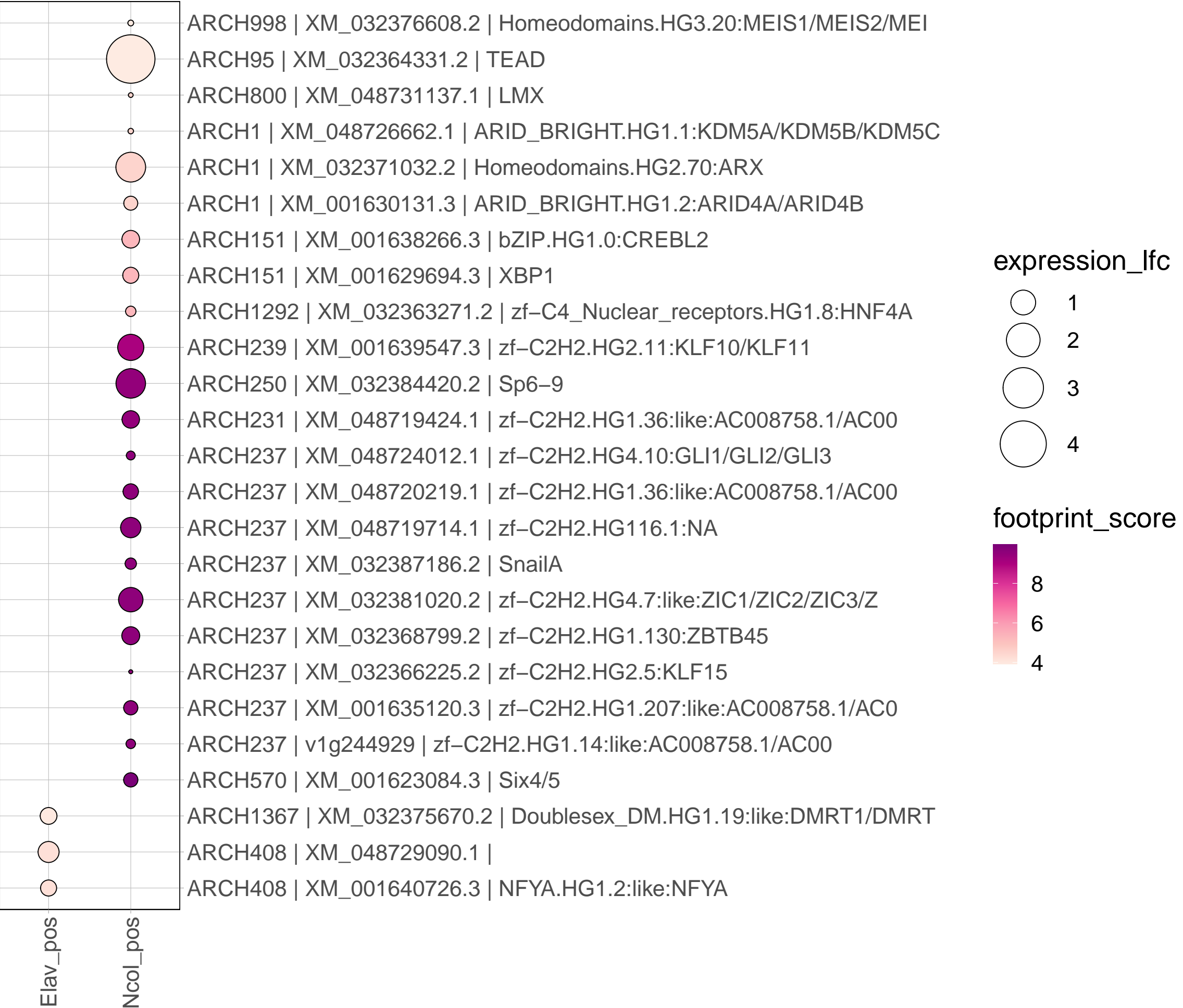
Nvec_vc1.1_XM_001631508.3
YTH:YTHDC1; YTH.HG1.0:YTHDC1



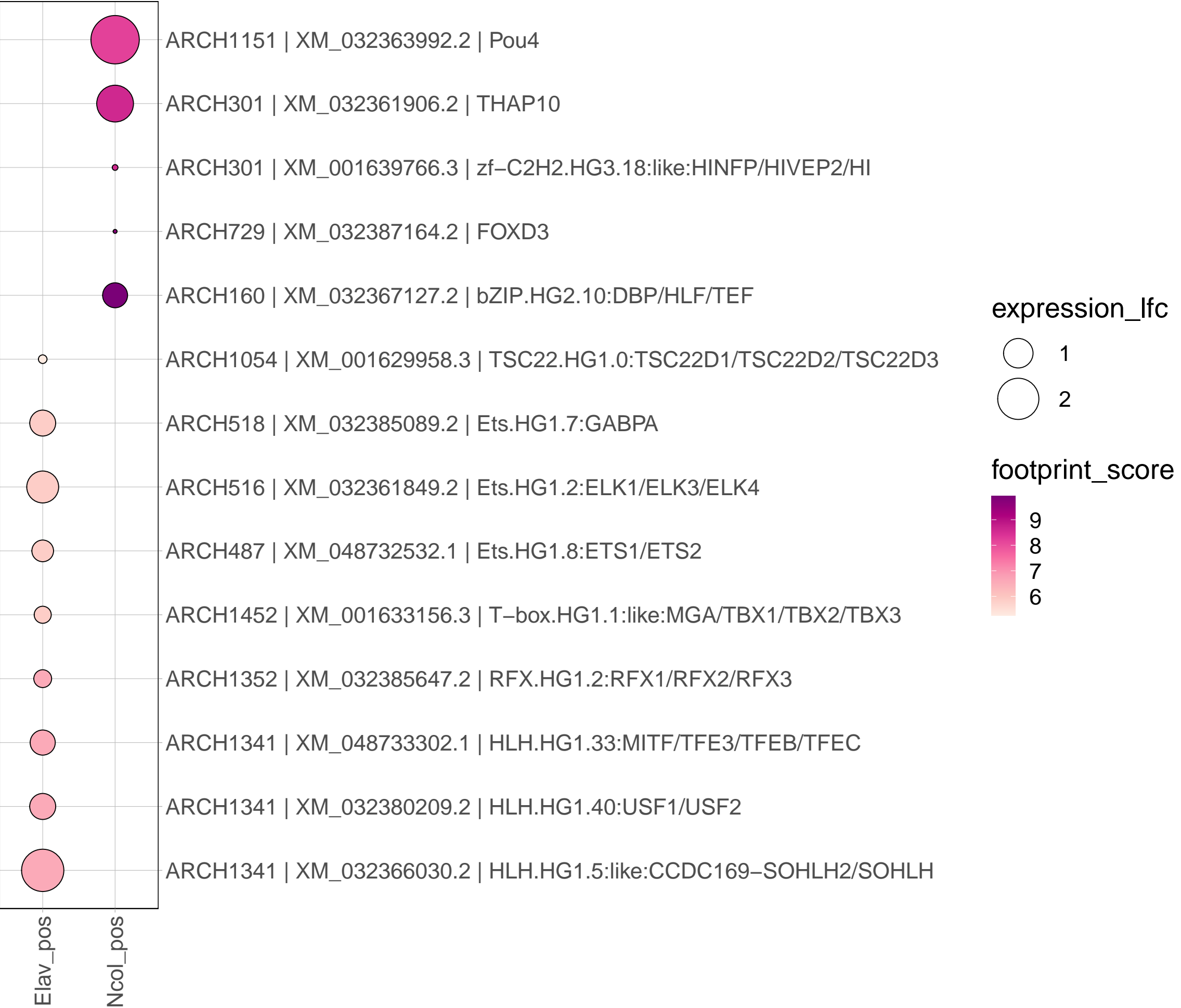
Nvec_vc1.1_XM_032370981.2
Synaptotagmin:SYT12; Synaptotagmin.HG1.38:SYT12



Nvec_vc1.1_XM_032384420.2
Sp6-9; SP6/SP7/SP8/SP9



Nvec_vc1.1_XM_001641986.3
GPCR–rhodopsin:NA; 7tm_1/7tm_1/



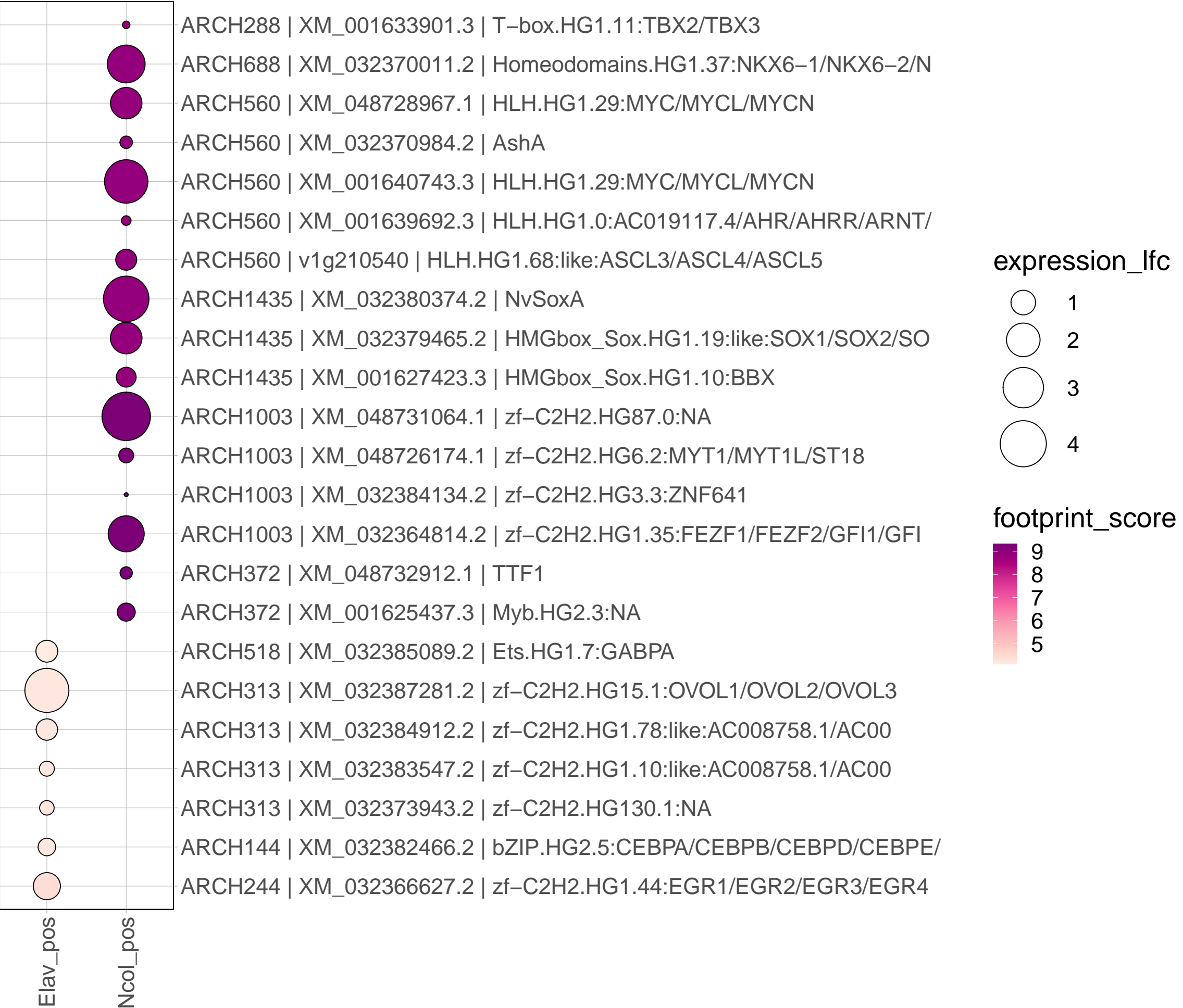
Nvec_vc1.1_XM_001637149.3
RAP1A, member of RAS oncogene famil; Ras/



Nvec_vc1.1_XM_001628711.3
RRM:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPC; RRM.HG17.3:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPC

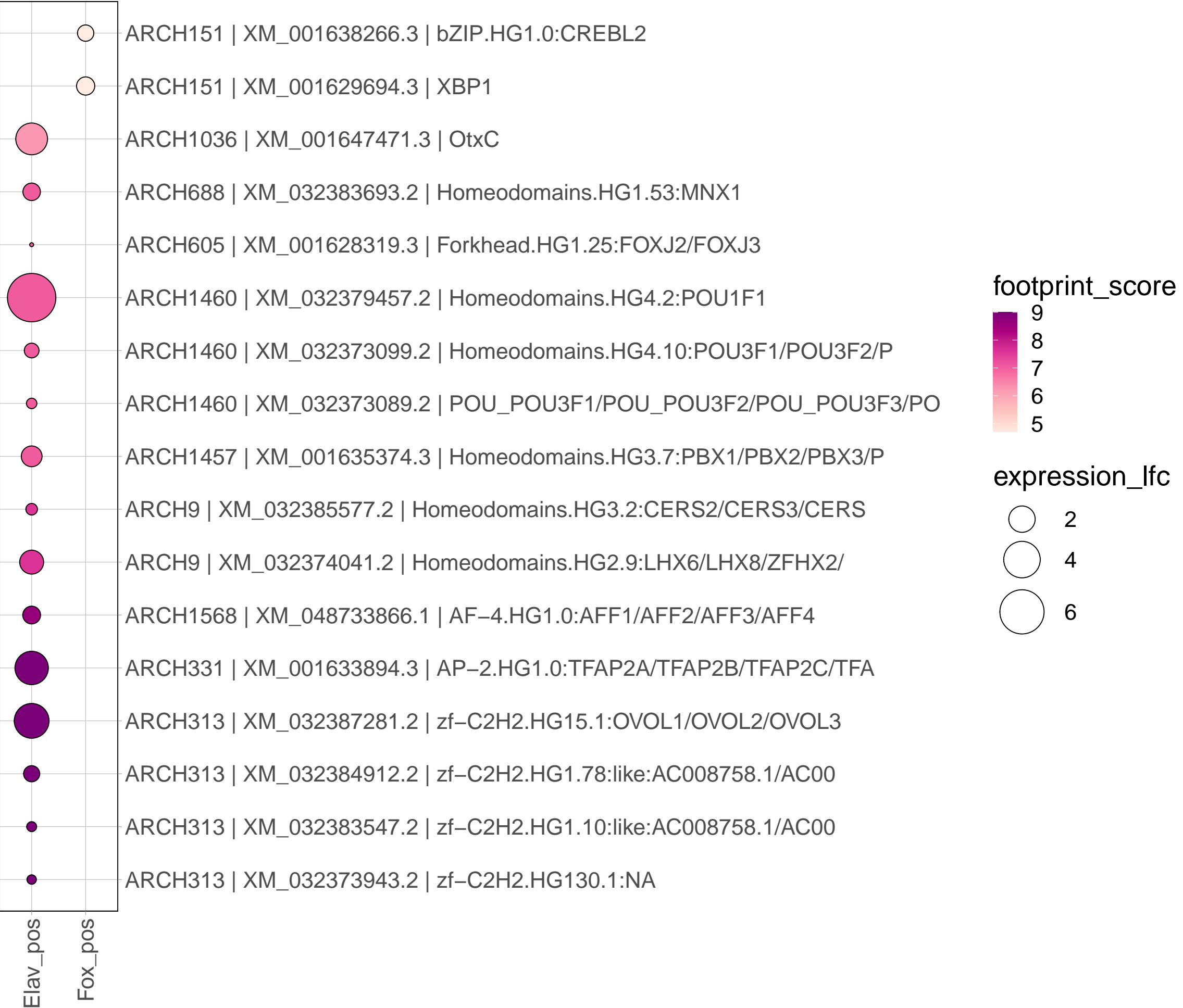


Nvec_vc1.1_XM_001630845.3
ion-glutamate:NA; Ion_trans_2/

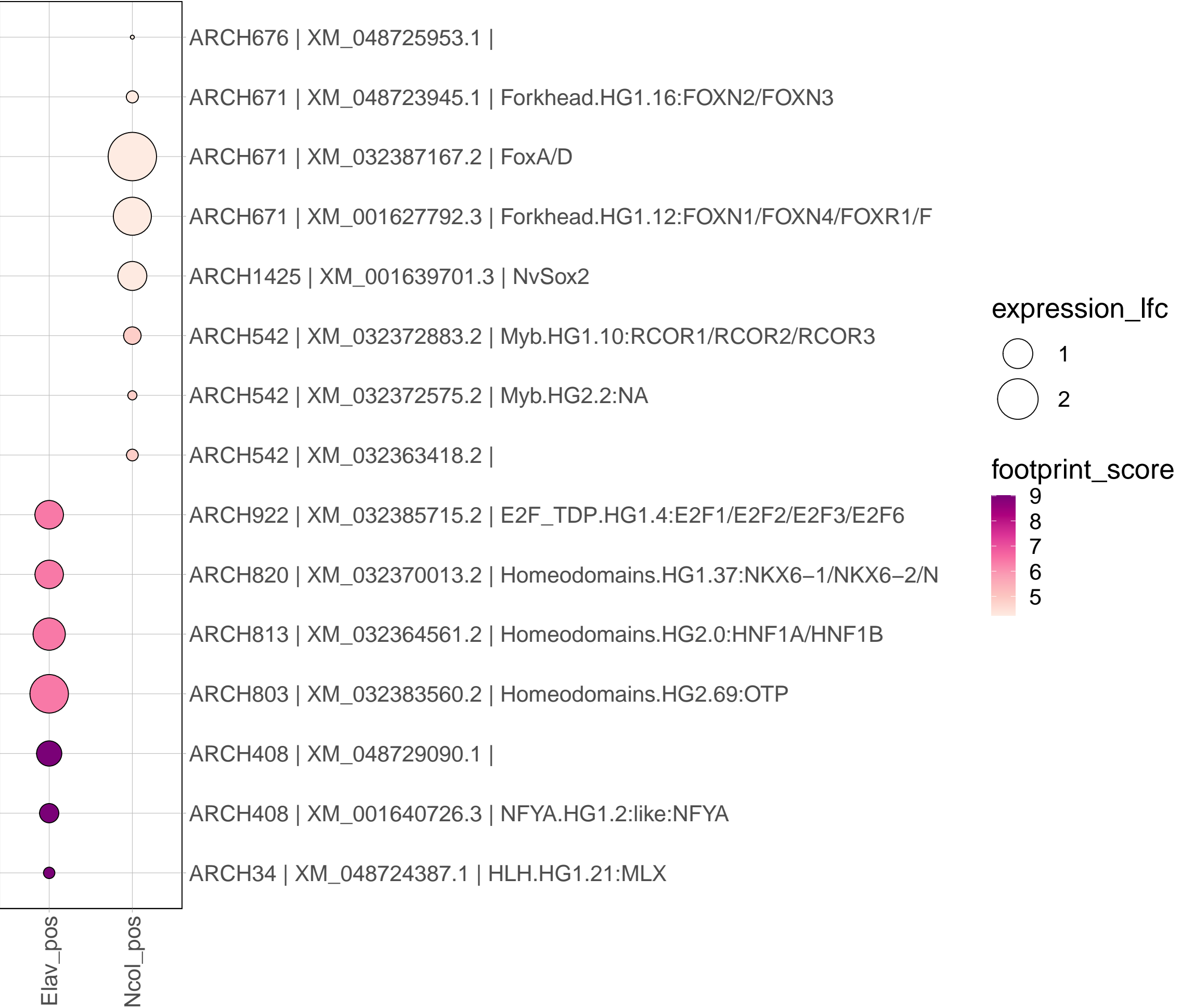


Nvec_vc1.1_XM_032379049.2

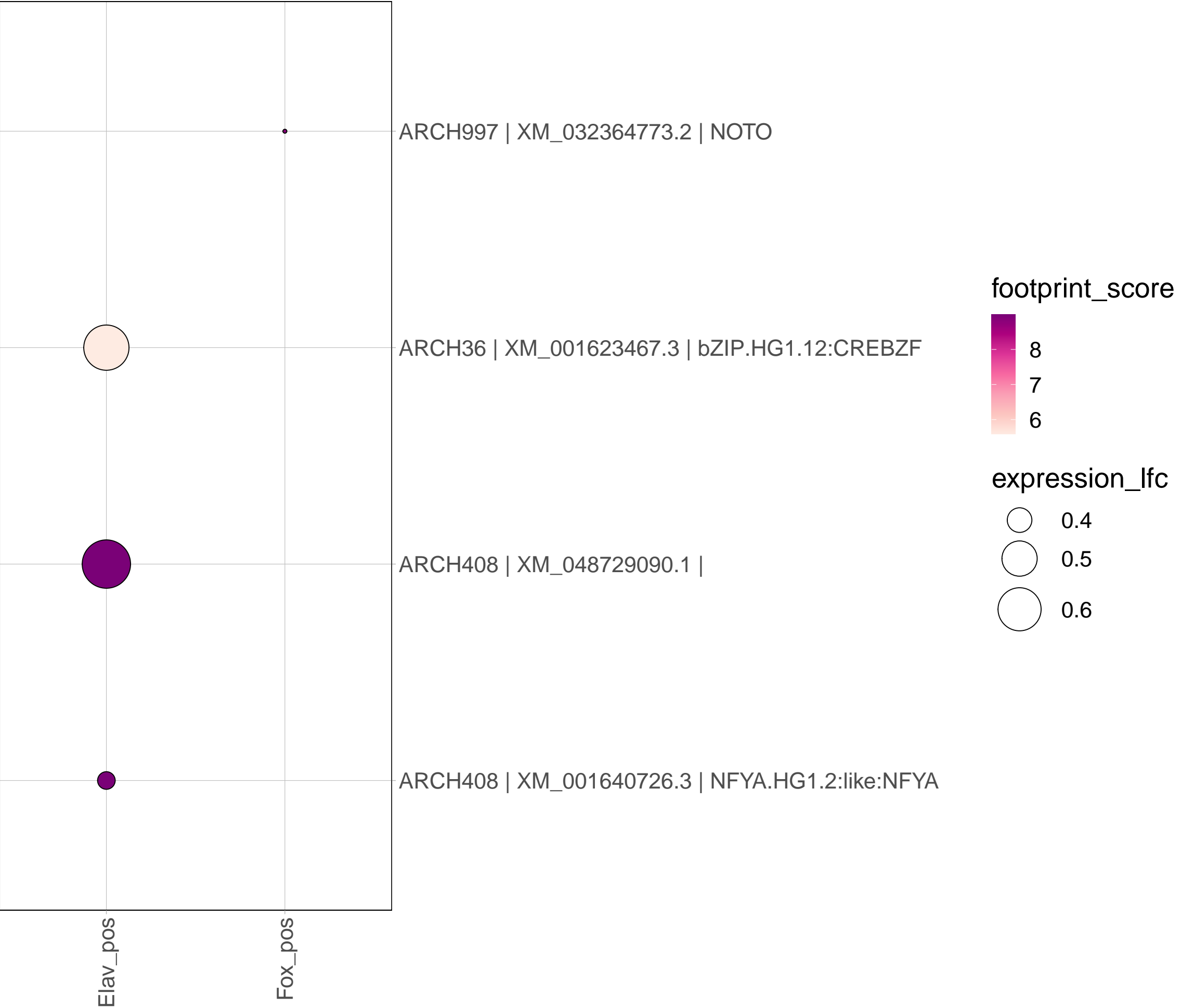
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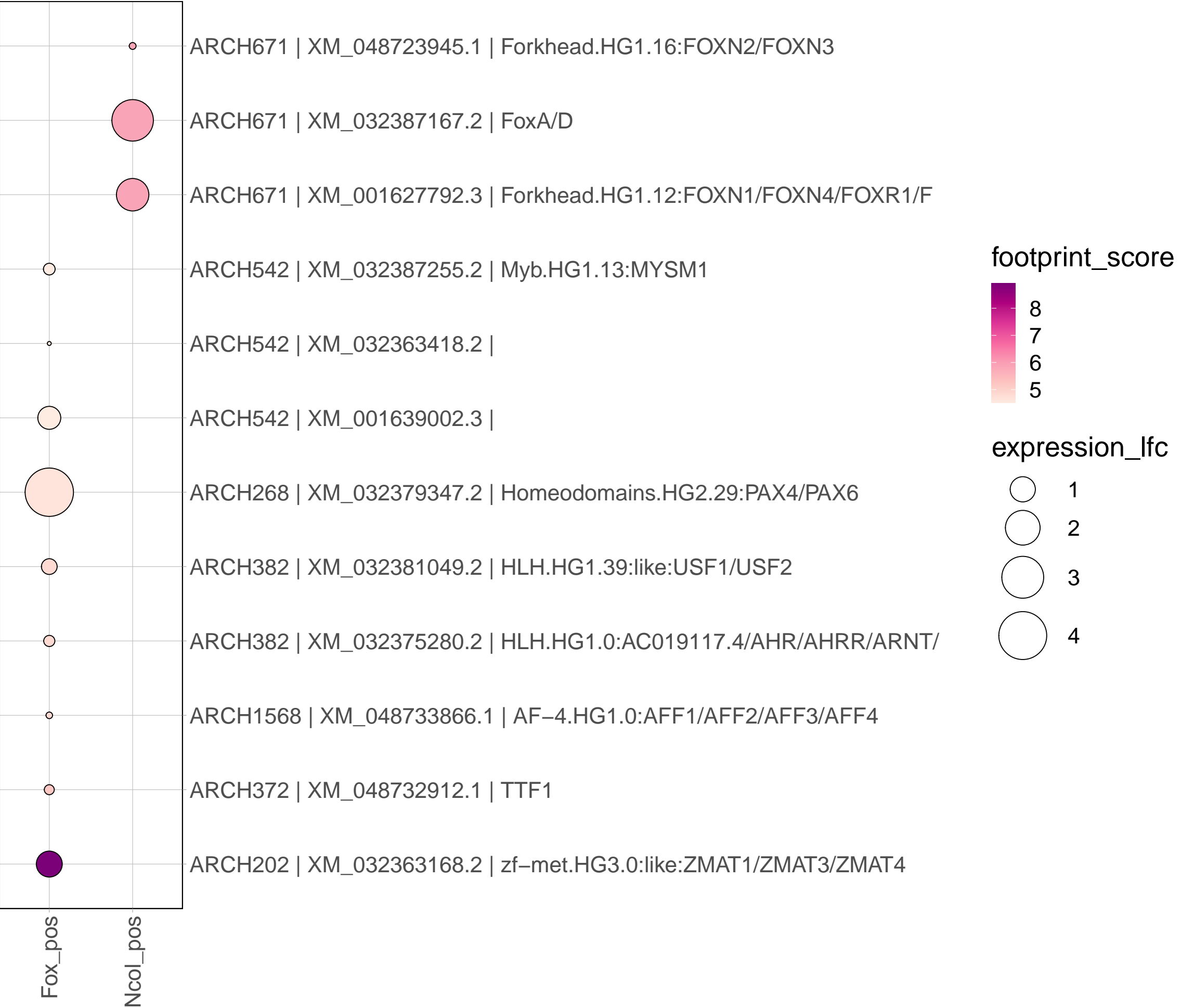
Nvec_vc1.1_XM_032376506.2
Zinc finger protein 862; Dimer_Tnp_hAT/



Nvec_vc1.1_XM_032370048.2
zf-CCCH:MKRN1/MKRN2/MKRN3; zf-CCCH.HG1.17:MKRN1/MKRN2/MKRN3



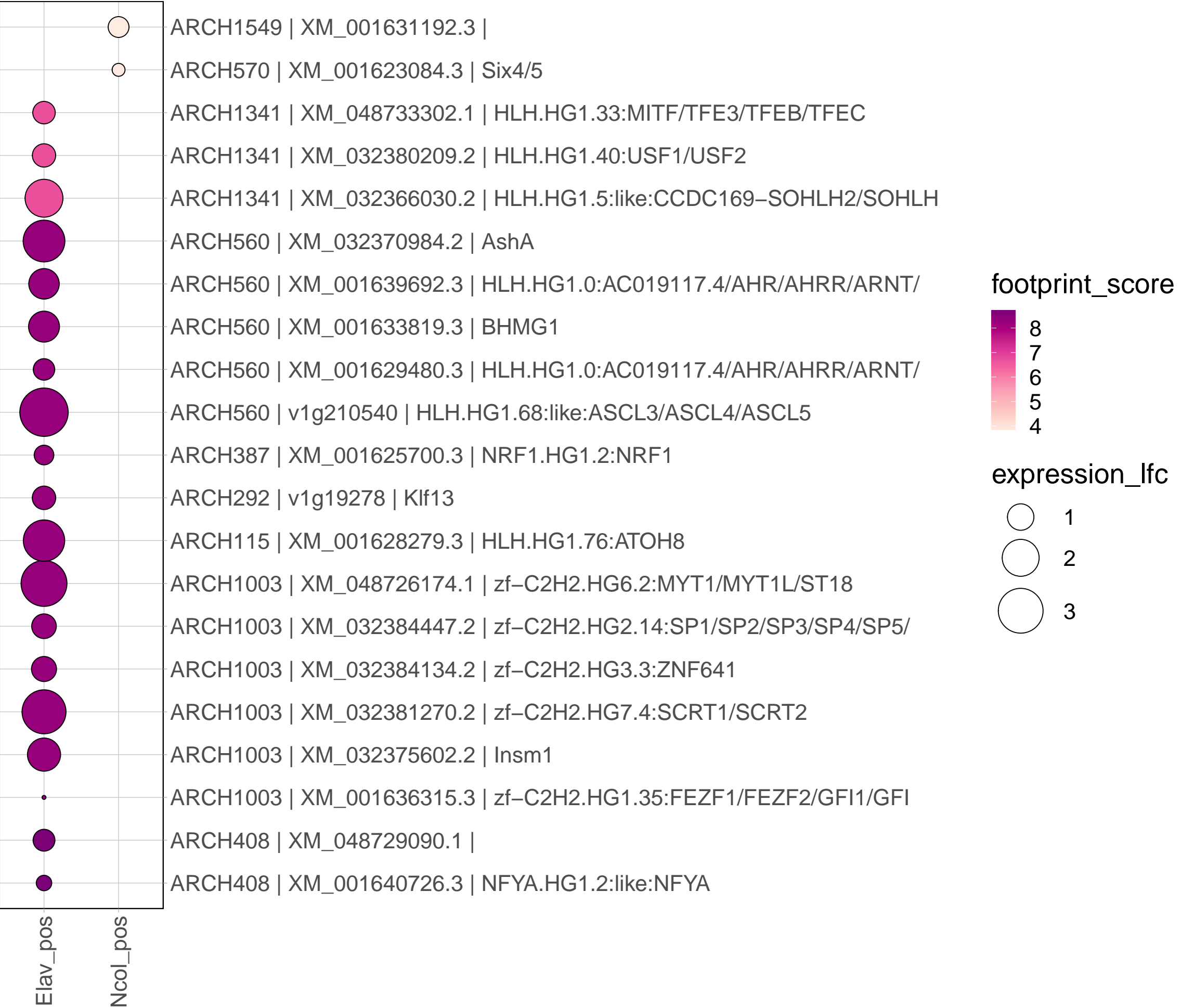
Nvec_vc1.1_XM_048731597.1
Allograft inflammatory factor 1; EF-hand_7/



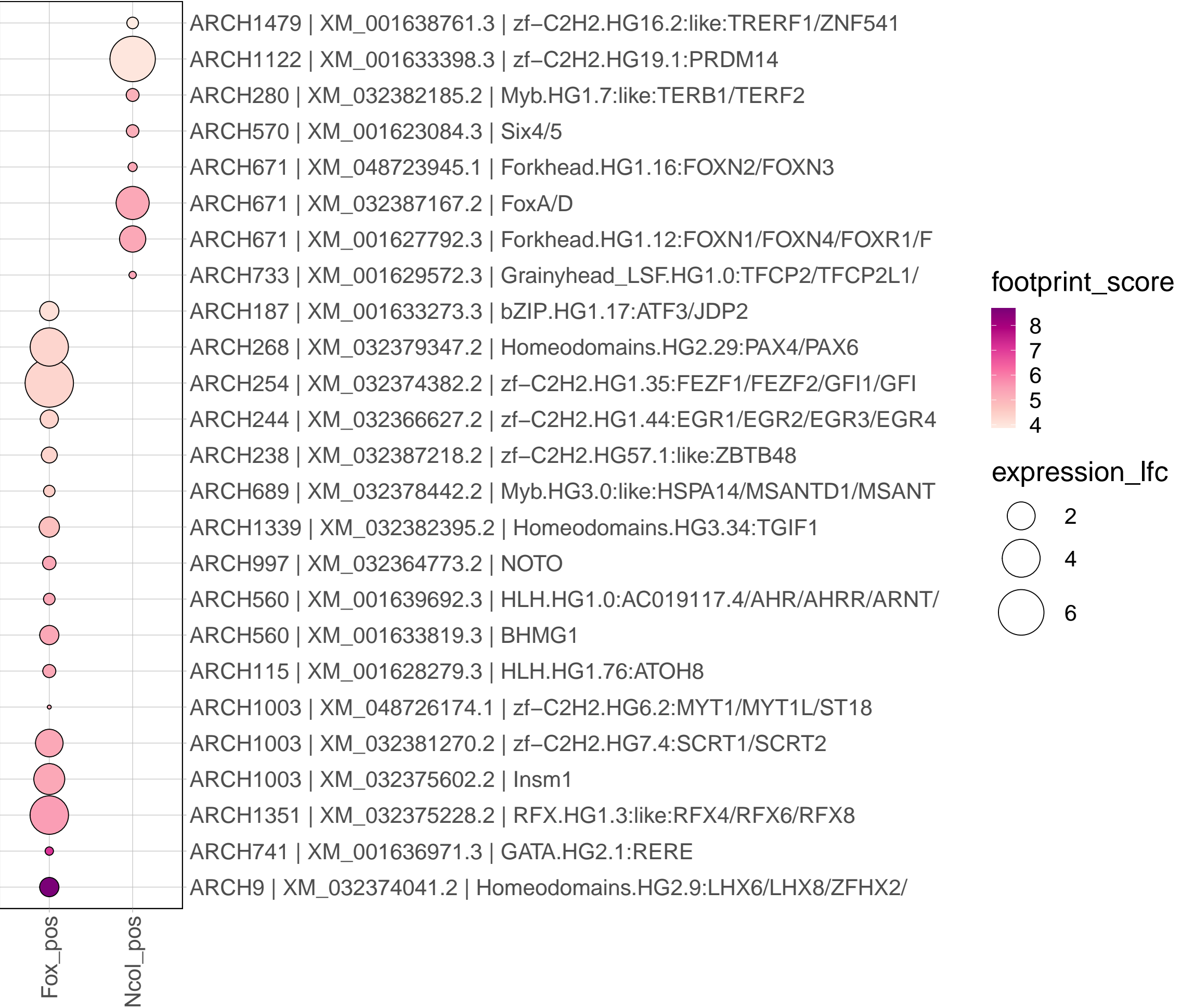
Nvec_vc1.1_XM_048731252.1
OTU domain-containing protein 3; OTU/



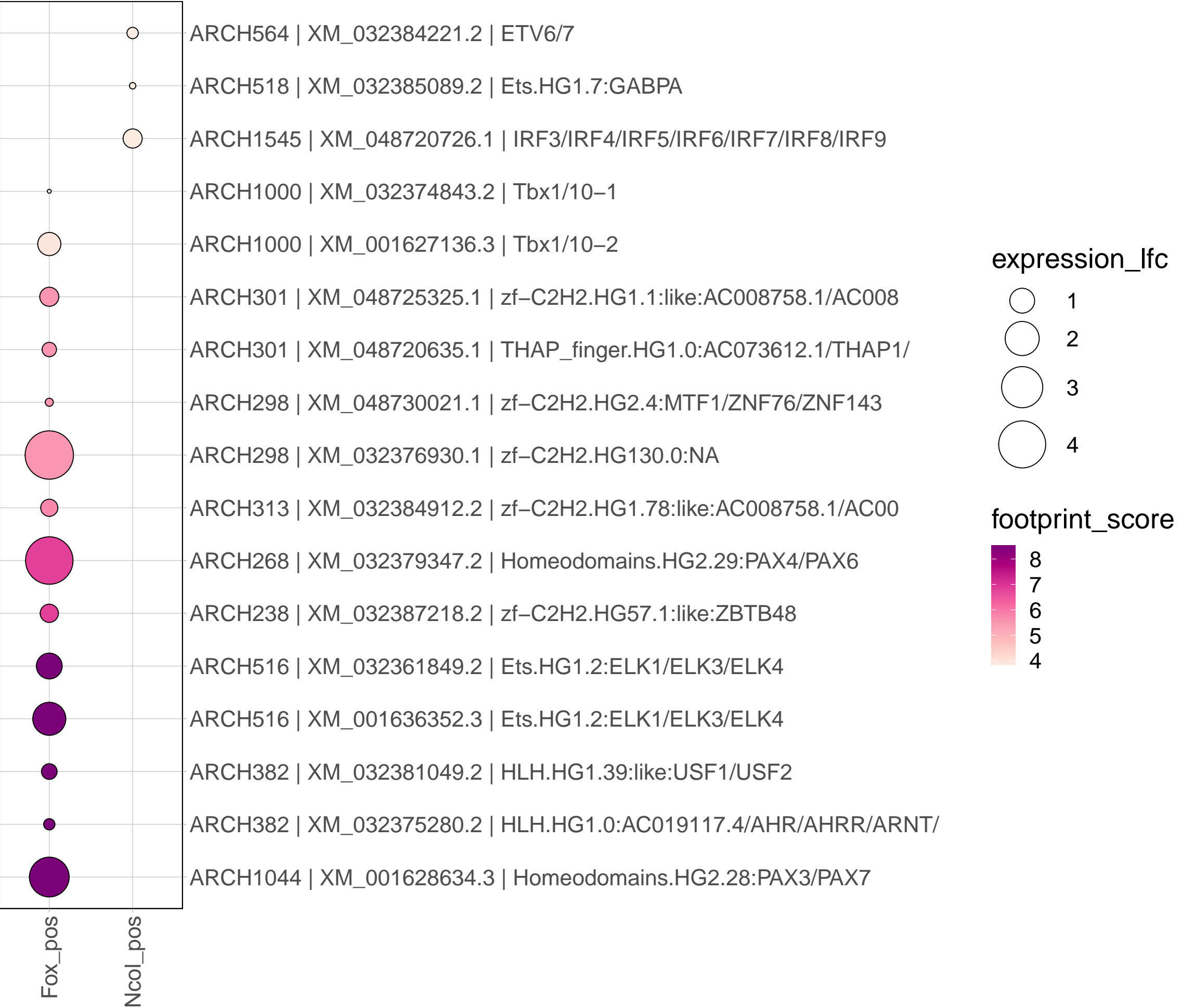
Nvec_vc1.1_XM_001633821.3



Nvec_vc1.1_XM_032379911.2
Sparc/osteonectin, cwcv and kazal-I; Thyroglobulin_1/



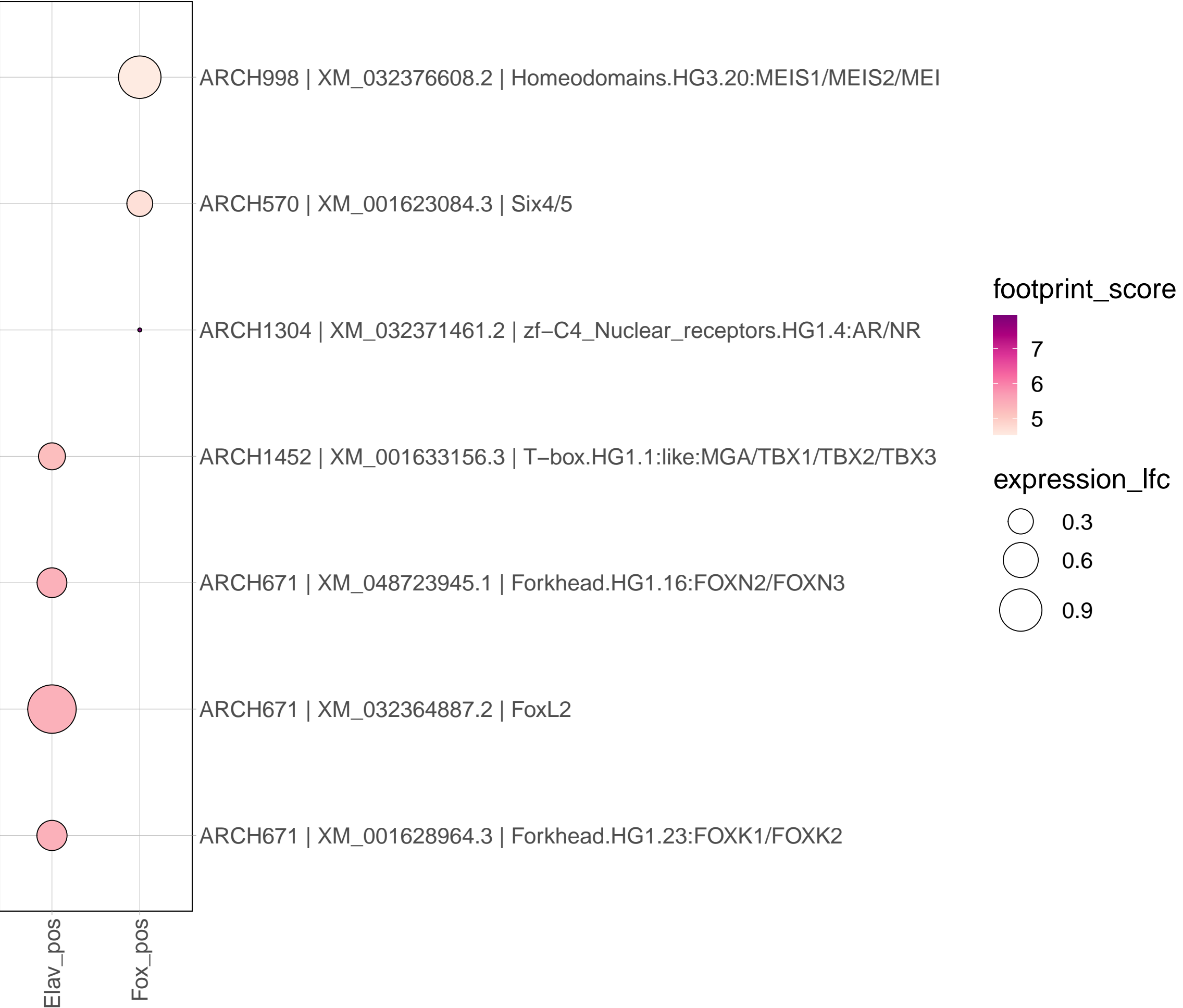
Nvec_vc1.1_XM_032370084.2



Nvec_vc1.1_XM_048723783.1
Cilia– and flagella–associated prot;

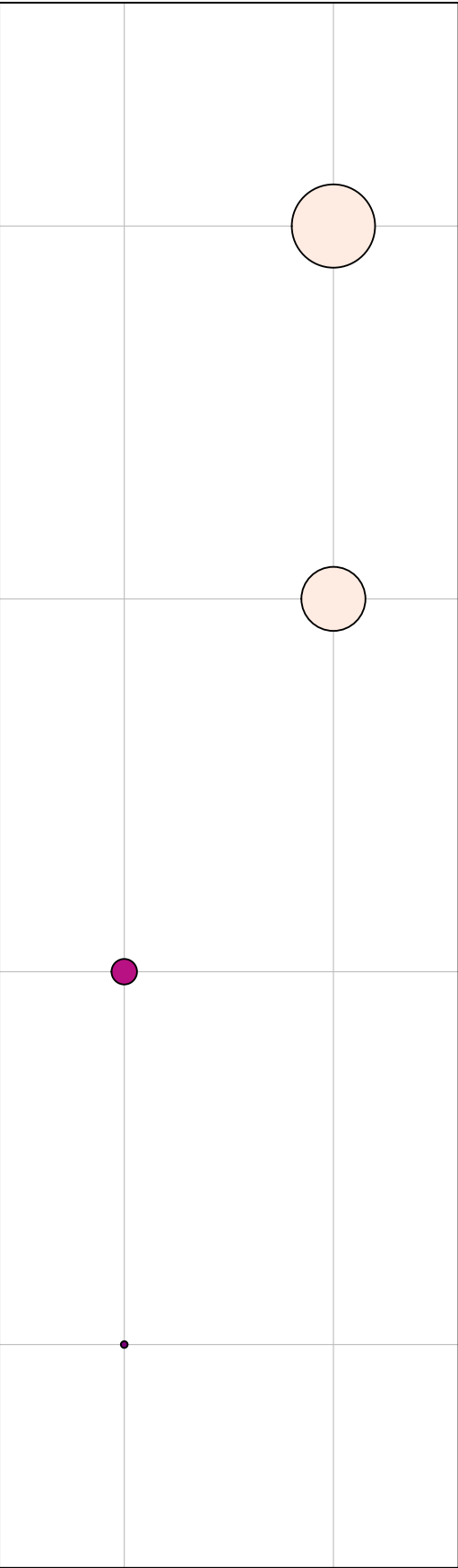


Nvec_vc1.1_XM_048725946.1
WD repeat-containing protein 49;



Nvec_vc1.1_XM_001636610.3

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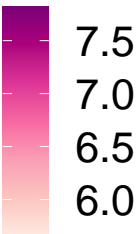
ARCH1351 | XM_032375228.2 | RFX.HG1.3:like:RFX4/RFX6/RFX8

ARCH1348 | XM_032380334.2 | RFX.HG1.6:like:RFX6

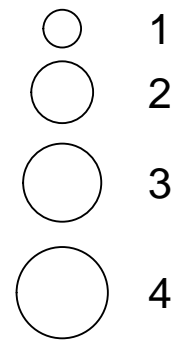
ARCH518 | XM_032385089.2 | Ets.HG1.7:GABPA

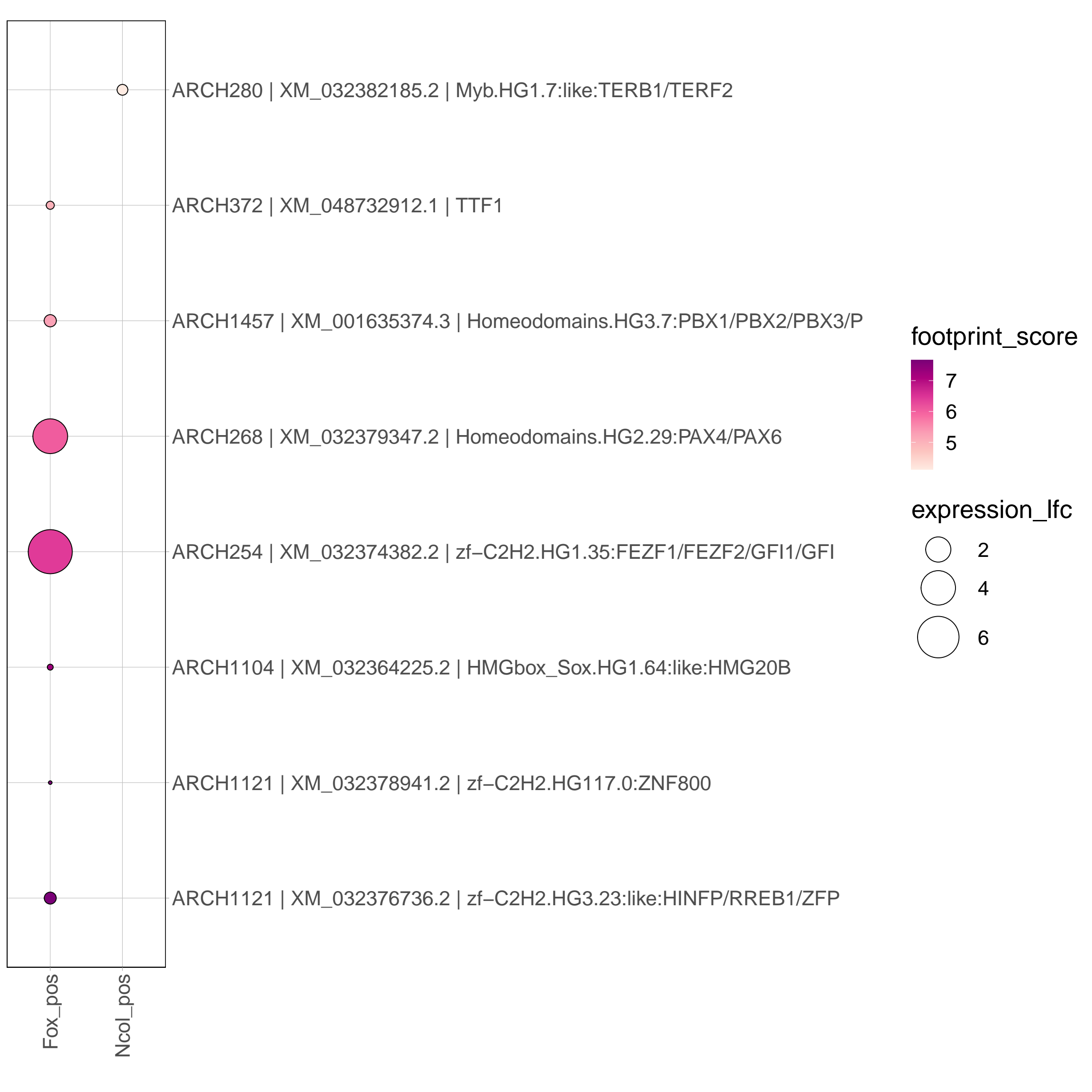
ARCH387 | XM_001625700.3 | NRF1.HG1.2:NRF1

footprint_score

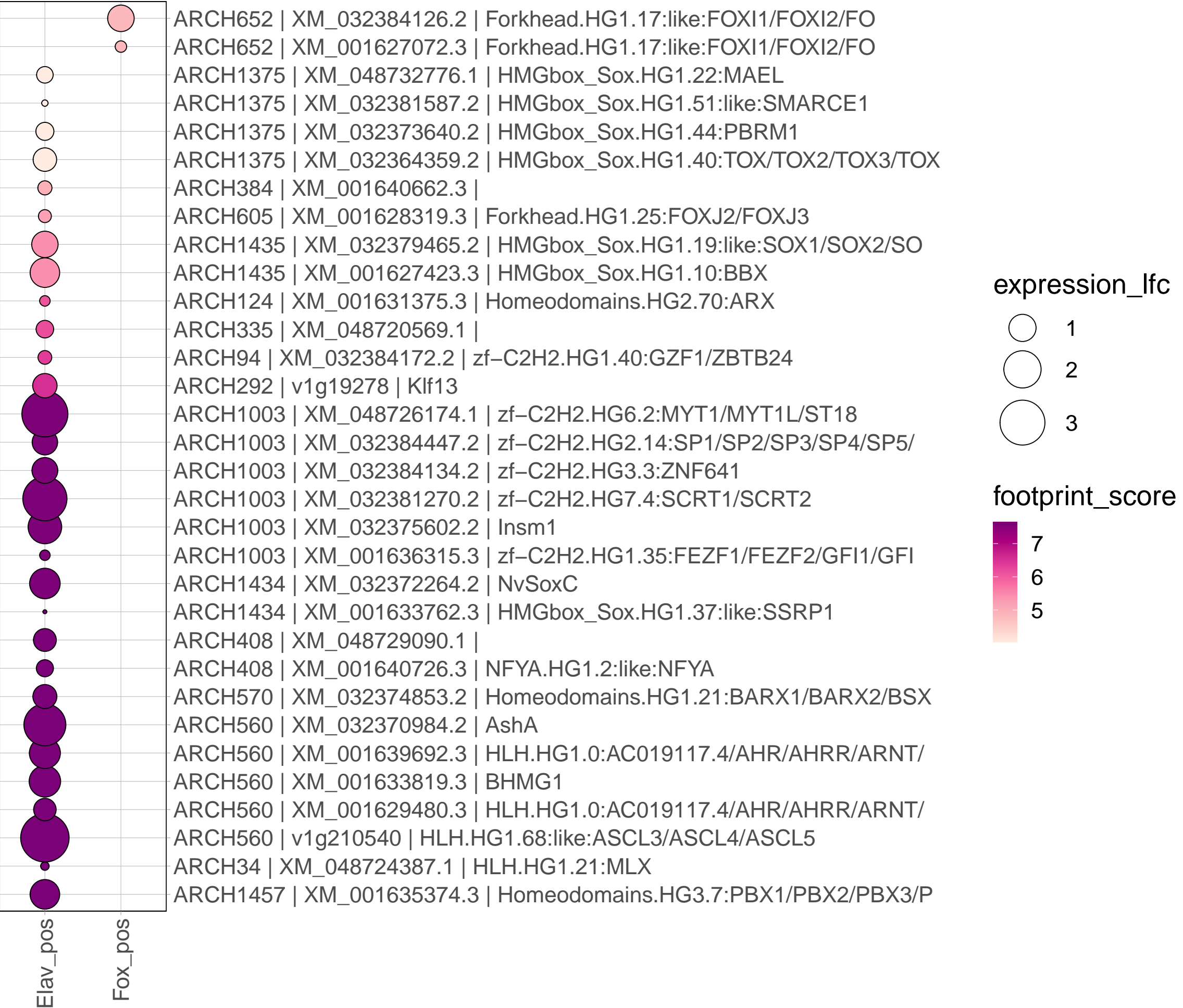


expression_lfc

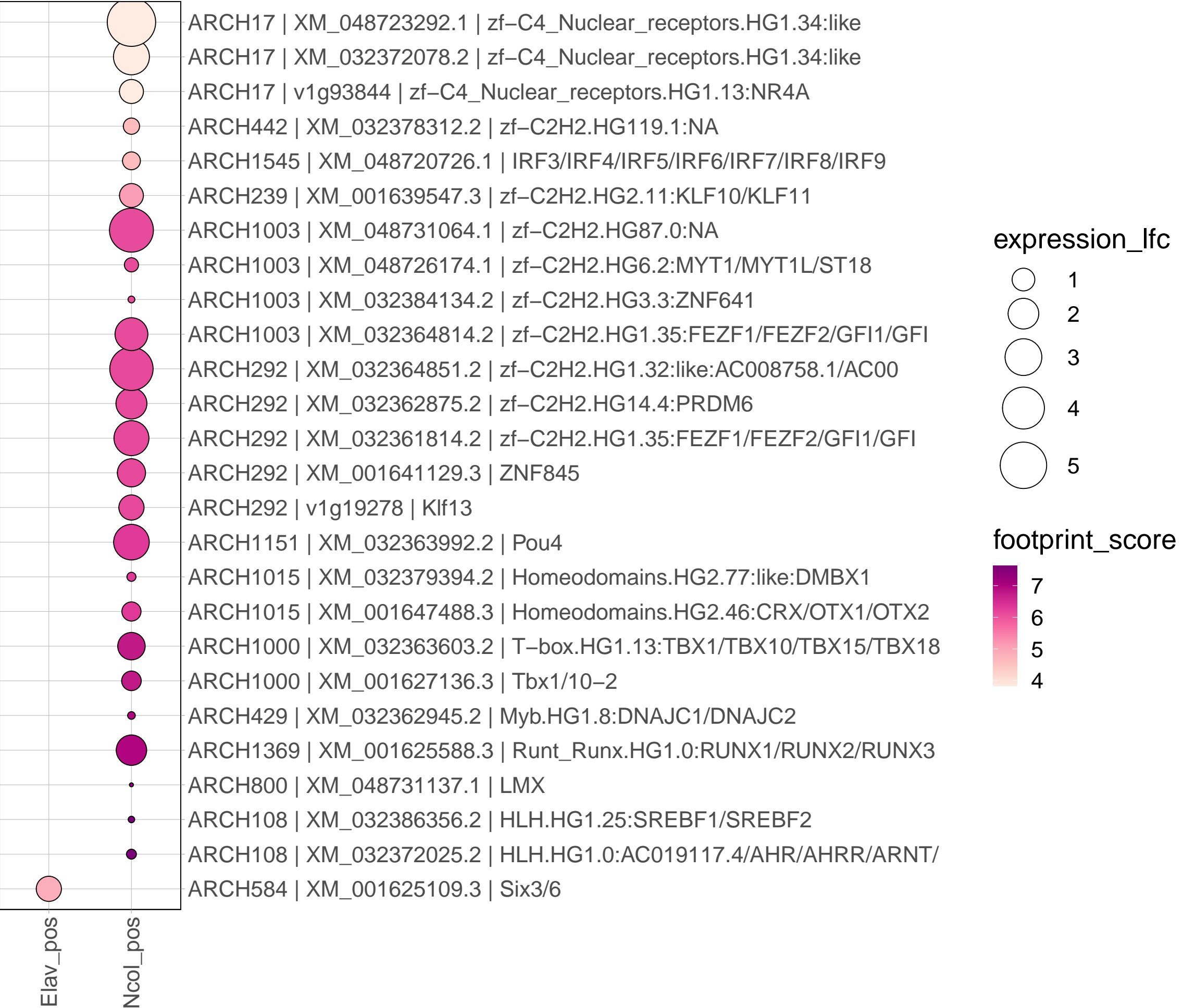




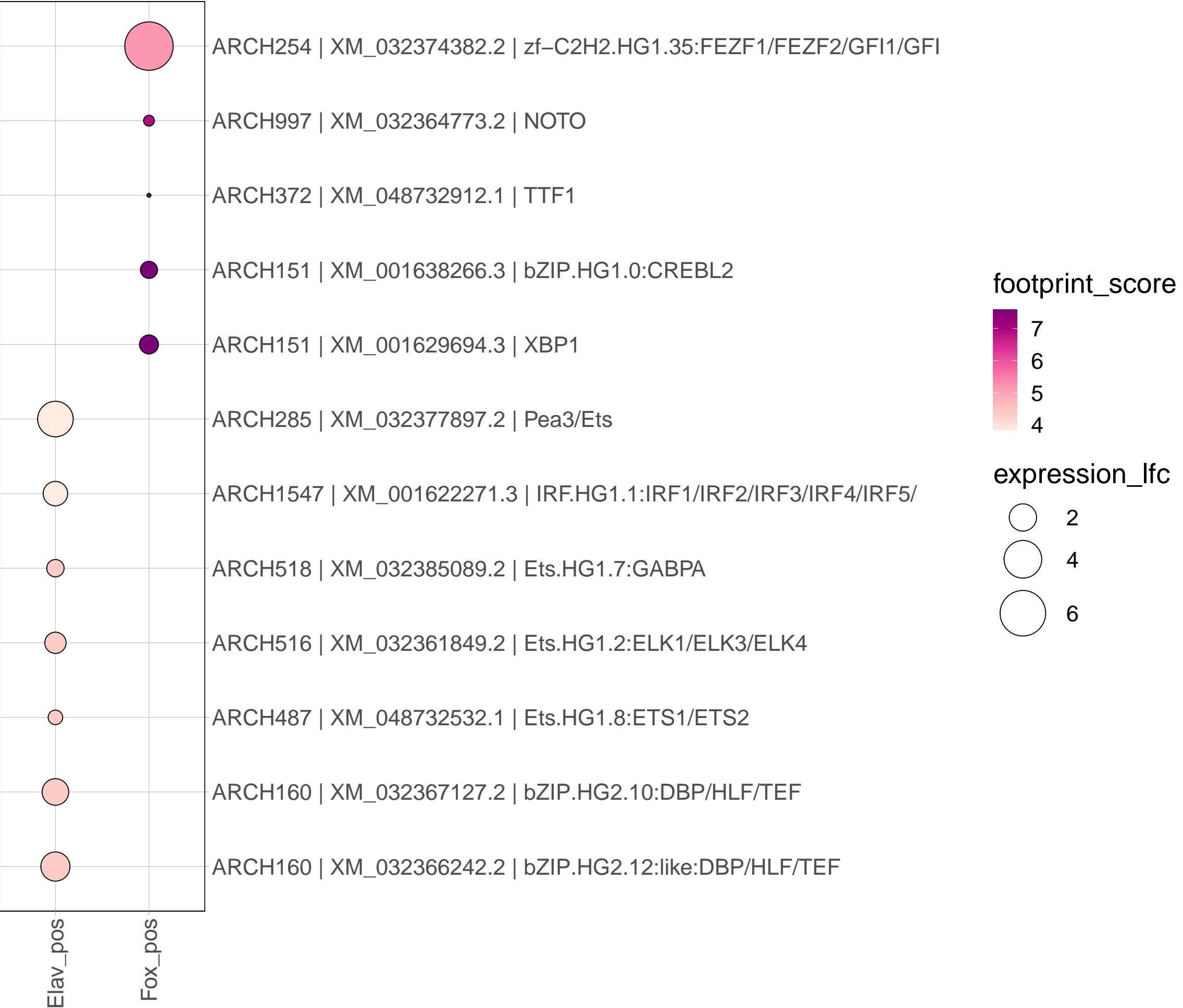
Nvec_vc1.1_XM_001628279.3
HLH.HG1.76:ATOH8; ATOH8



Nvec_vc1.1_XM_032377736.2
Protein FAM173B; Methyltransf_23/



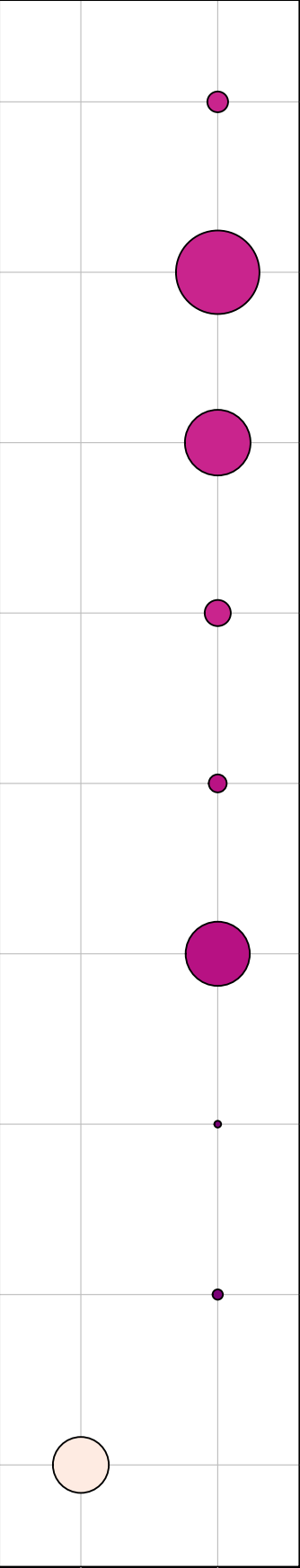
Nvec_vc1.1_XM_032361849.2
Ets.HG1.2:ELK1/ELK3/ELK4; ELK1/ELK3/ELK4



Nvec_vc1.1_XM_048731003.1
ion-NaKCa:NA; Ion_trans_2/



Nvec_vc1.1_XM_001639611.1
; EF-hand_7/



expression_lfc

1

2

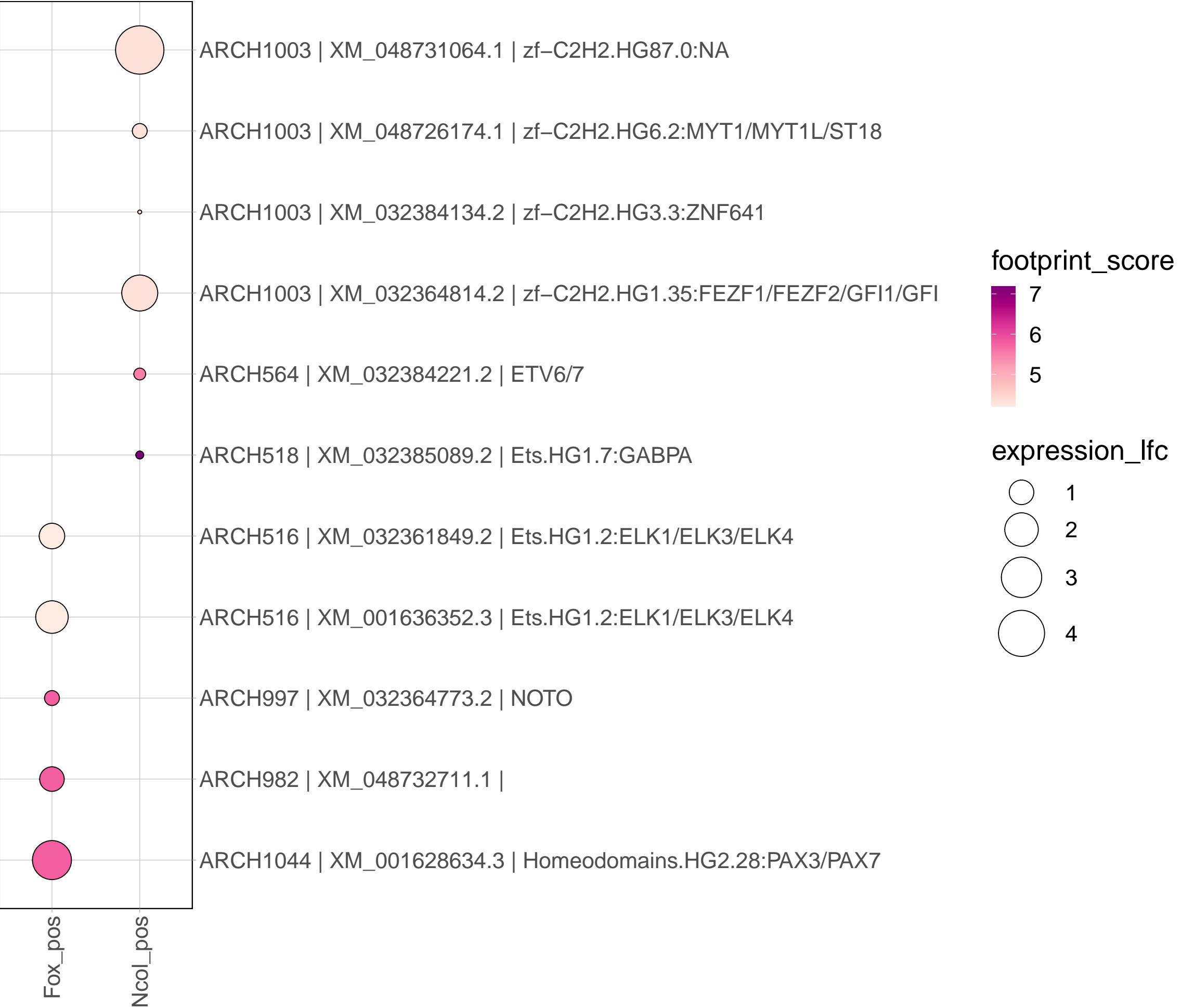
footprint_score

7

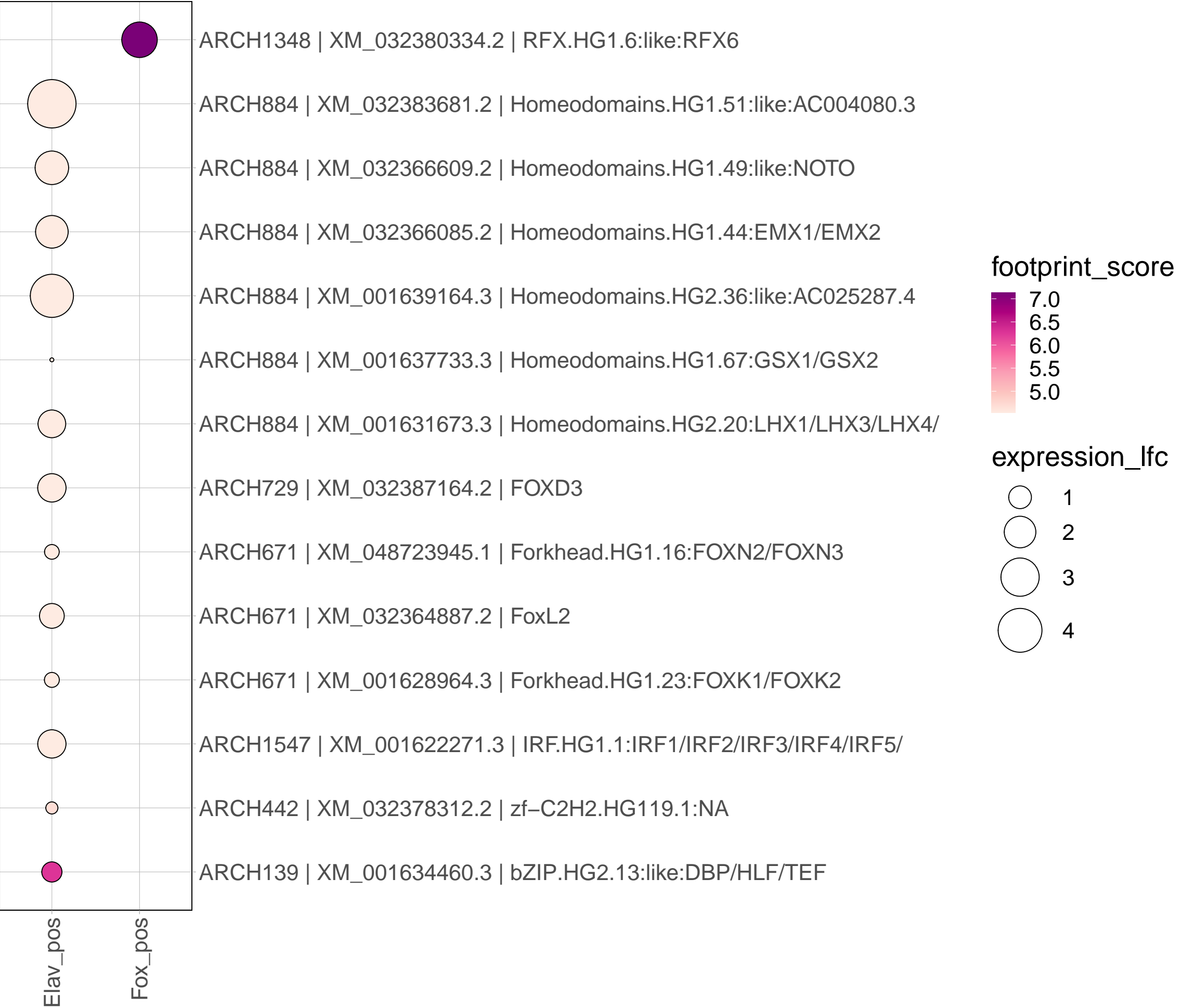
6

5

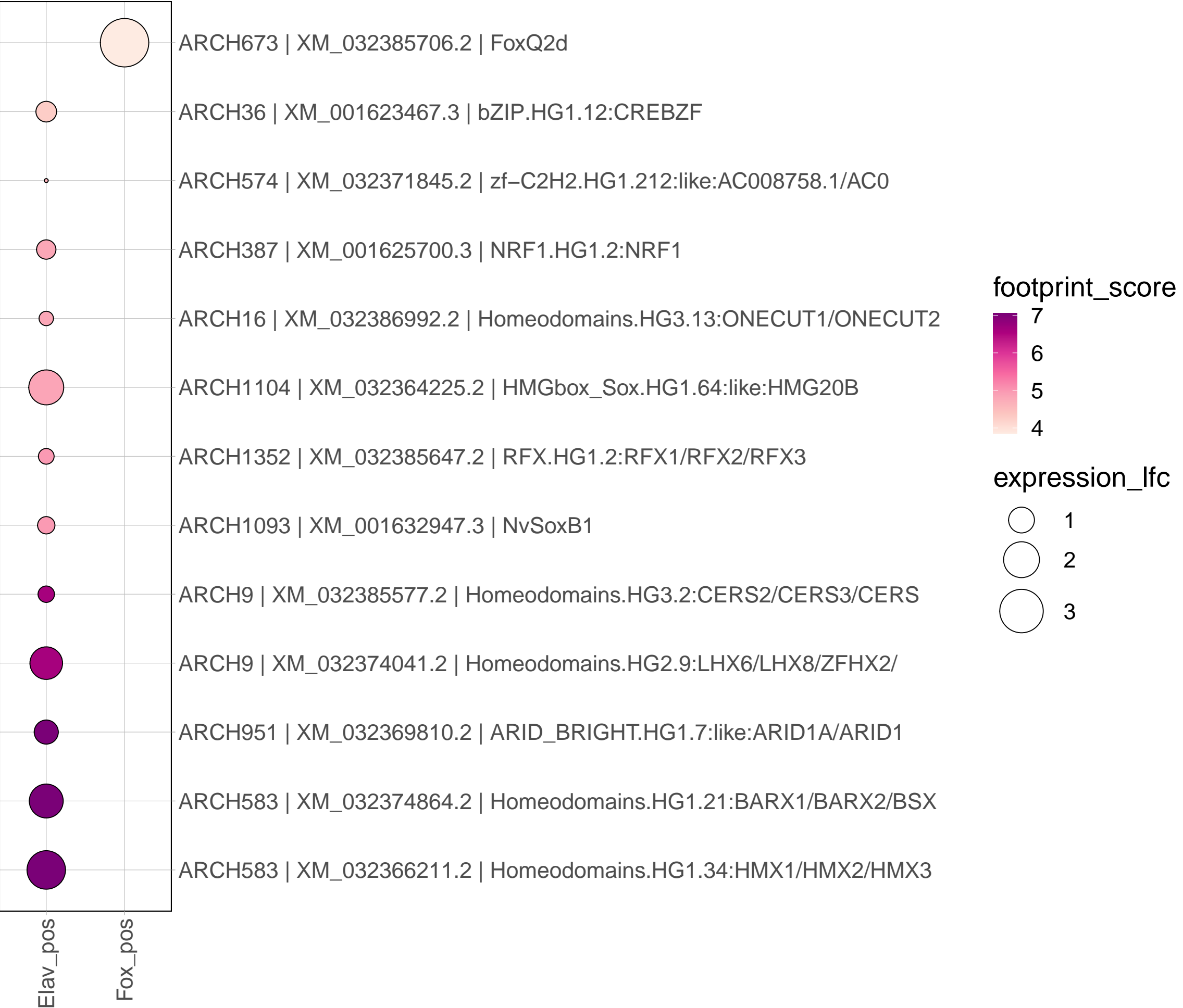
Nvec_vc1.1_XM_032362682.2
CEA protein; V-set/Ig_2/



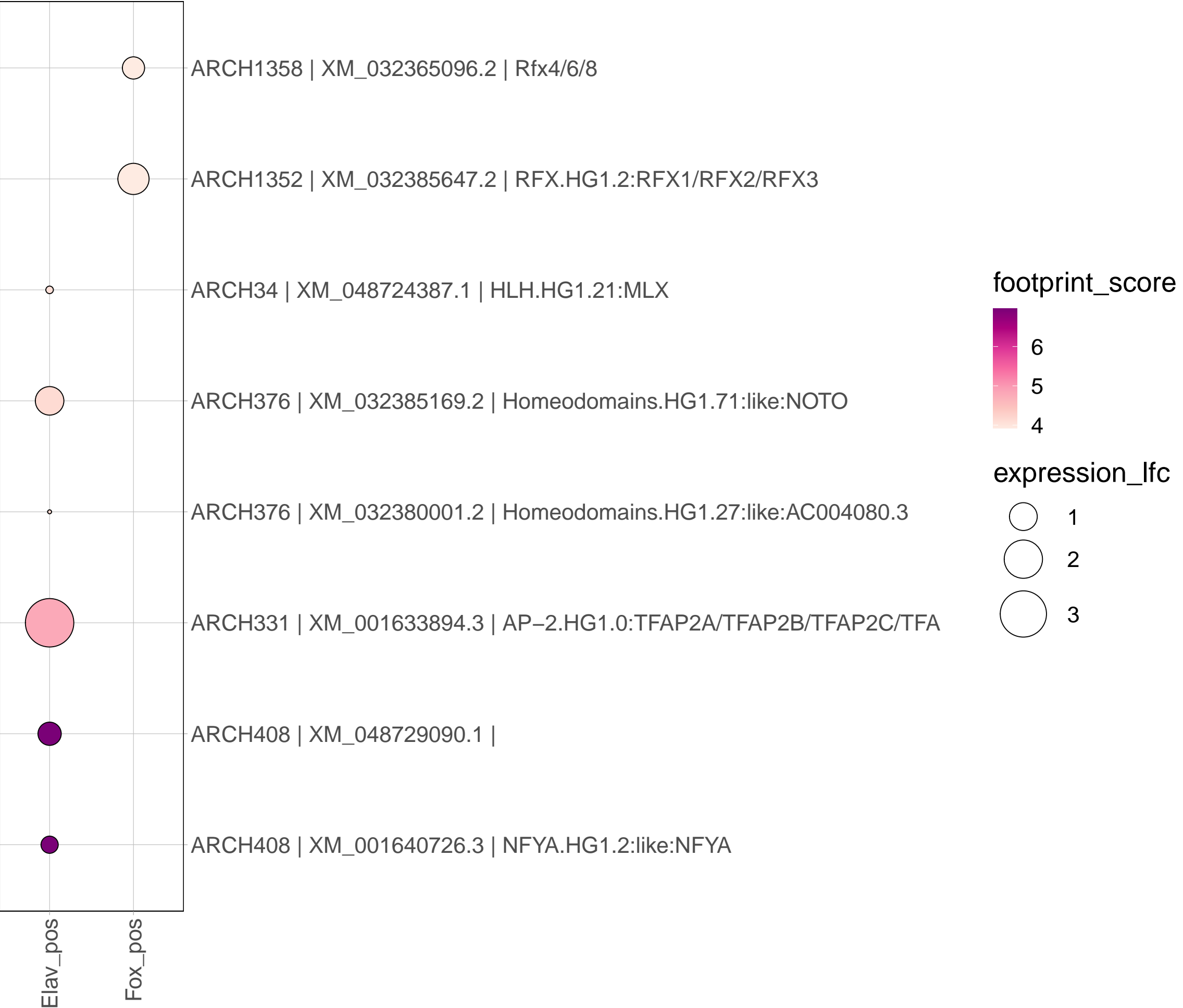
Nvec_vc1.1_XM_032380524.2
Pkinase:PIK3R4; Pkinase.HG180.0:PIK3R4



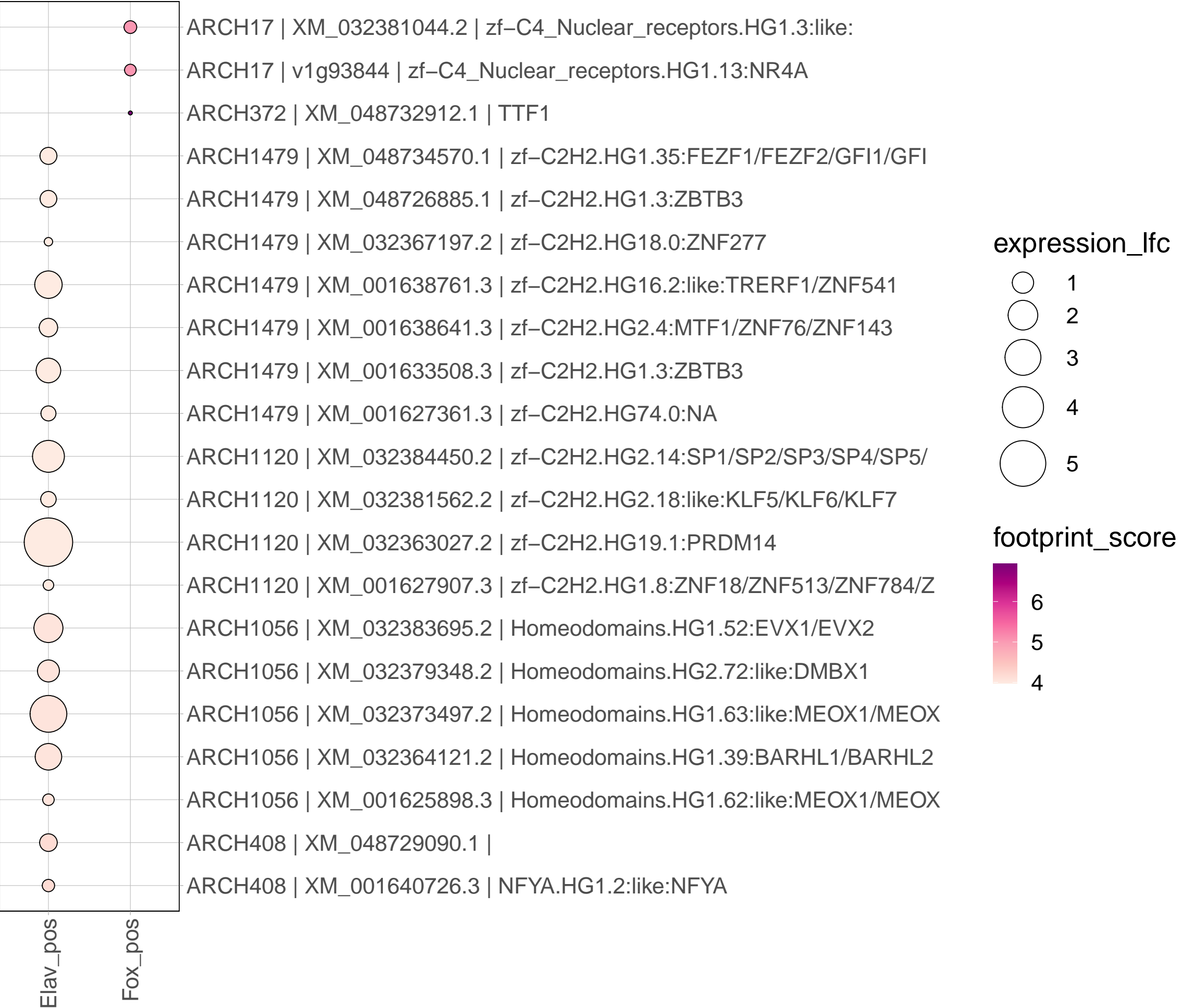
Nvec_vc1.1_XM_032378372.2
GPCR–rhodopsin:like:ADORA2B:likeclu; 7tm_1/



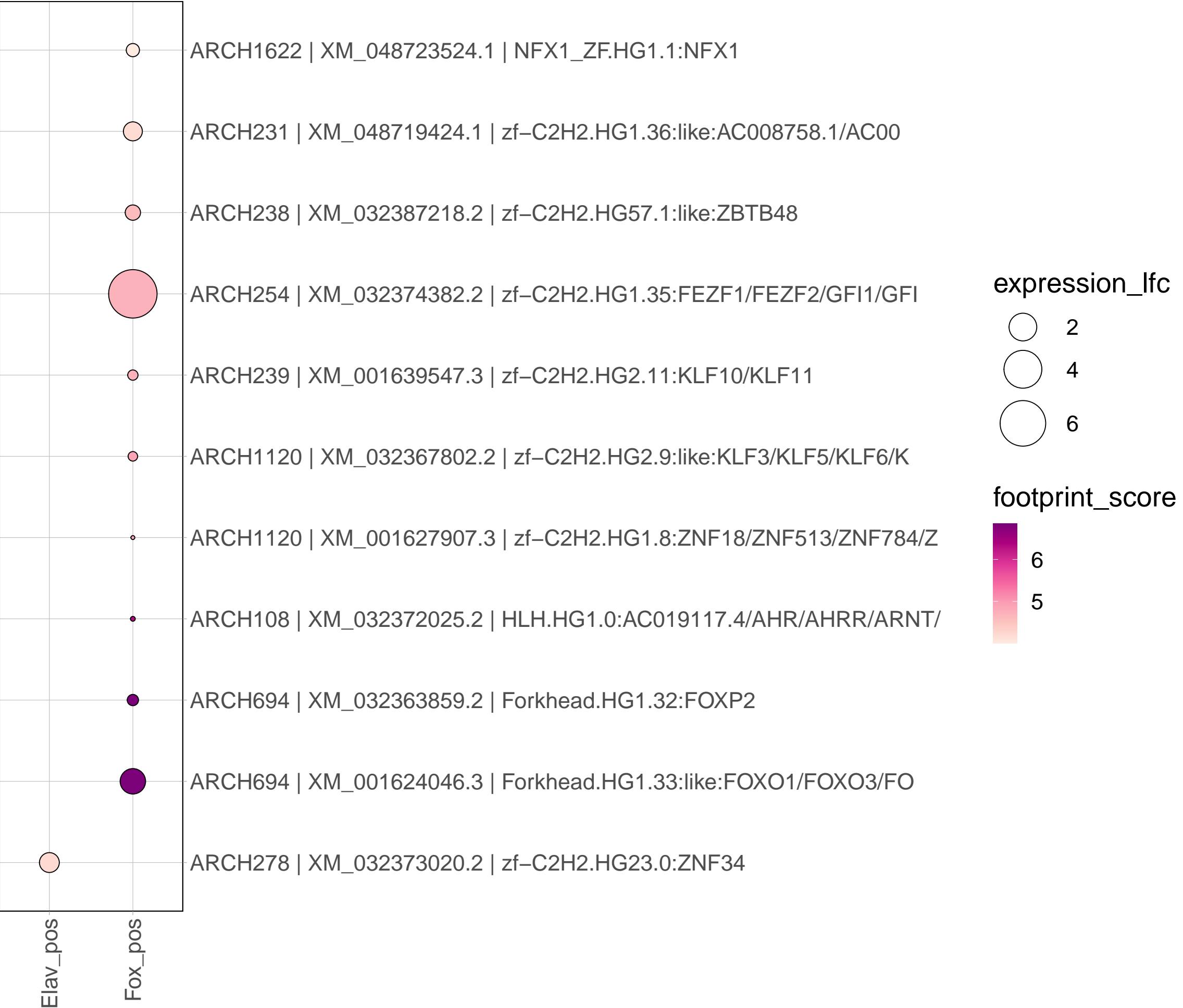
Nvec_vc1.1_XM_032386263.2
Rhombotin-1; LIM/LIM/



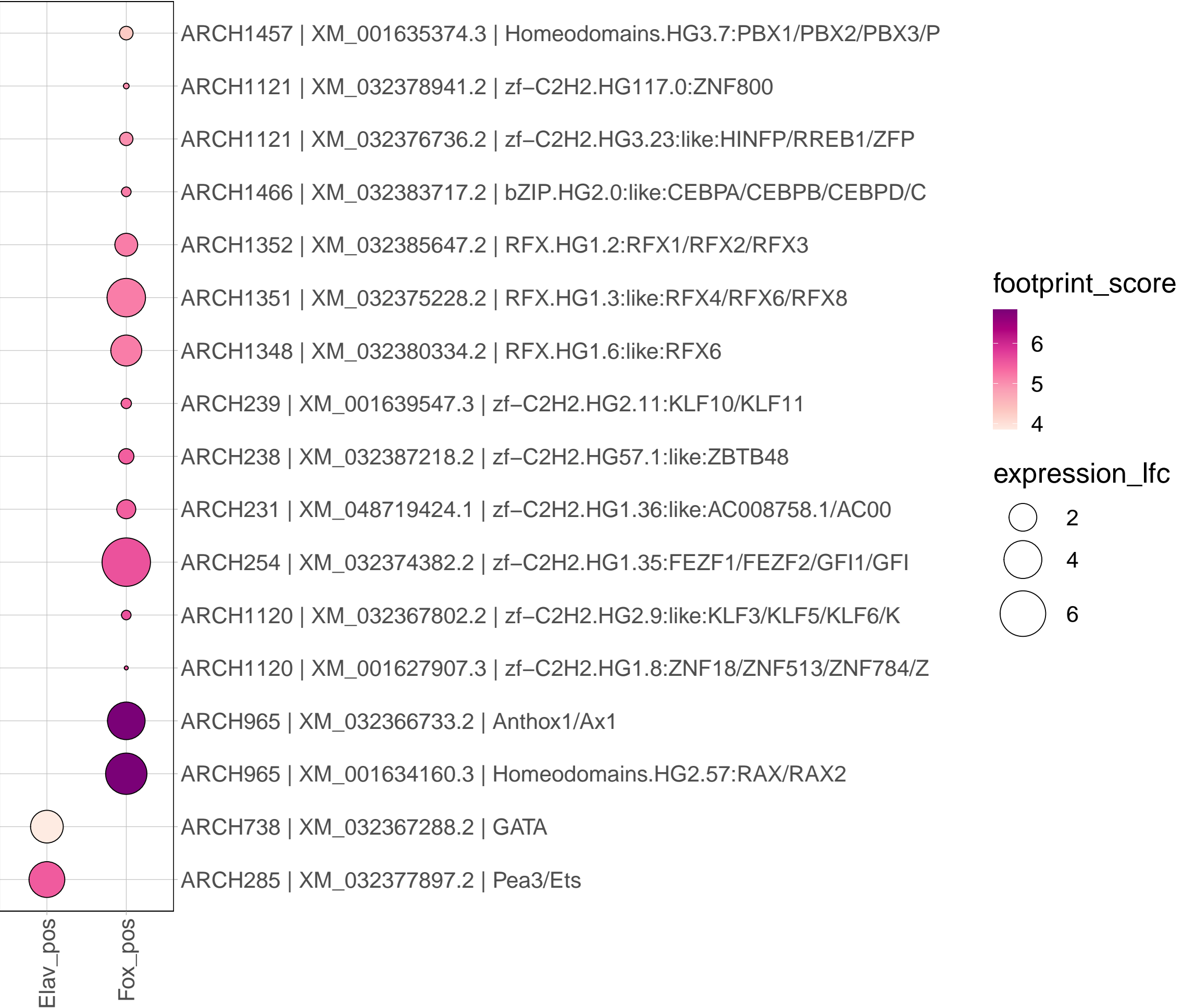
Nvec_vc1.1_XM_032378457.2
GPCR-rhodopsin:NA; 7TM_GPCR_Srw/



Nvec_vc1.1_XM_032379543.2
; DUF885/

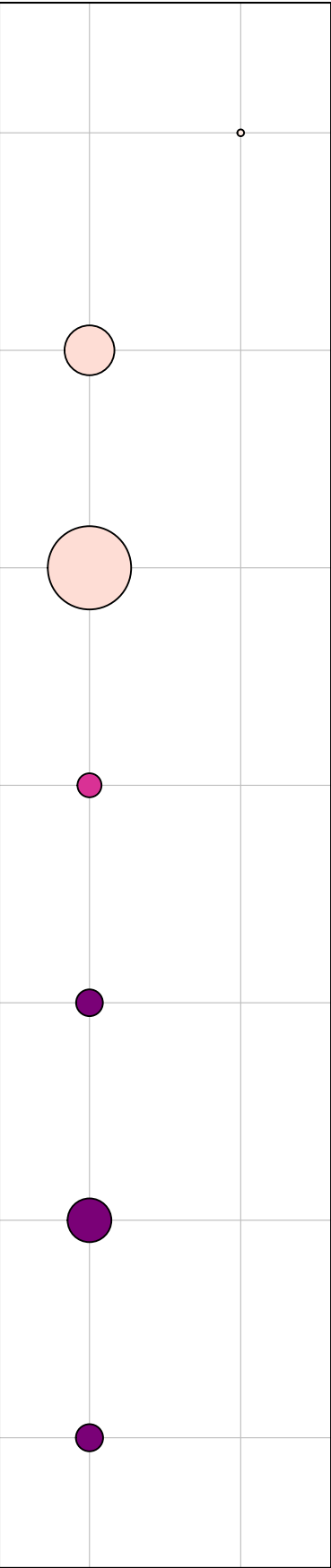


Nvec_vc1.1_XM_001633143.3
Replication factor C subunit 5; AAA/Rep_fac_C/



Nvec_vc1.1_XM_048720909.1

;



Elav_pos

Fox_pos

ARCH873 | XM_032380190.2 | Homeodomains.HG3.20:MEIS1/MEIS2/MEI

ARCH1547 | XM_001622271.3 | IRF.HG1.1:IRF1/IRF2/IRF3/IRF4/IRF5/

ARCH1272 | XM_032383641.2 | zf-C4_Nuclear_receptors.HG1.9:like:

ARCH1452 | XM_001633156.3 | T-box.HG1.1:like:MGA/TBX1/TBX2/TBX3

ARCH671 | XM_048723945.1 | Forkhead.HG1.16:FOXN2/FOXN3

ARCH671 | XM_032364887.2 | FoxL2

ARCH671 | XM_001628964.3 | Forkhead.HG1.23:FOXK1/FOXK2

expression_lfc



1



2



3



4

footprint_score



6.5

6.0

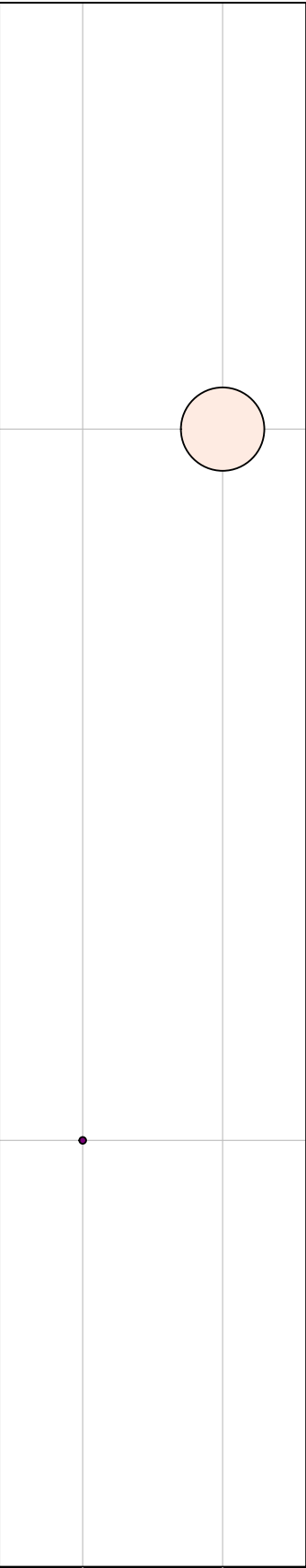
5.5

5.0

4.5

Nvec_vc1.1_XM_048729009.1

;



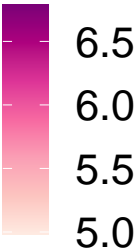
ARCH731 | XM_032386243.2 | Grainyhead

ARCH993 | XM_001634845.3 | zf-CXXC.HG1.8:CXXC4/CXXC5/TET1/TET3

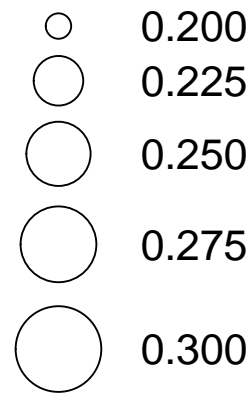
Fox_pos

Ncol_pos

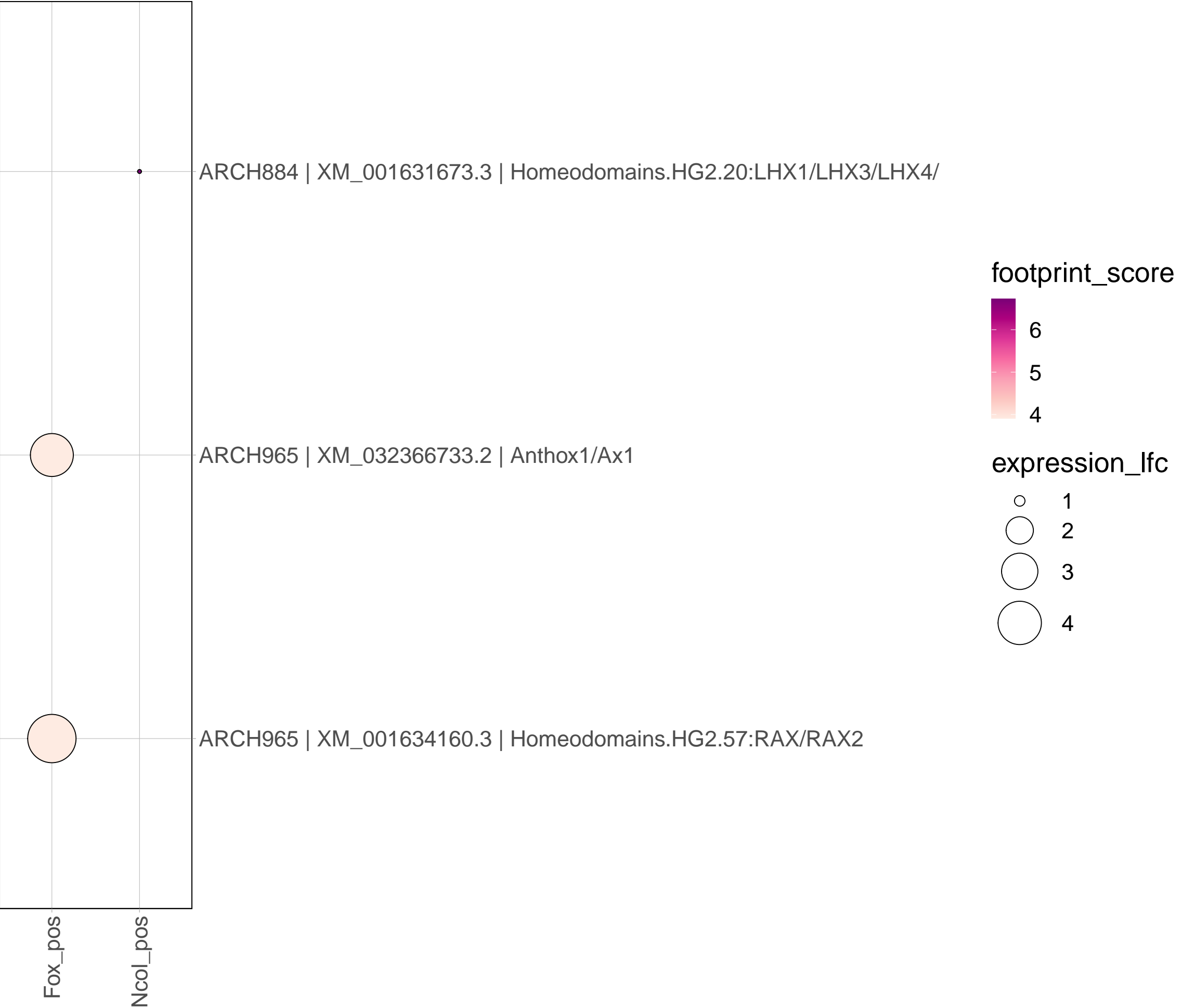
footprint_score



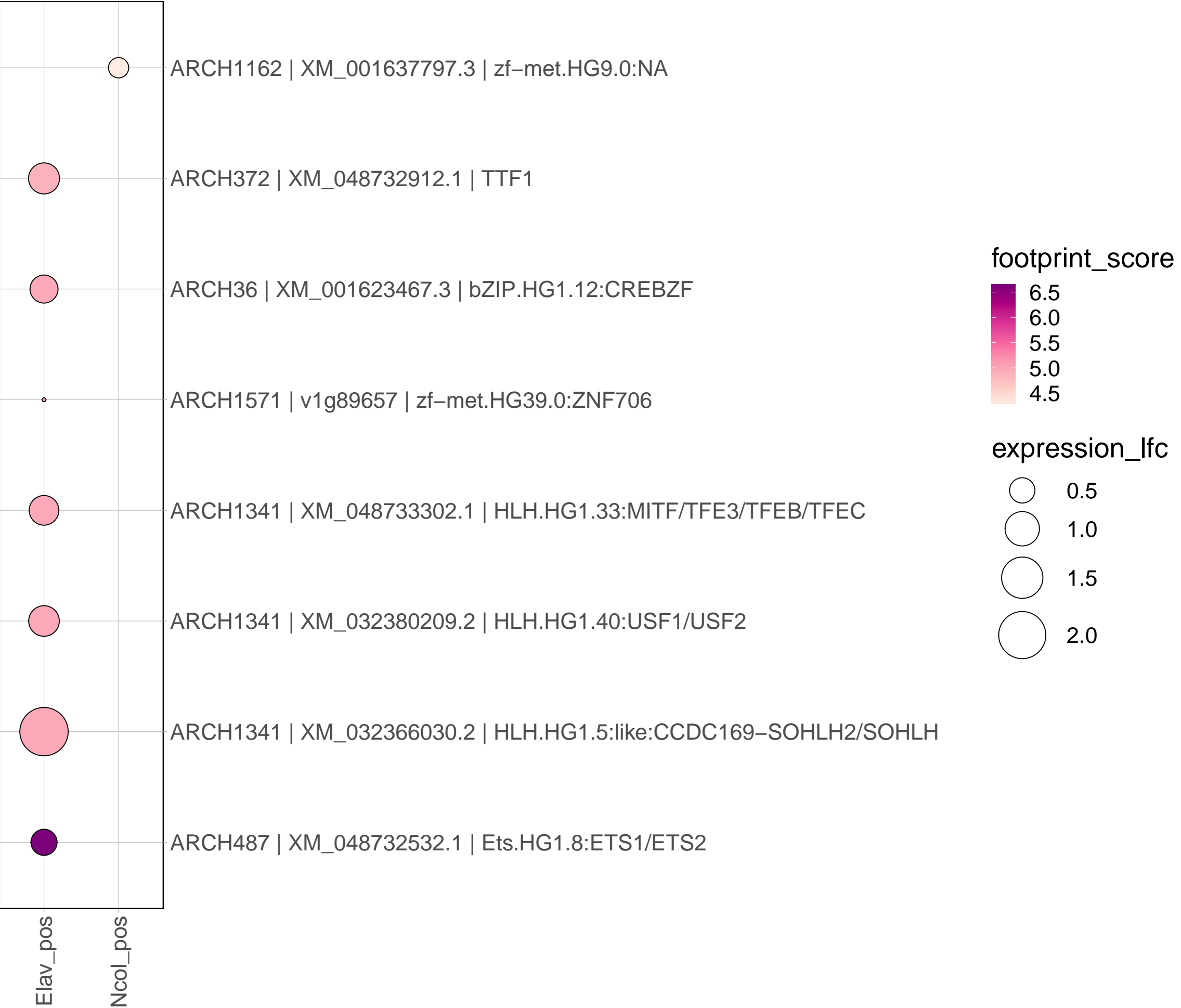
expression_lfc



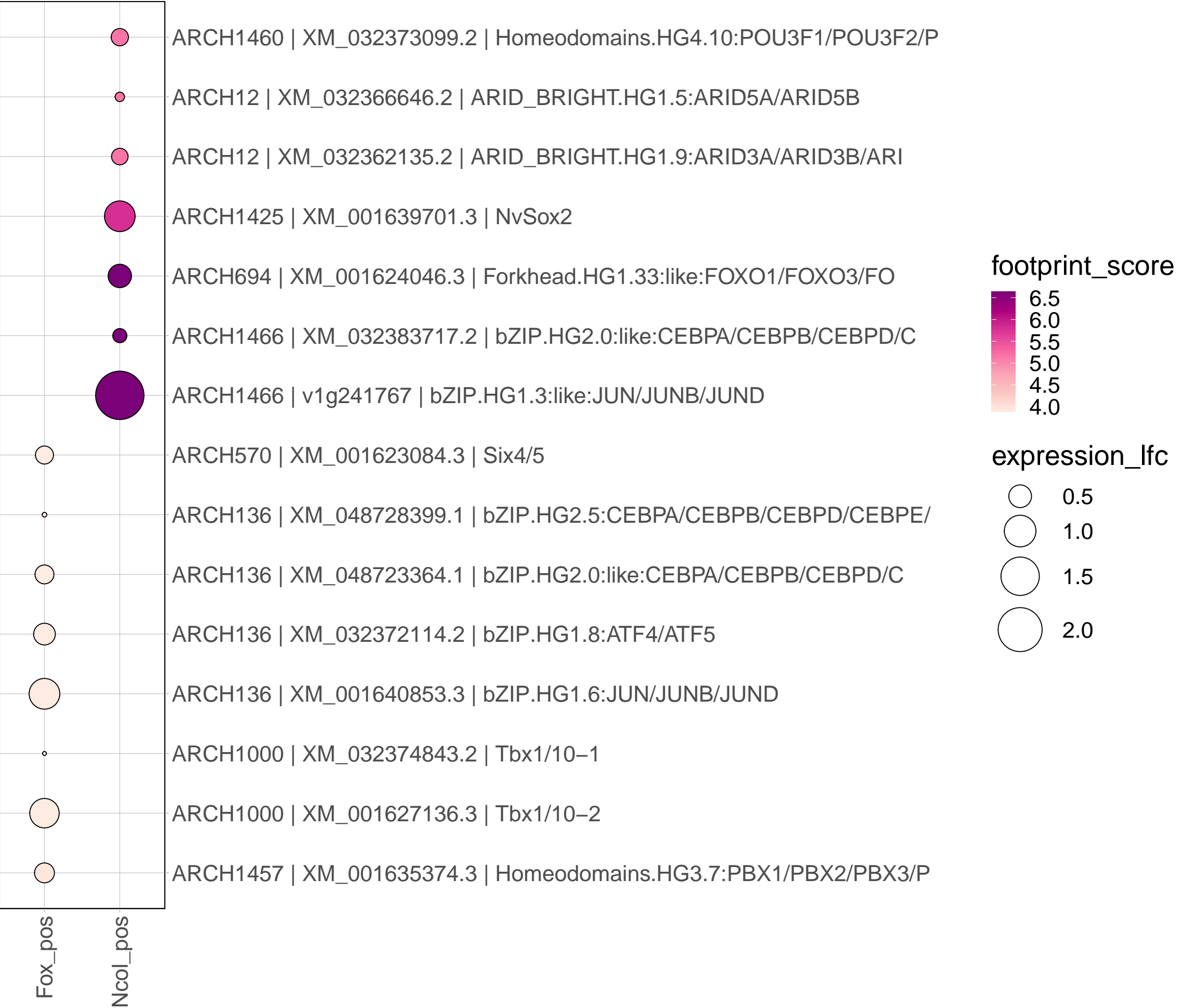
Nvec_vc1.1_XM_001623762.3
Vesicular glutamate transporter 3; MFS_1/



Nvec_vc1.1_XM_001627181.3
Myeloid leukemia factor 1; Mlf1IP/



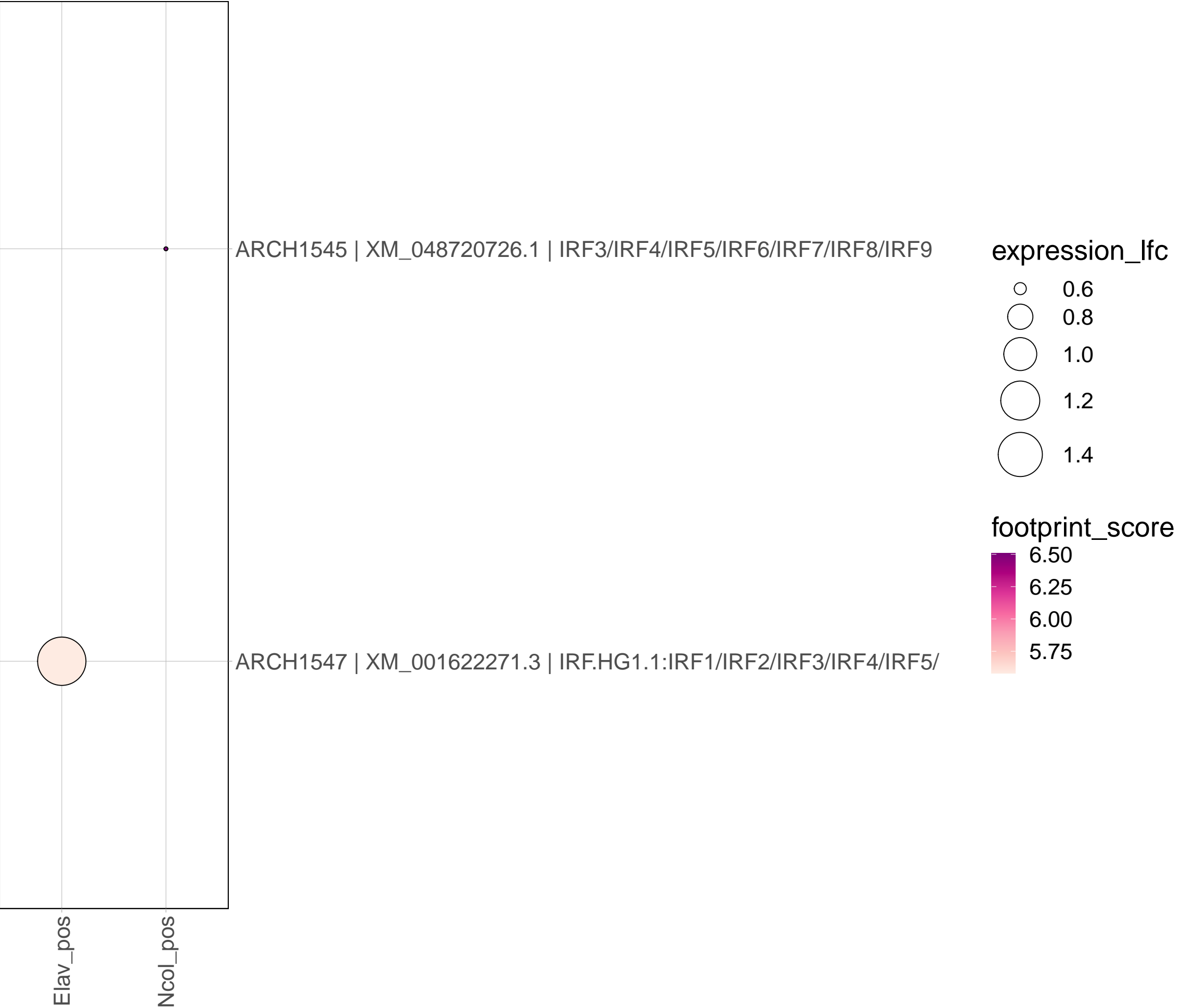
Nvec_vc1.1_XM_032372769.2
fibronectin:like:AXL/MERTK/NPHS1/RO; I-set/Ig_2/fn3/fn3/



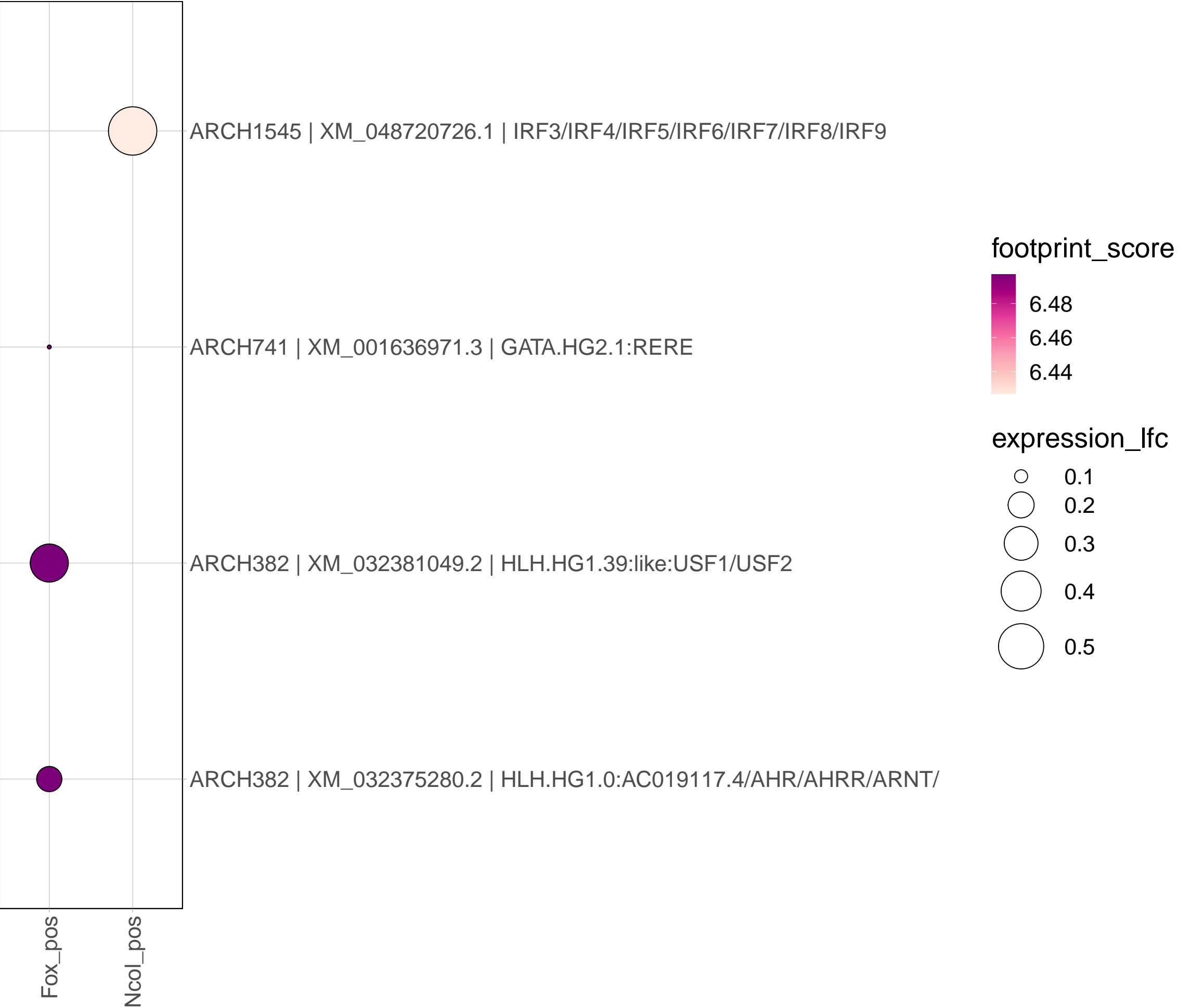
Nvec_vc1.1_XM_048734459.1
E3 ubiquitin–protein ligase RNF19A;



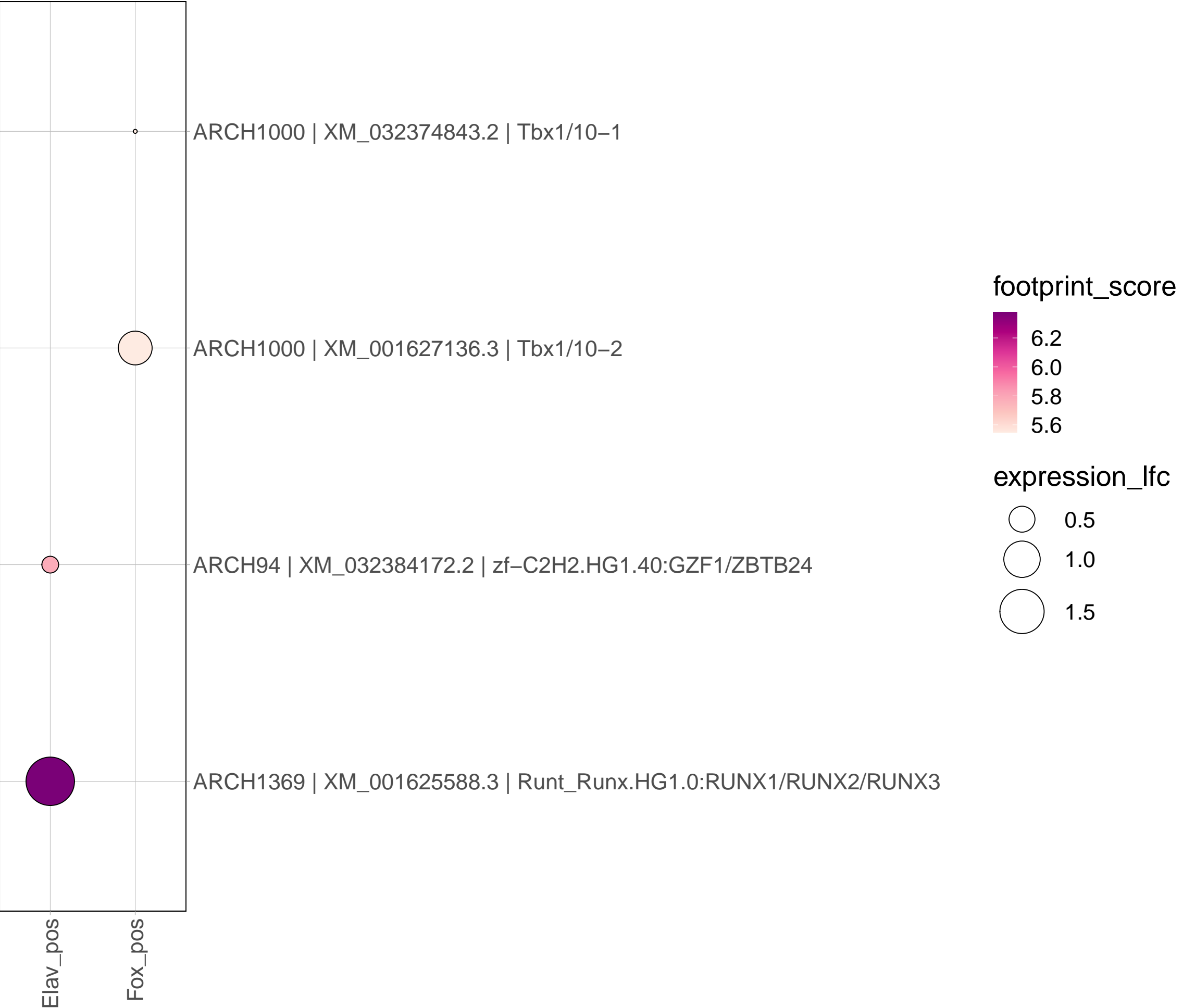
Nvec_vc1.1_XM_001636661.3
Serine/threonine–protein phosphatas; Metallophos/



Nvec_vc1.1_XM_001639688.3
Peroxisomal acyl-coenzyme A oxidase; Acyl-CoA_dh_M/Acyl-CoA_dh_1/ACOX

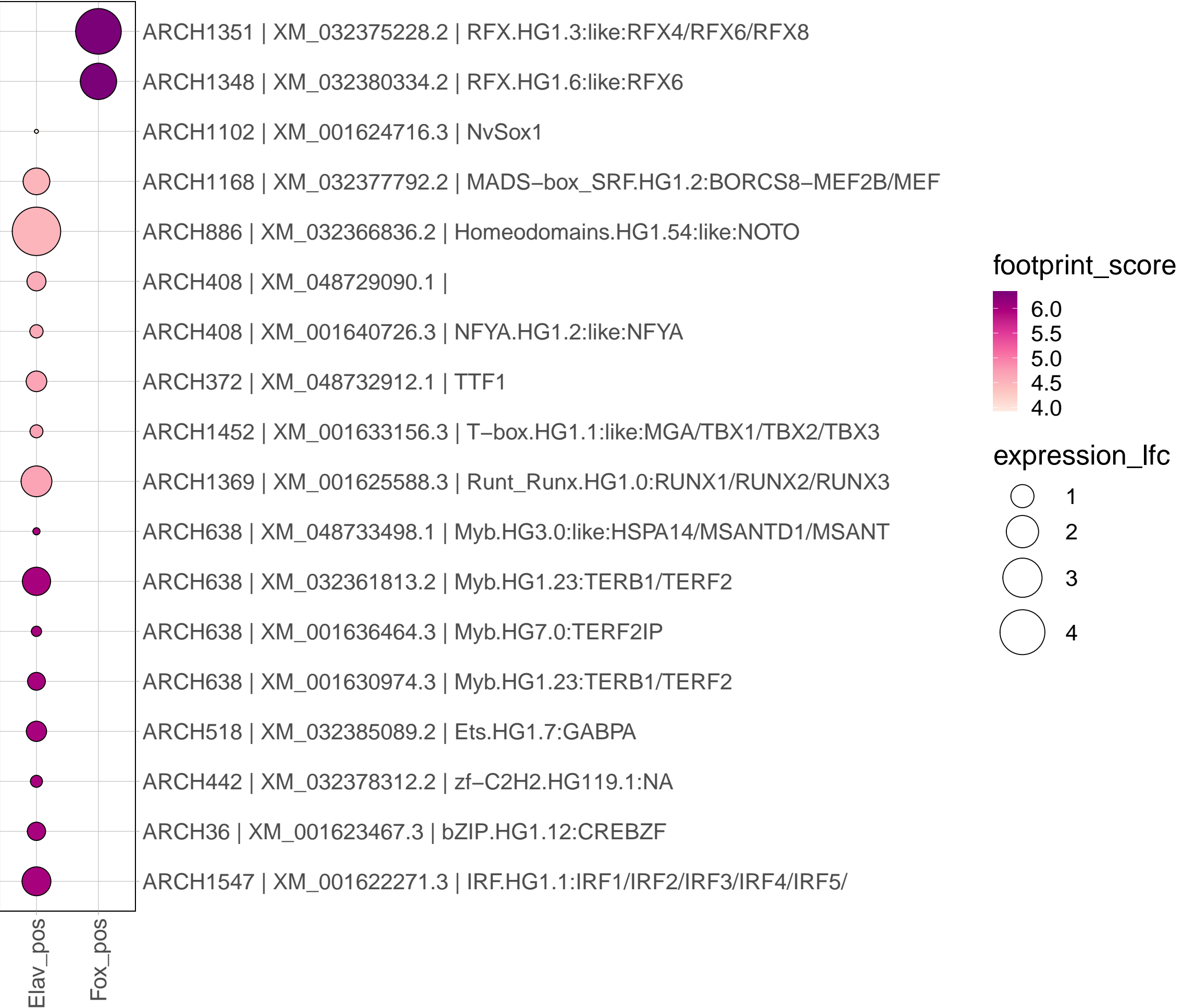


Nvec_vc1.1_XM_032374307.2
Major facilitator superfamily domai; MFS_2/

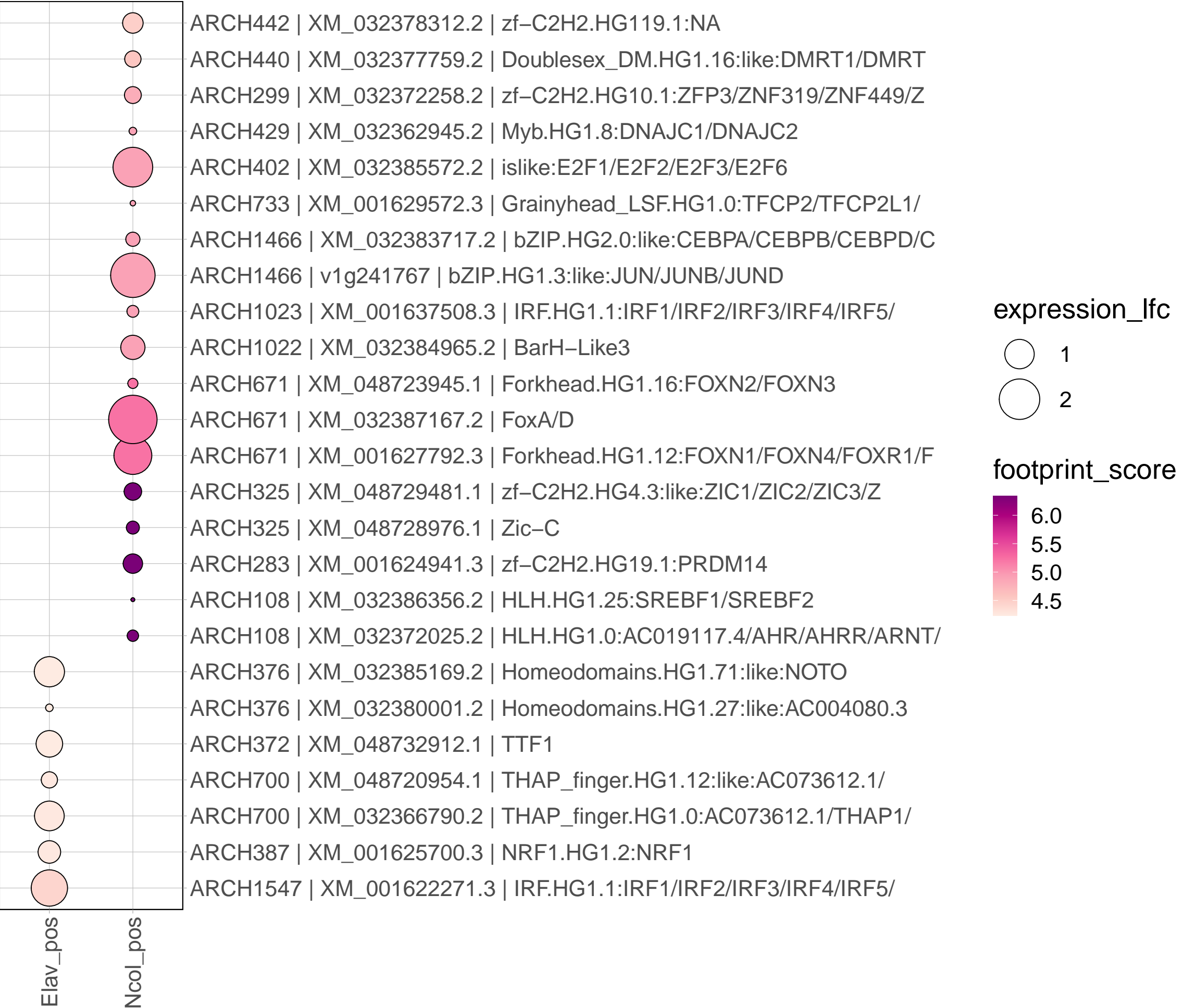


Nvec_vc1.1_XM_032365655.2

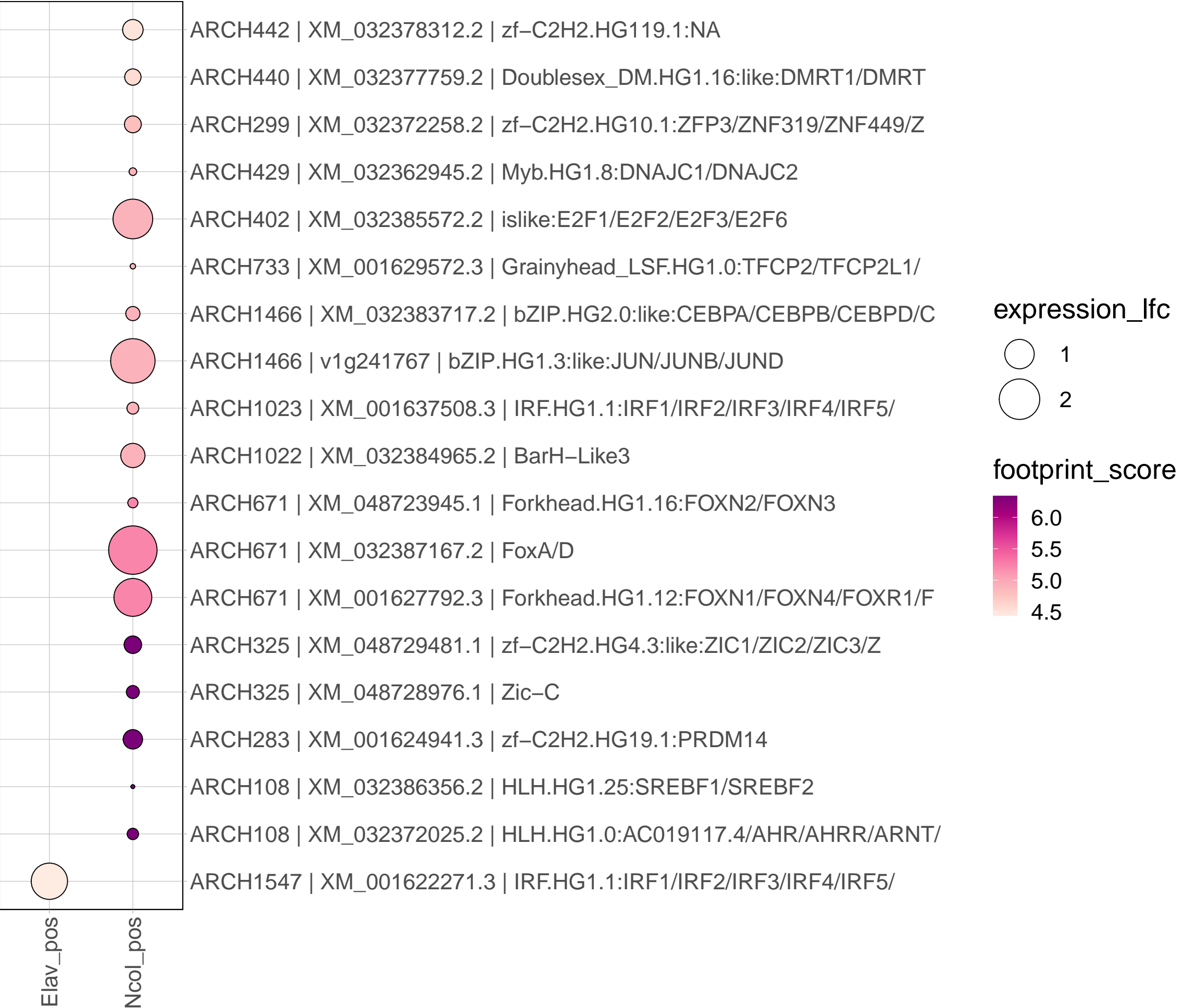
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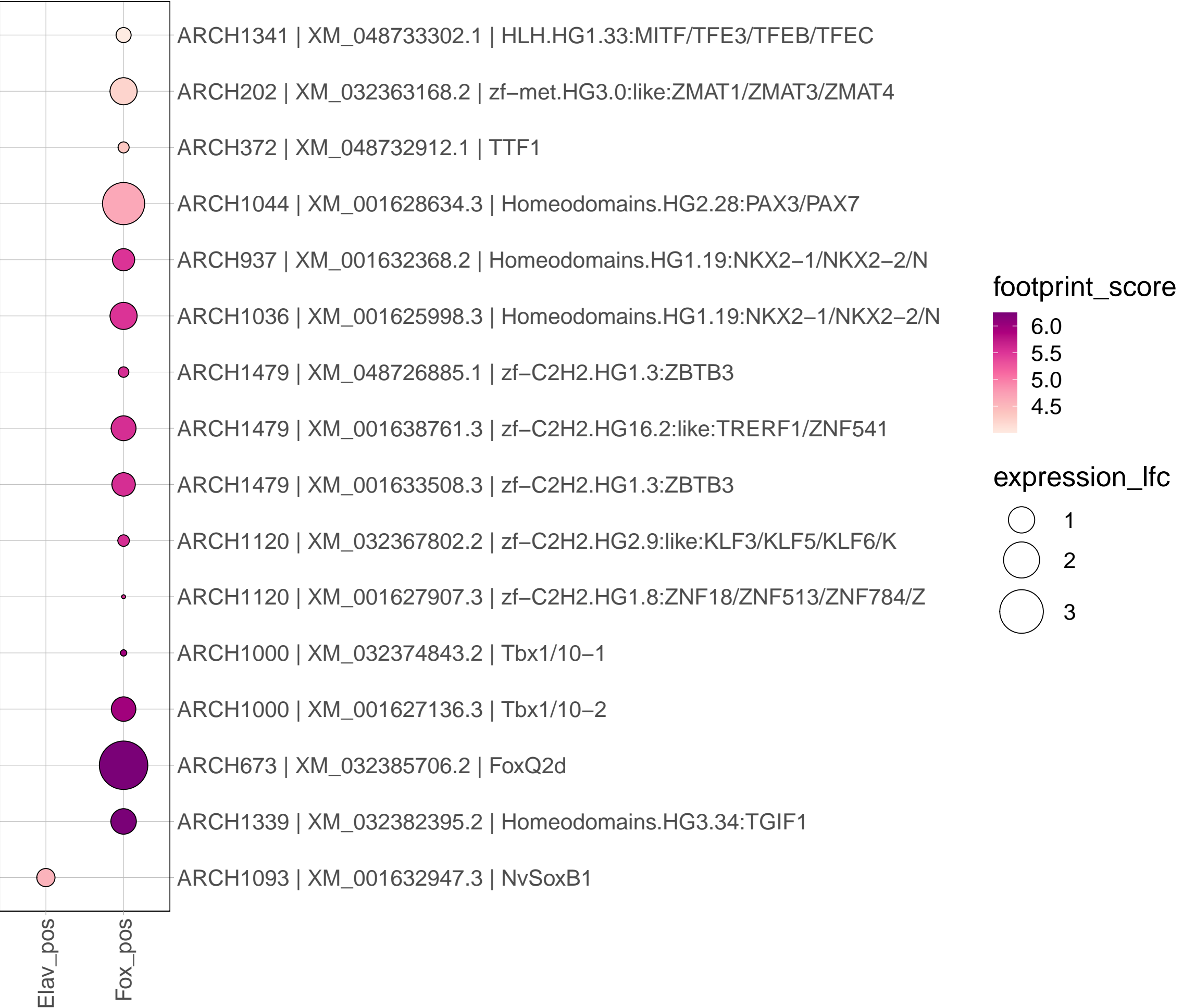
Nvec_vc1.1_XM_048720751.1
cDNA FLJ33589 fis, clone BRAMY20125; Asparaginase_2/



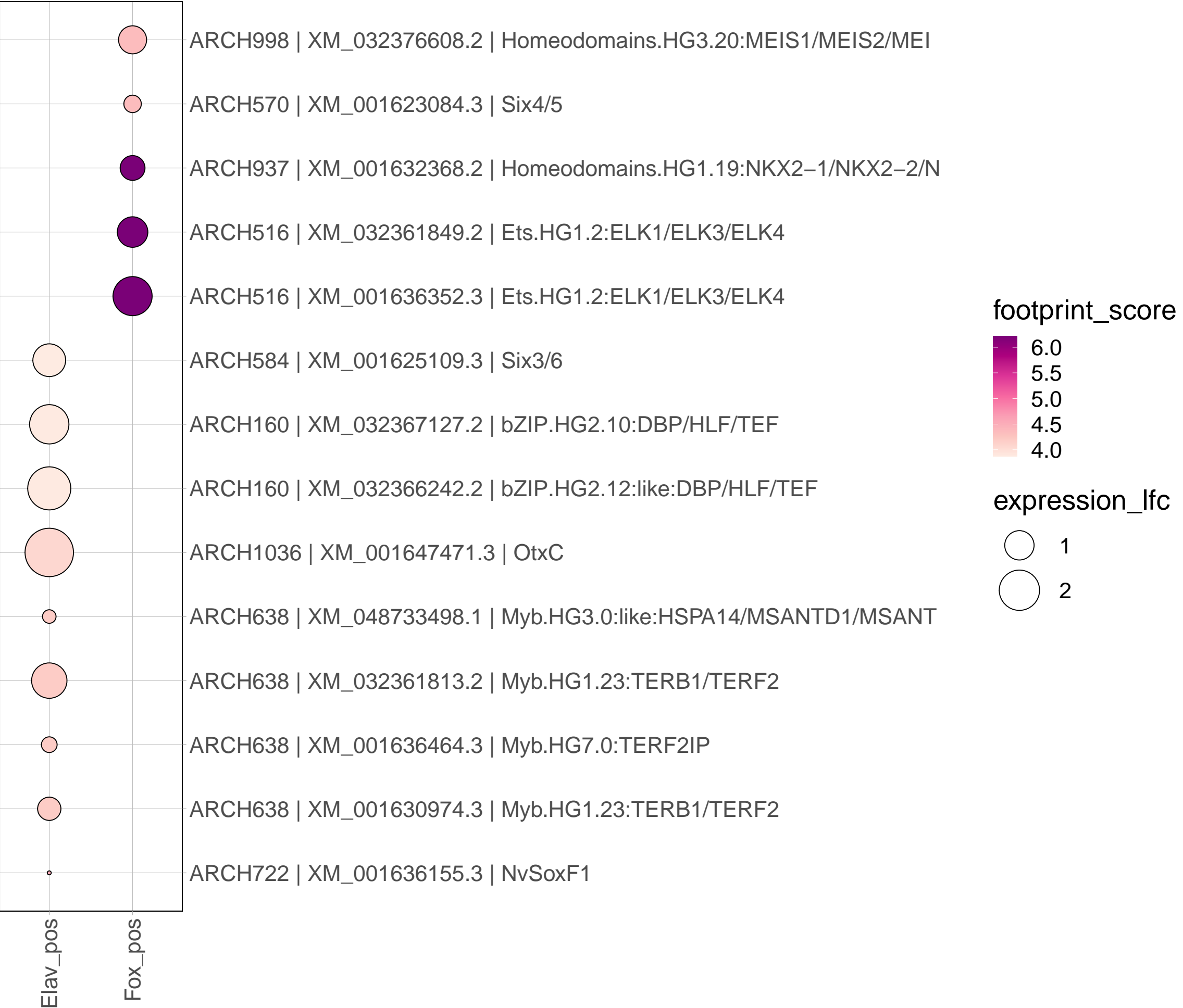
Nvec_vc1.1_XM_048720757.1
cDNA FLJ33589 fis, clone BRAMY20125; Asparaginase_2/Asparaginase_2/



Nvec_vc1.1_XM_032362262.2
GPCR–rhodopsin:like:CCKAR/CCKBR/GPR; 7tm_1/

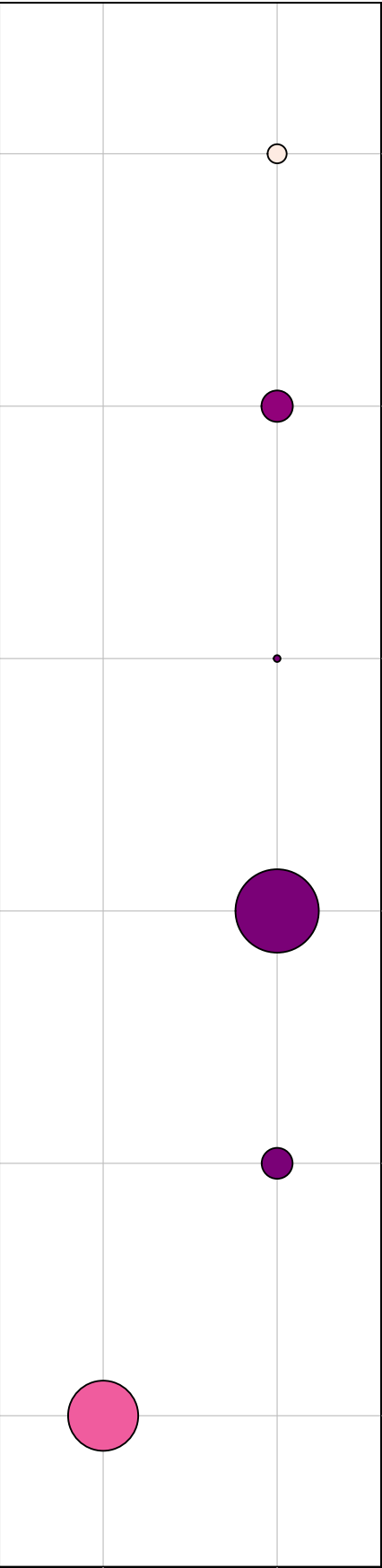


Nvec_vc1.1_XM_048734599.1
GPCR–rhodopsin:like:ADORA2B:likeclu; 7tm_1/



Nvec_vc1.1_XM_048723891.1

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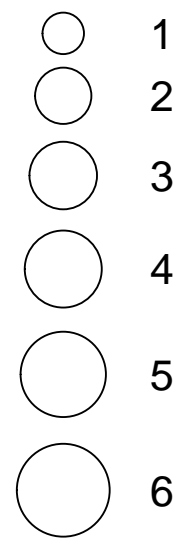
Fox_pos

Ncol_pos

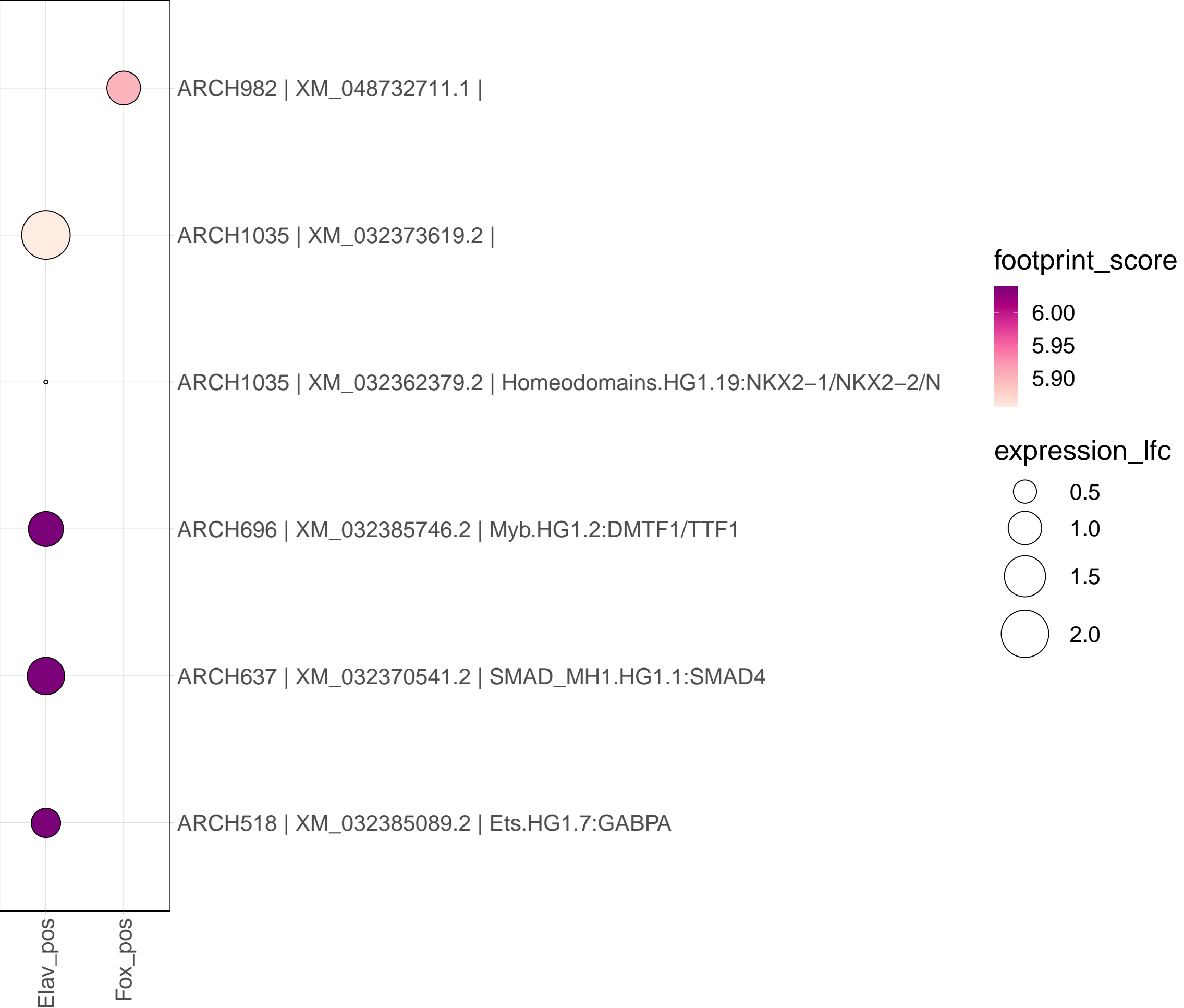
footprint_score



expression_lfc

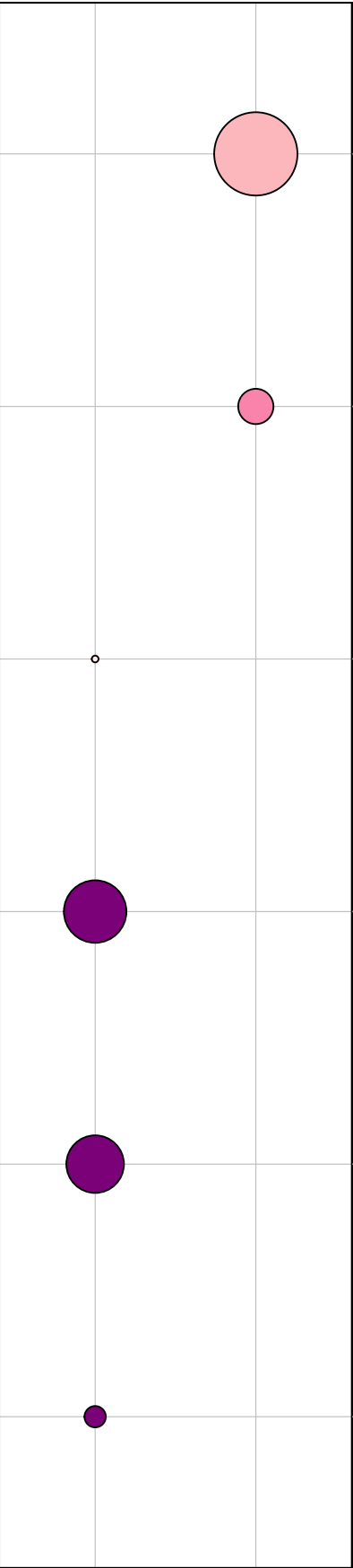


Nvec_vc1.1_XM_032386096.2
Exportin-T; Xpo1/



Nvec_vc1.1_XM_032363157.2

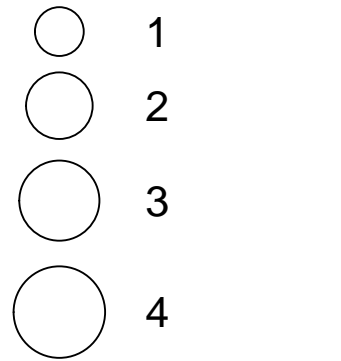
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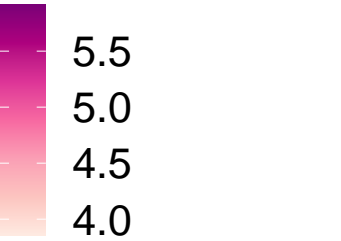
Elav_pos

Fox_pos

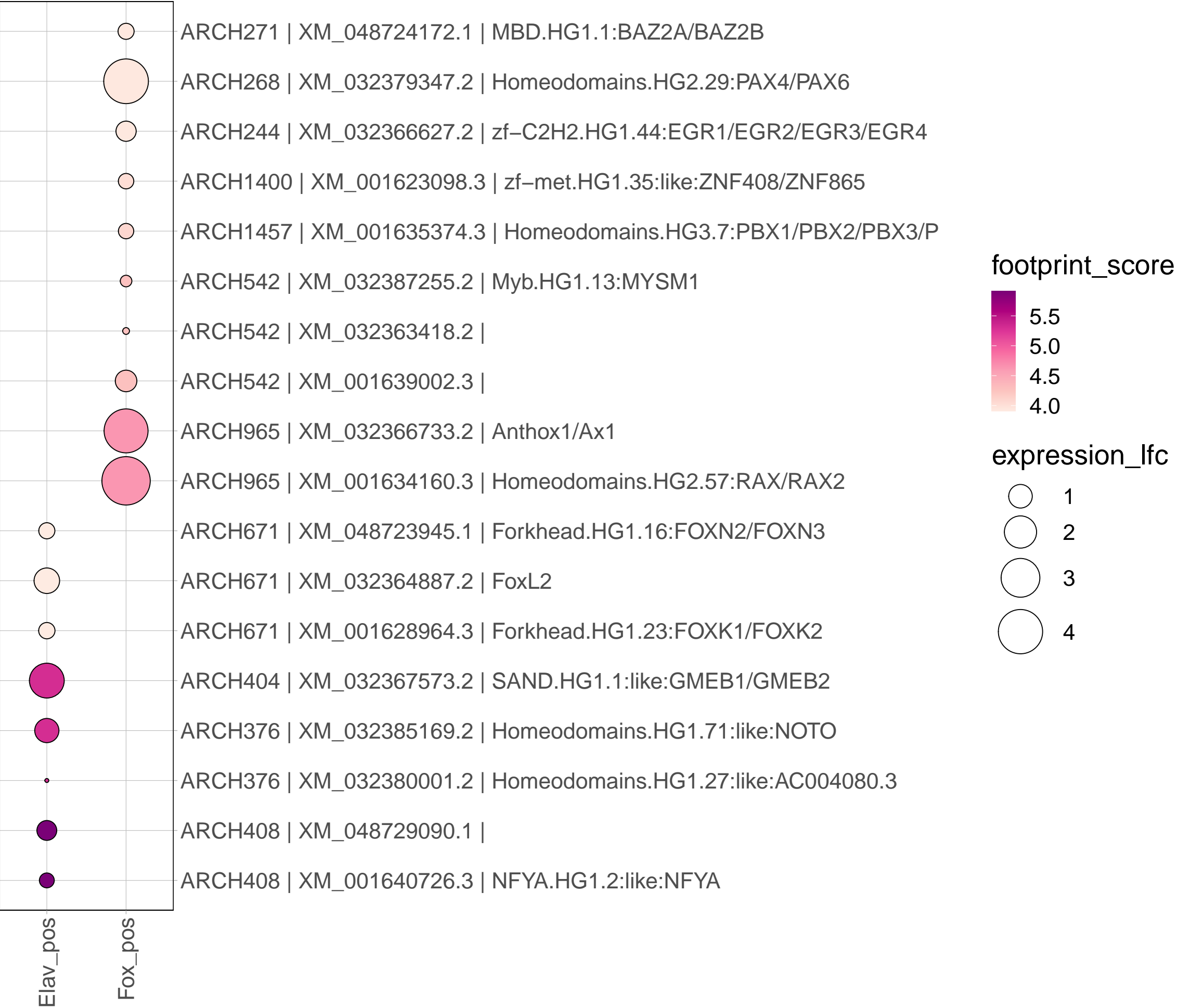
expression_lfc



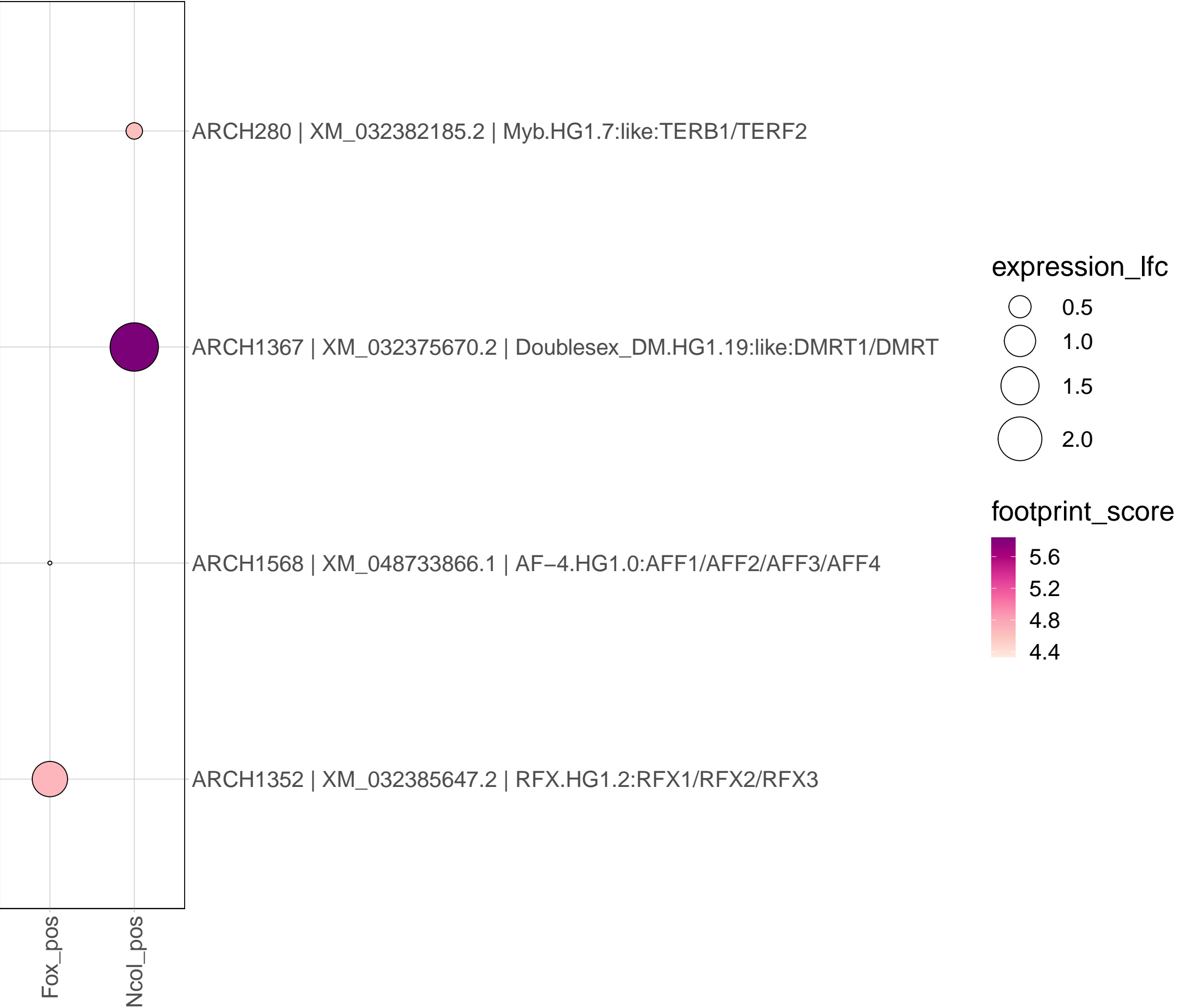
footprint_score



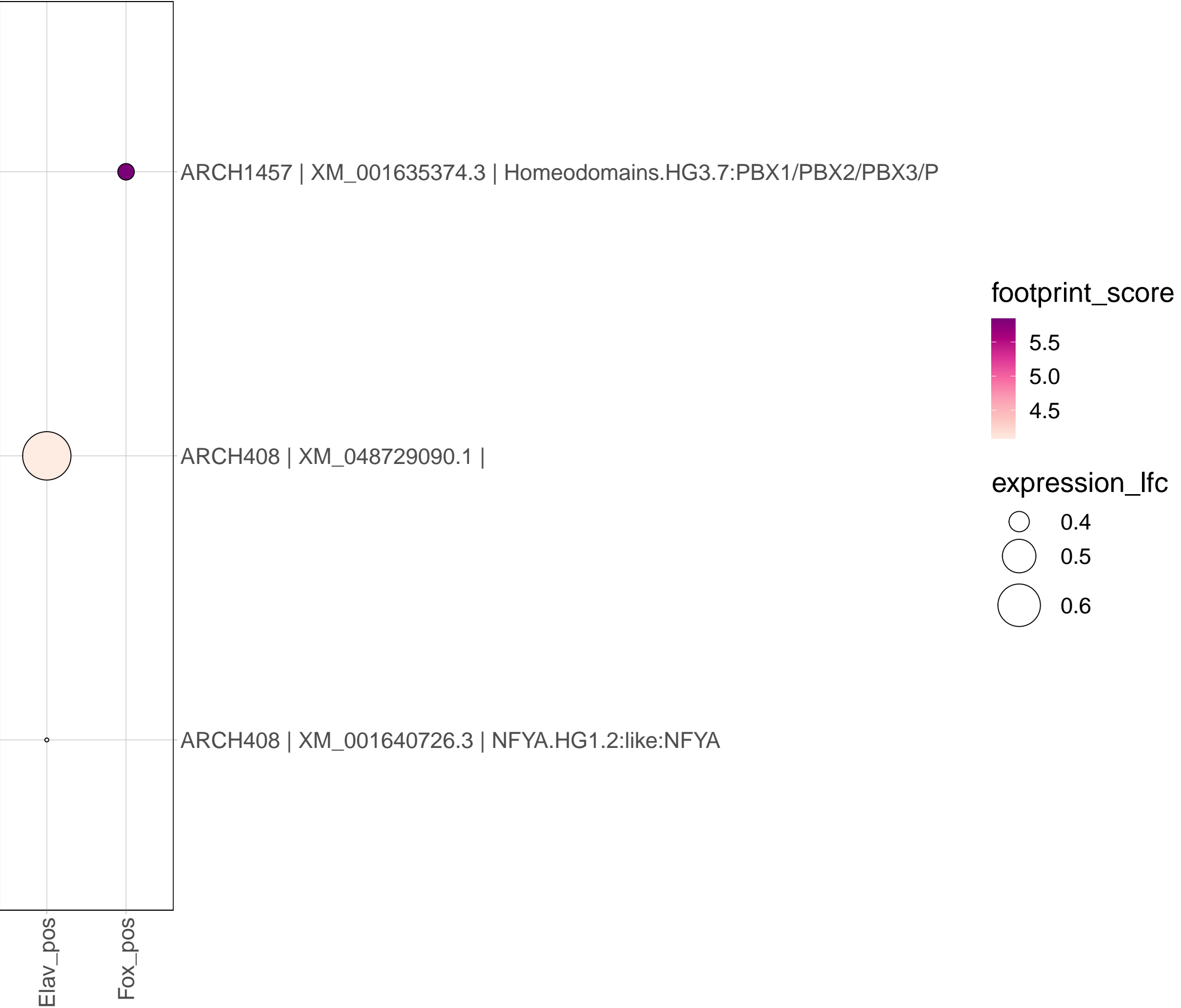
Nvec_vc1.1_XM_001633244.3
Synaptotagmin:C2CD4C/C2CD4D/TC2N; Synaptotagmin.HG1.17:C2CD4C/C2CD4D/TC2N



Nvec_vc1.1_XM_001639954.3
GPCR–rhodopsin:like:GPR27/GPR50/GPR; 7tm_1/



Nvec_vc1.1_XM_032372557.2
Synaptotagmin:like:FO681492/SYT15:l; C2/C2/

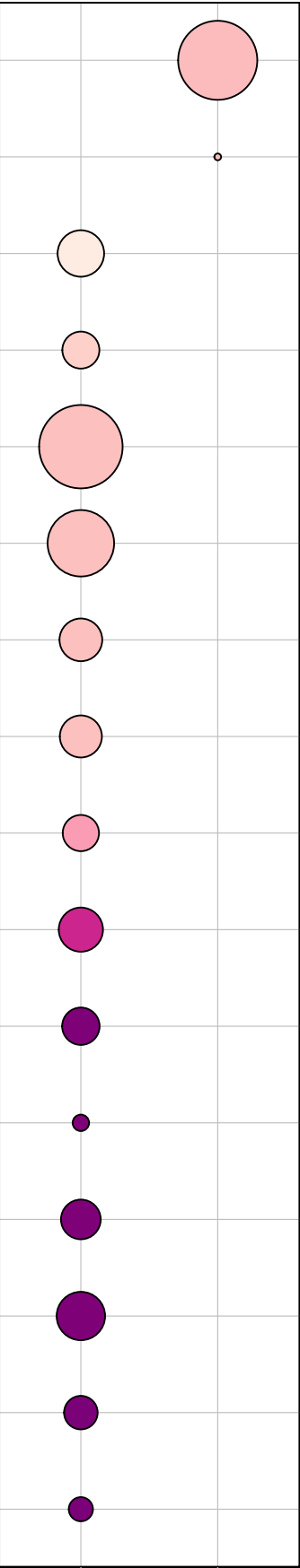


Nvec_vc1.1_XM_001636882.3
ADP-ribosylation factor-like 4C; Arf/



Nvec_vc1.1_XM_048719664.1

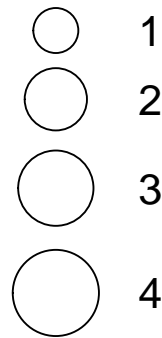
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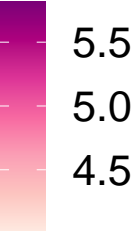
Elav_pos

Ncol_pos

expression_lfc

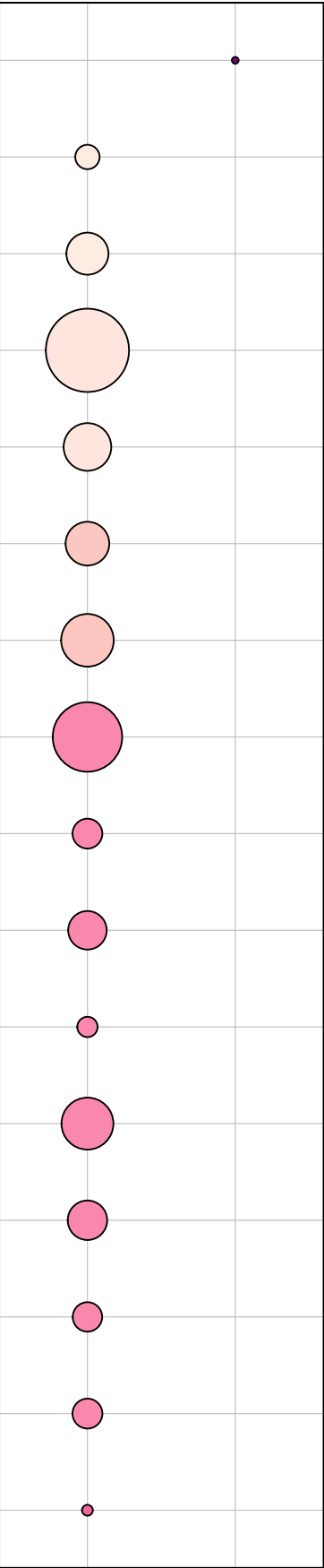


footprint_score



Nvec_vc1.1_XM_048725869.1

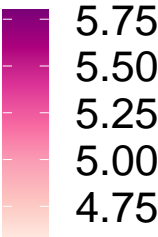
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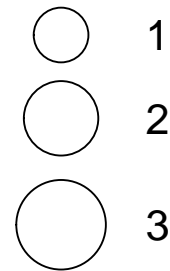
Elav_pos

Fox_pos

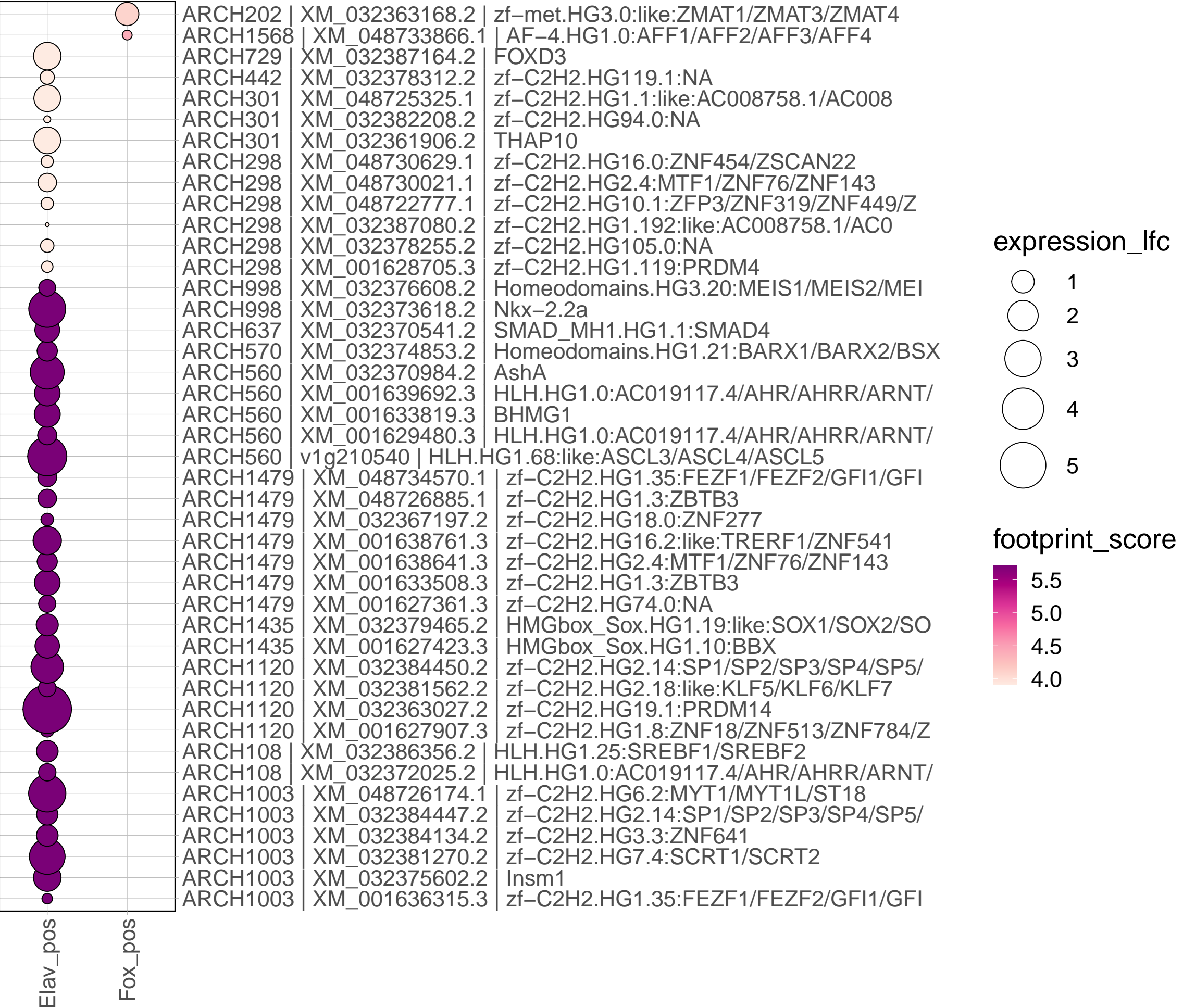
footprint_score



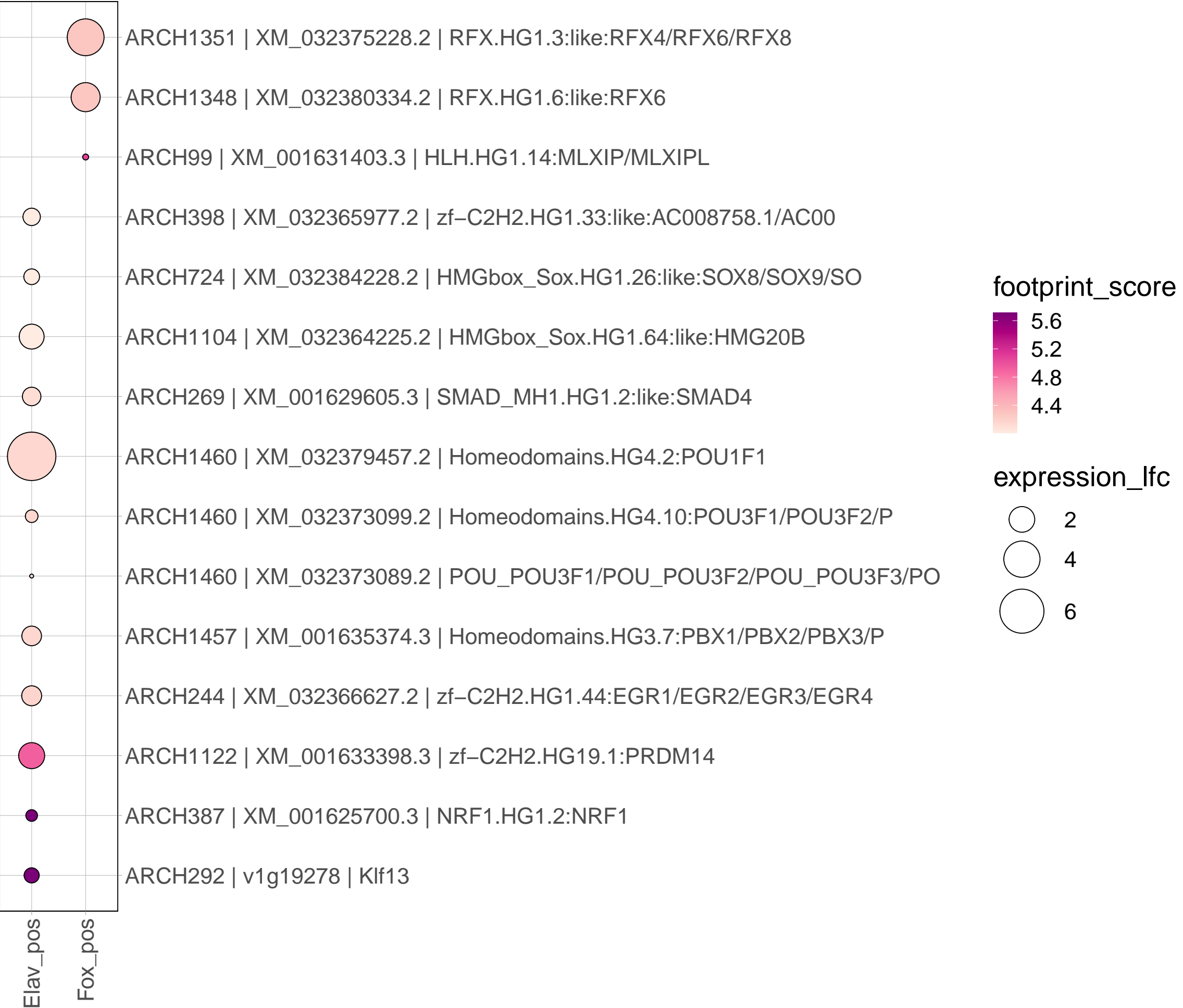
expression_lfc



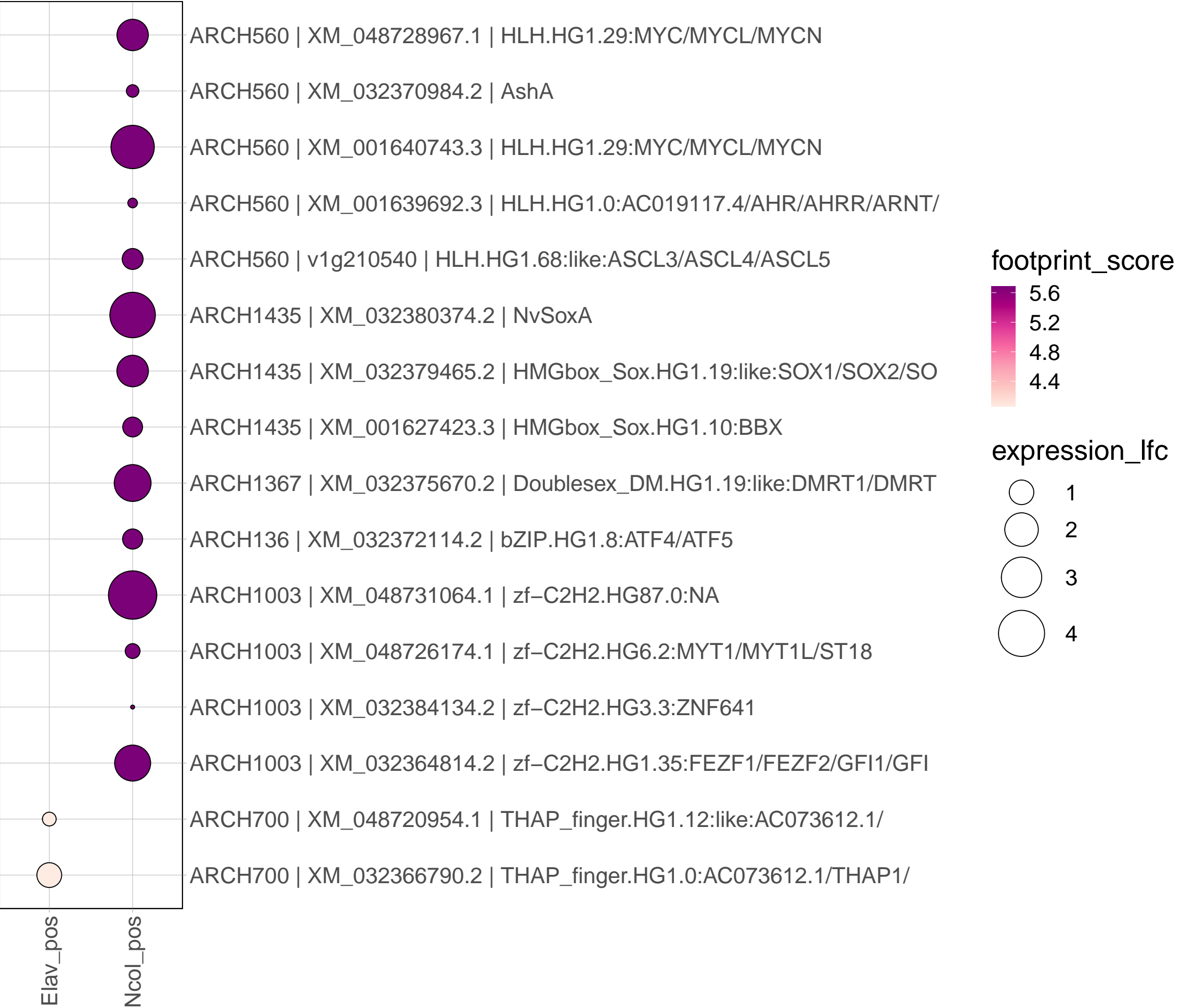
Nvec_vc1.1_XM_001632492.3
Alpha-(1,3)-fucosyltransferase 4; Glyco_transf_10/



Nvec_vc1.1_XM_001639681.3
Potassium channel, subfamily K, mem; Ion_trans_2/Ion_trans_2/

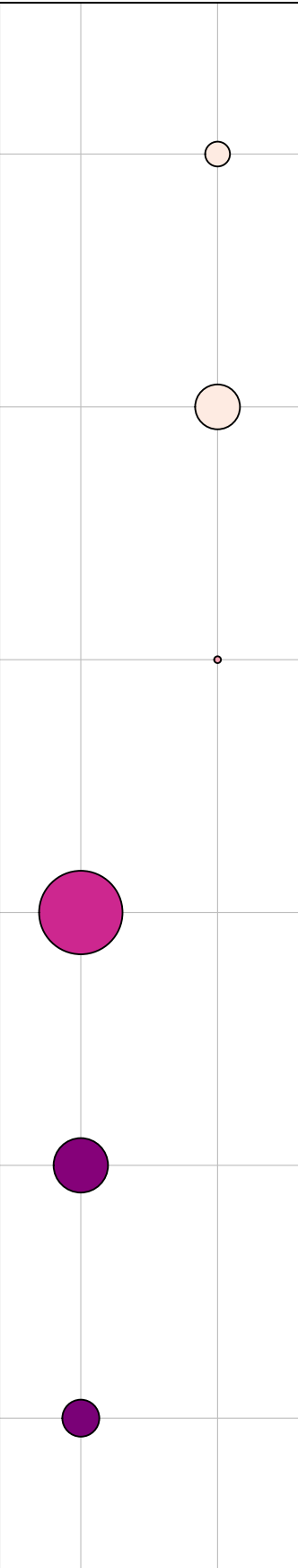


Nvec_vc1.1_XM_048729870.1
60 kDa lysophospholipase; Asparaginase/Ank_2/



Nvec_NVE13928

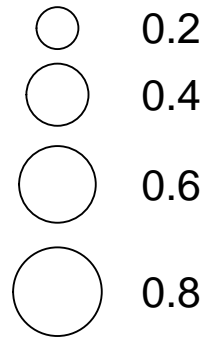
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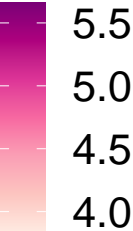
Fox_pos

Ncol_pos

expression_lfc



footprint_score



ARCH12 | XM_032366646.2 | ARID_BRIGHT.HG1.5:ARID5A/ARID5B

ARCH12 | XM_032362135.2 | ARID_BRIGHT.HG1.9:ARID3A/ARID3B/ARI

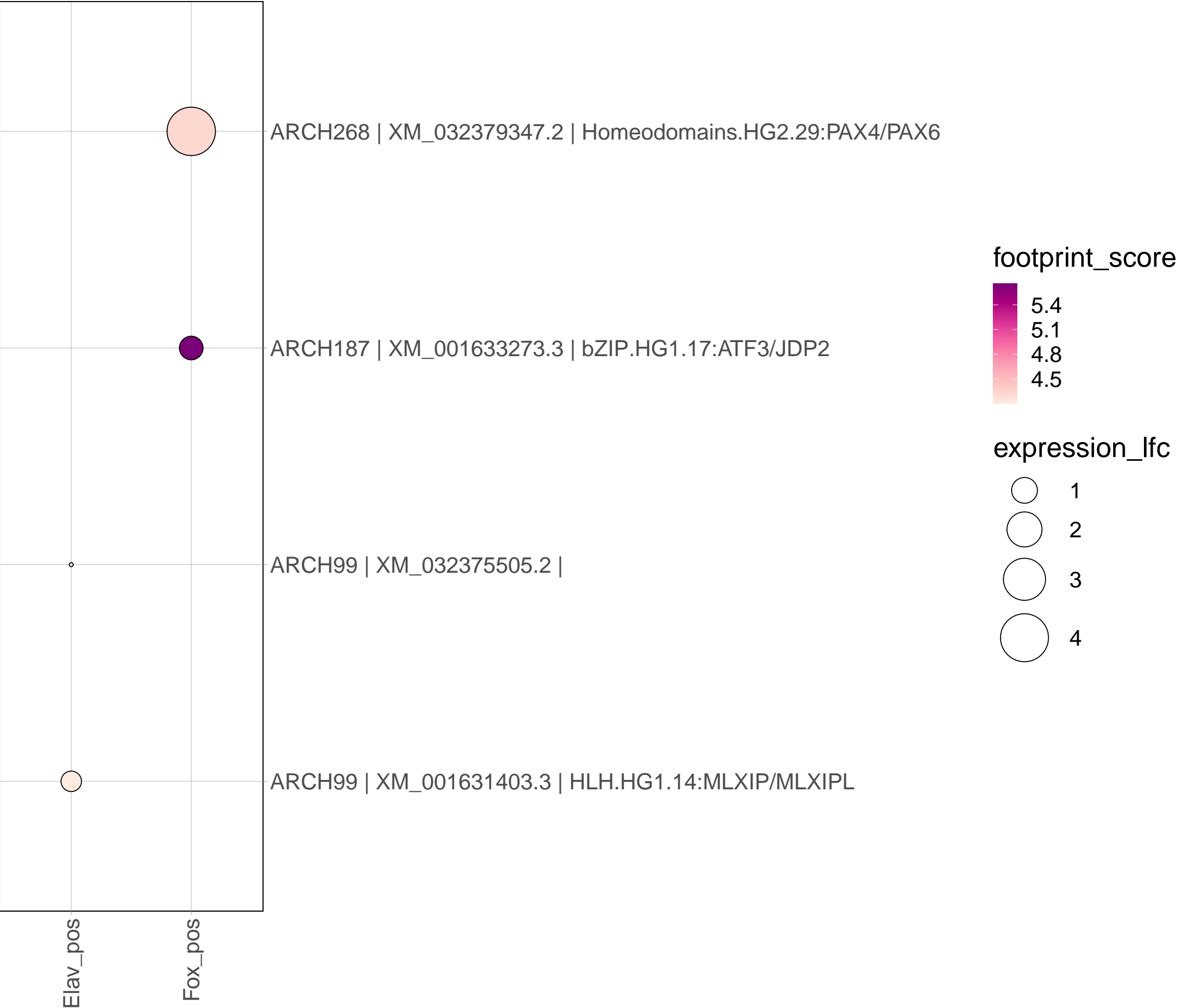
ARCH429 | XM_032362945.2 | Myb.HG1.8:DNAJC1/DNAJC2

ARCH1434 | XM_032372264.2 | NvSoxC

ARCH1457 | XM_001635374.3 | Homeodomains.HG3.7:PBX1/PBX2/PBX3/P

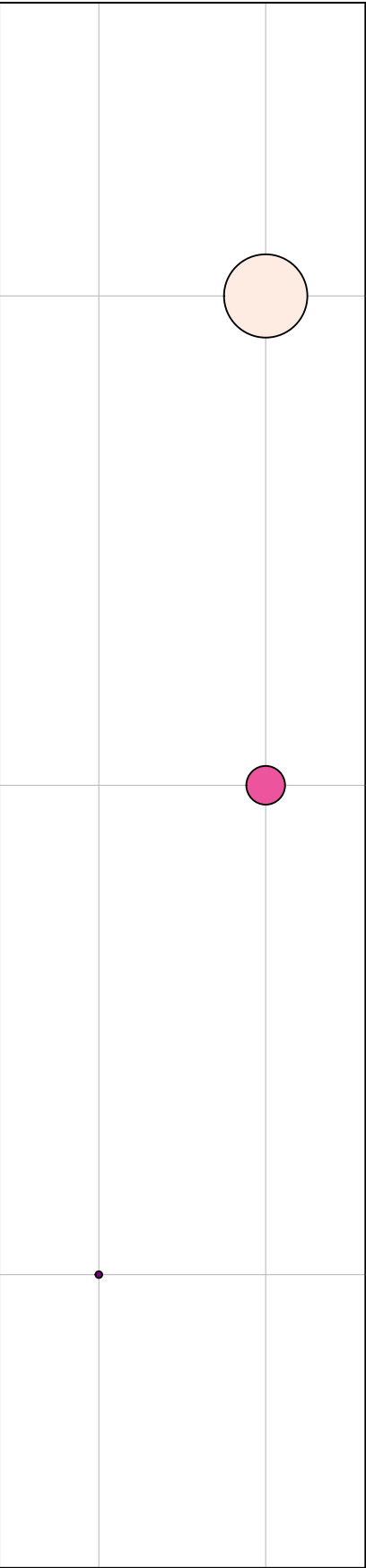
ARCH993 | XM_001634845.3 | zf-CXXC.HG1.8:CXXC4/CXXC5/TET1/TET3

Nvec_vc1.1_XM_032364349.2
Aspartate aminotransferase; Aminotran_1_2/



Nvec_vc1.1_XM_032376979.2

;

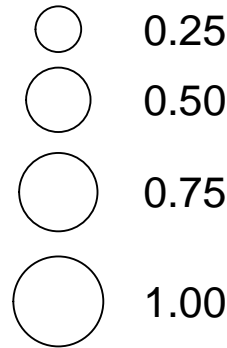


ARCH700 | XM_032366790.2 | THAP_finger.HG1.0:AC073612.1/THAP1/

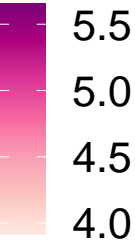
ARCH325 | XM_032385905.2 | zf-C2H2.HG1.3:ZBTB3

ARCH722 | XM_001636155.3 | NvSoxF1

expression_lfc



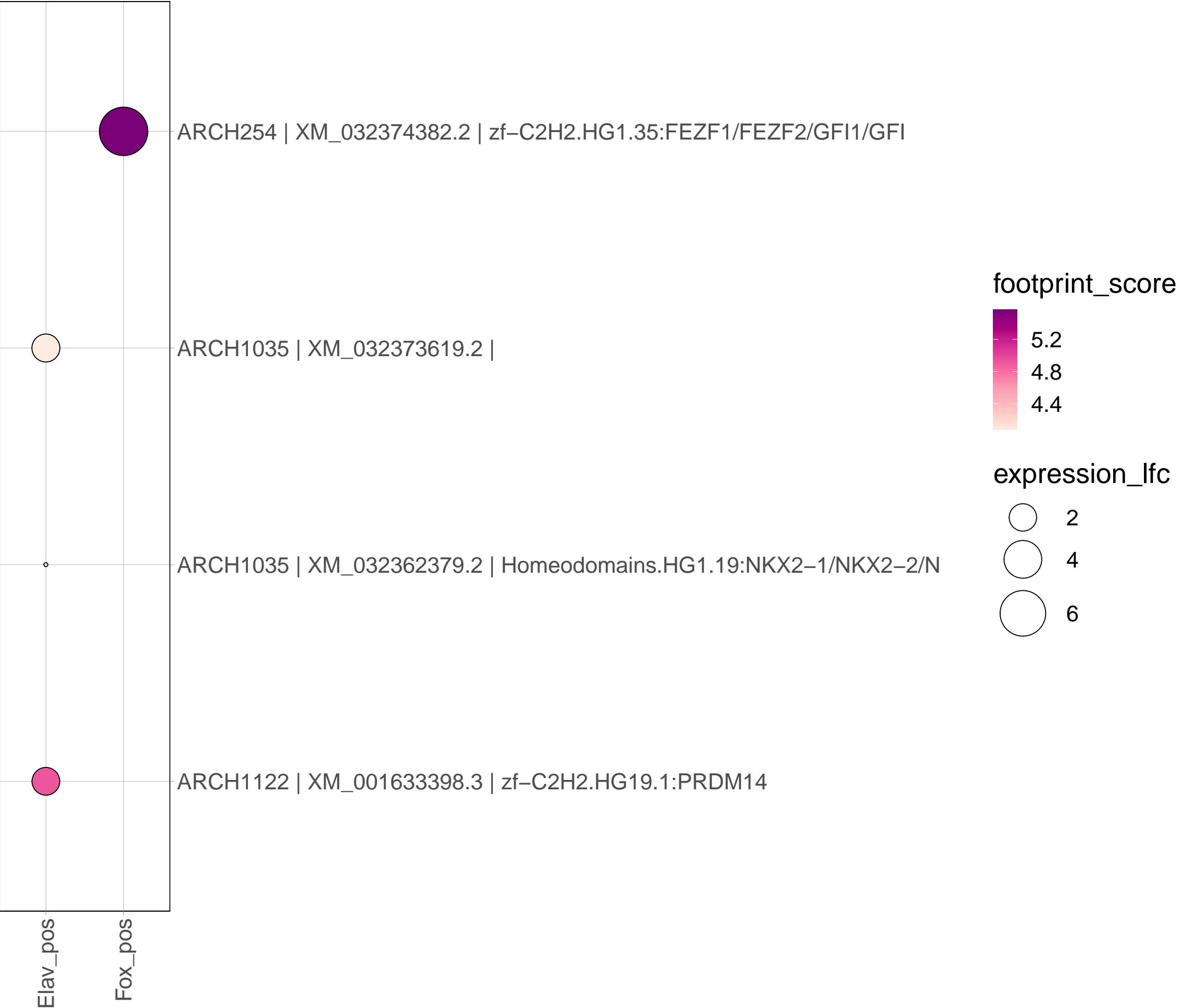
footprint_score



Elav_pos

Fox_pos

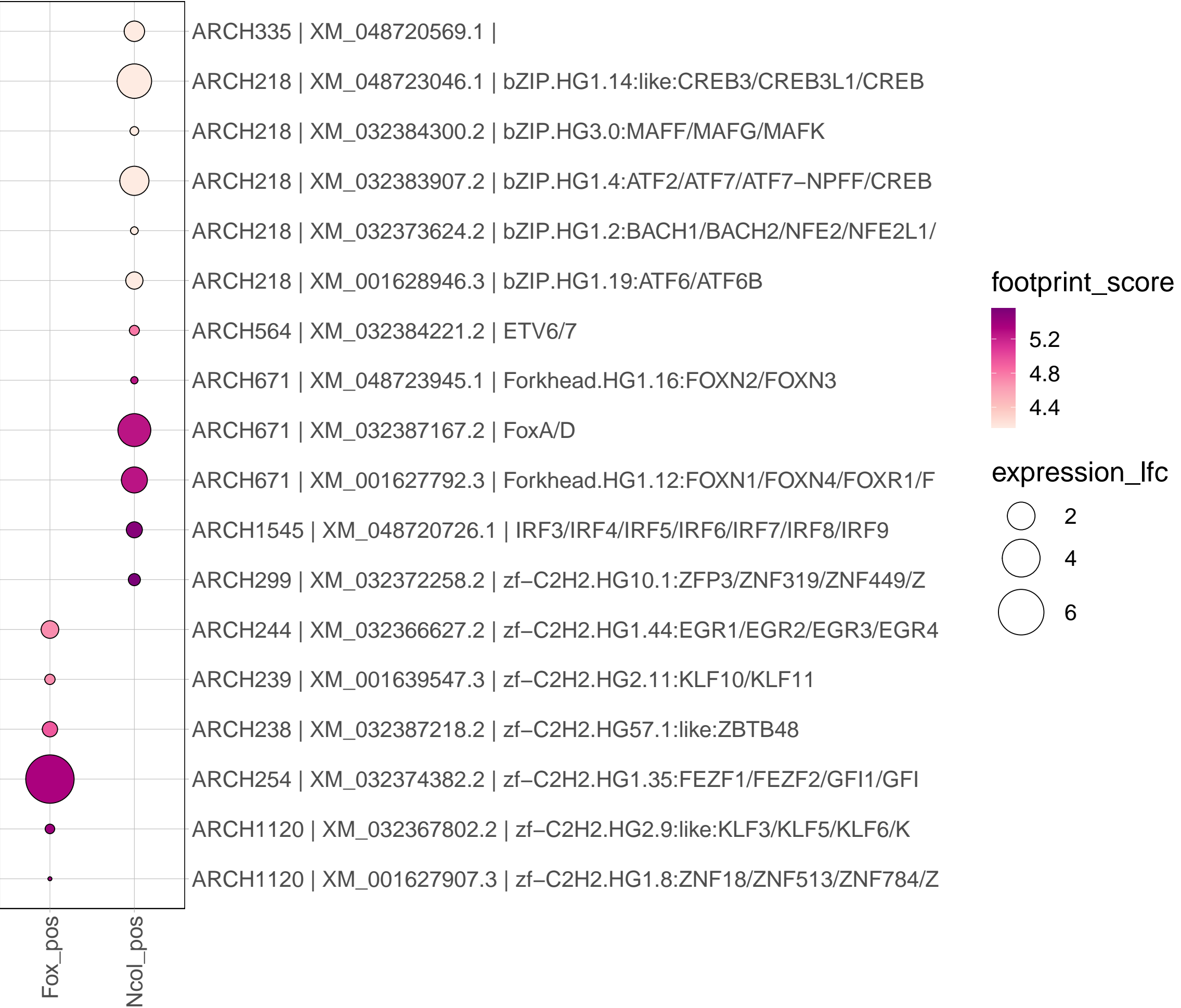
Nvec_vc1.1_XM_048730108.1
GPCR-rhodopsin:NA; 7tm_1/



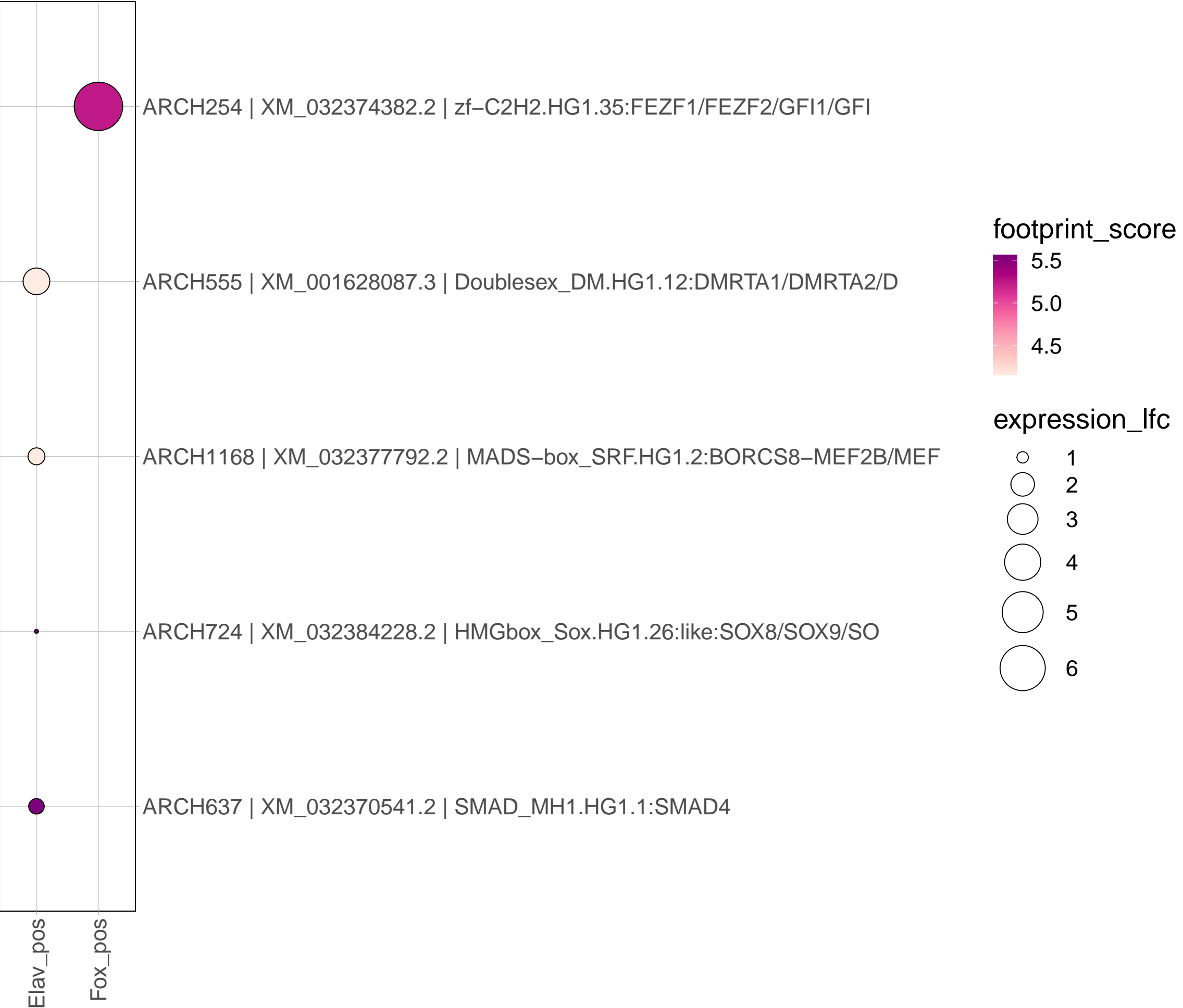
Nvec_vc1.1_XM_032382511.2
Serine incorporator 1; Serinc/Serinc/



Nvec_vc1.1_XM_032382117.2
GPCR–rhodopsin:like:GPR176:likeclu:; 7tm_1/

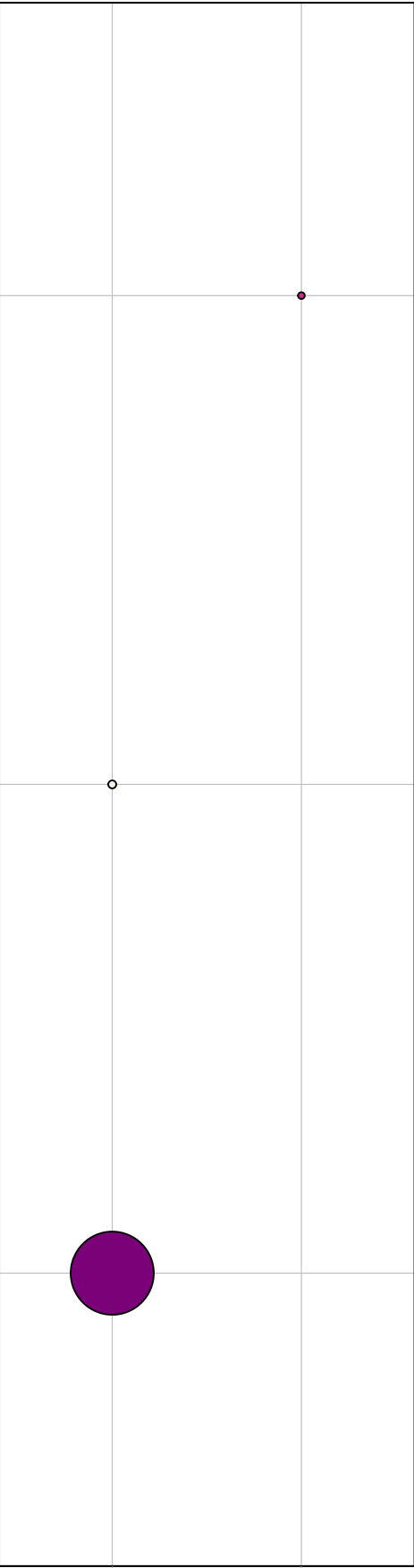


Nvec_vc1.1_XM_032386298.2
GPCR–rhodopsin:NA; 7tm_1/



Nvec_vc1.1_XM_032362299.2

;
;

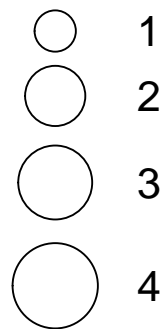


ARCH269 | XM_001629605.3 | SMAD_MH1.HG1.2:like:SMAD4

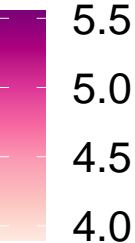
ARCH442 | XM_032378312.2 | zf-C2H2.HG119.1:NA

ARCH1272 | XM_032383641.2 | zf-C4_Nuclear_receptors.HG1.9:like:

expression_lfc



footprint_score



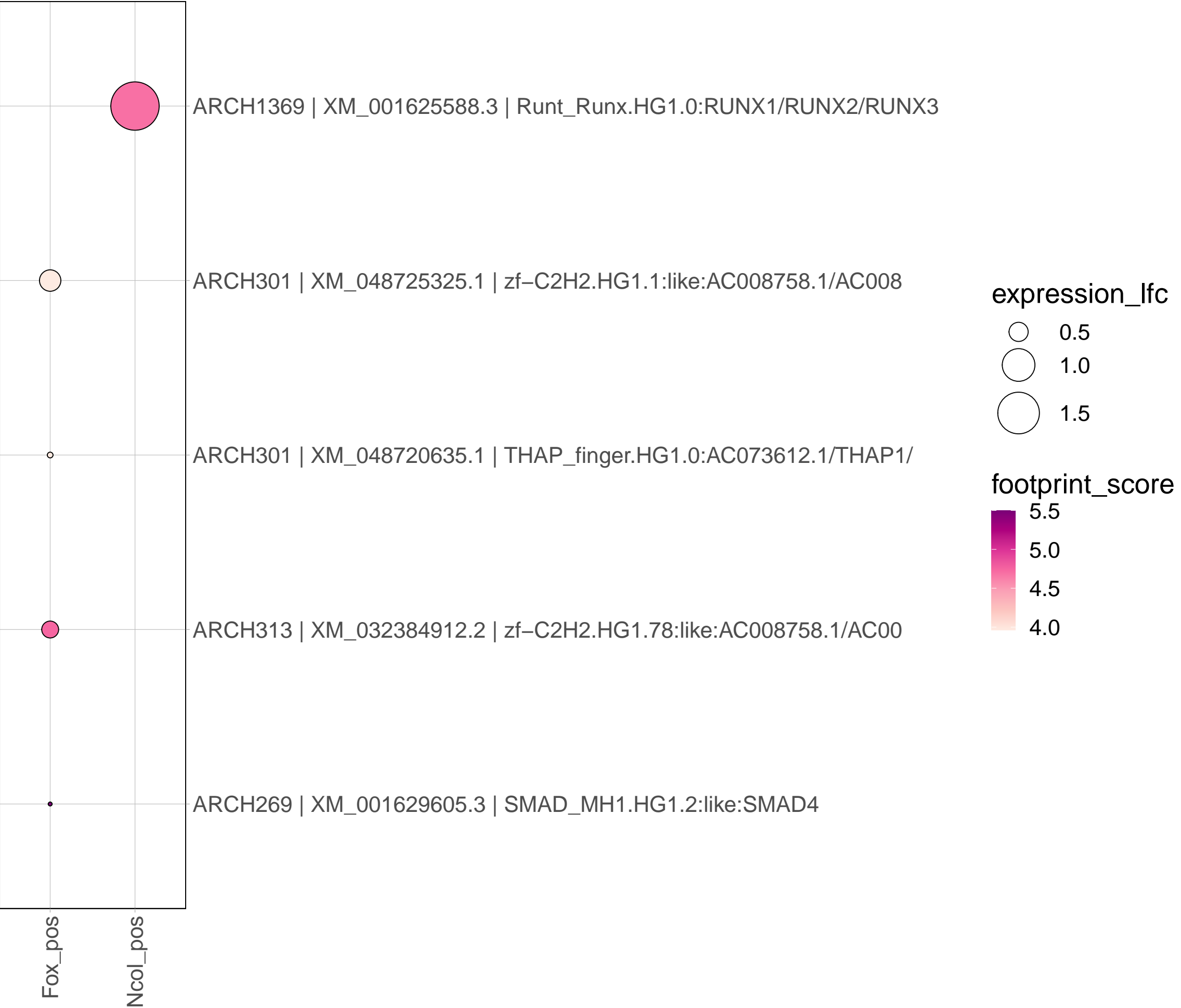
Elav_pos

Fox_pos

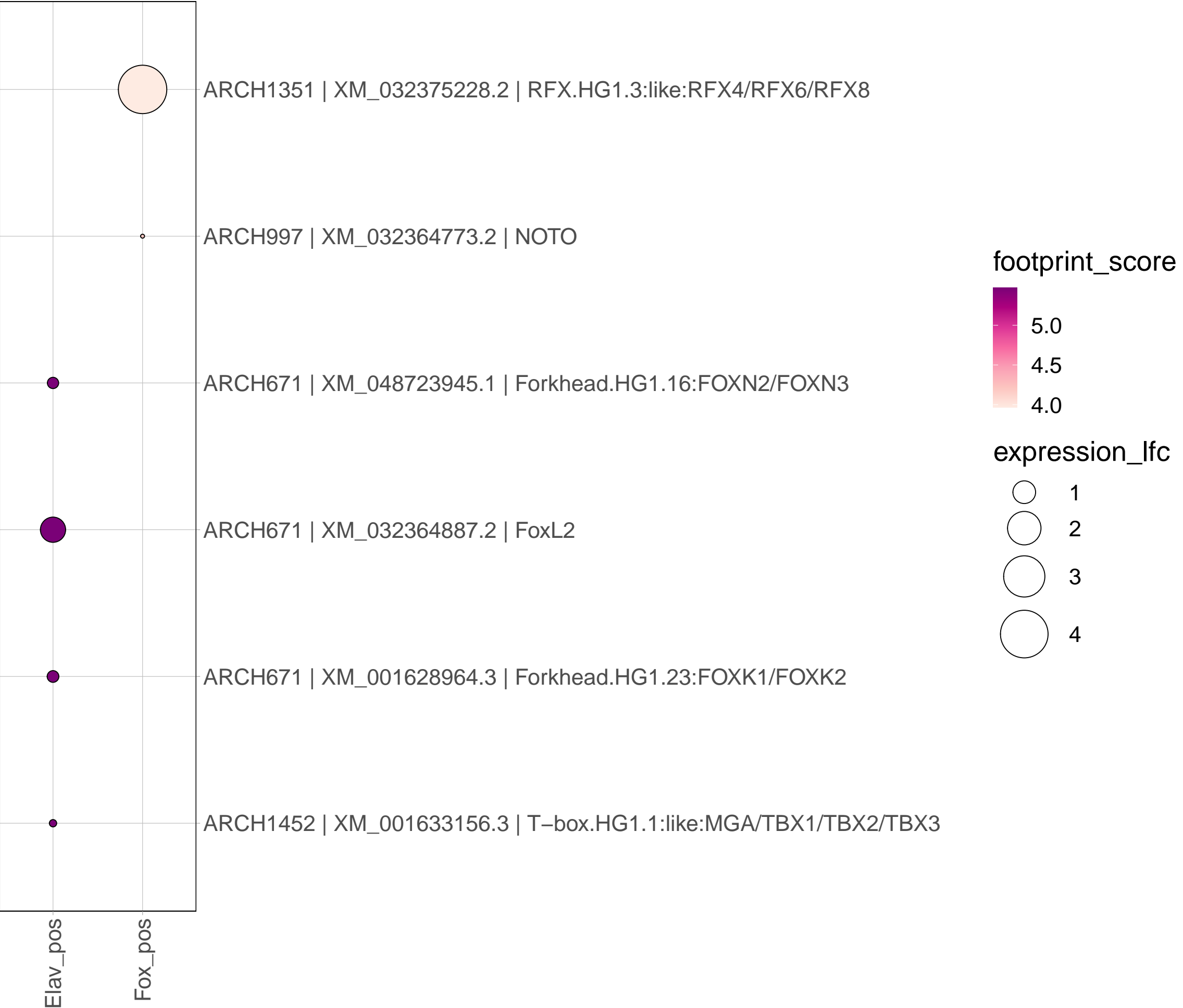
Nvec_vc1.1_XM_001633615.3
HCG2044780; Kua-UEV1_localIn/



Nvec_vc1.1_XM_001636545.3
; RNase_P_Rpp14/



Nvec_vc1.1_XM_048720443.1
C-Maf-inducing protein;

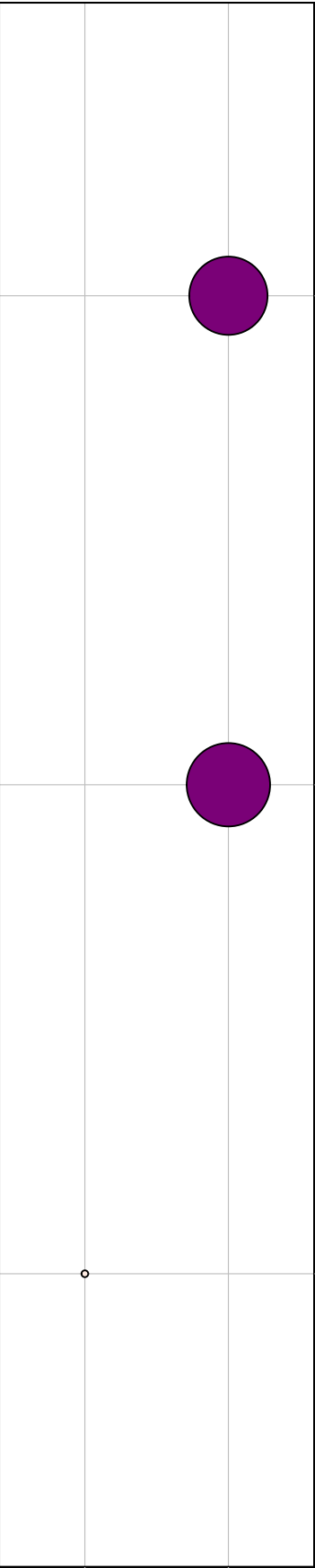


Nvec_vc1.1_XM_032384695.2
Monocarboxylate transporter 10; MFS_1/



Nvec_vc1.1_XM_048731044.1

;



Elav_pos

Ncol_pos

ARCH688 | XM_032370011.2 | Homeodomains.HG1.37:NKX6-1/NKX6-2/N

ARCH1151 | XM_032363992.2 | Pou4

ARCH637 | XM_032370541.2 | SMAD_MH1.HG1.1:SMAD4

footprint_score



5.0

4.5

4.0

expression_lfc



1.6

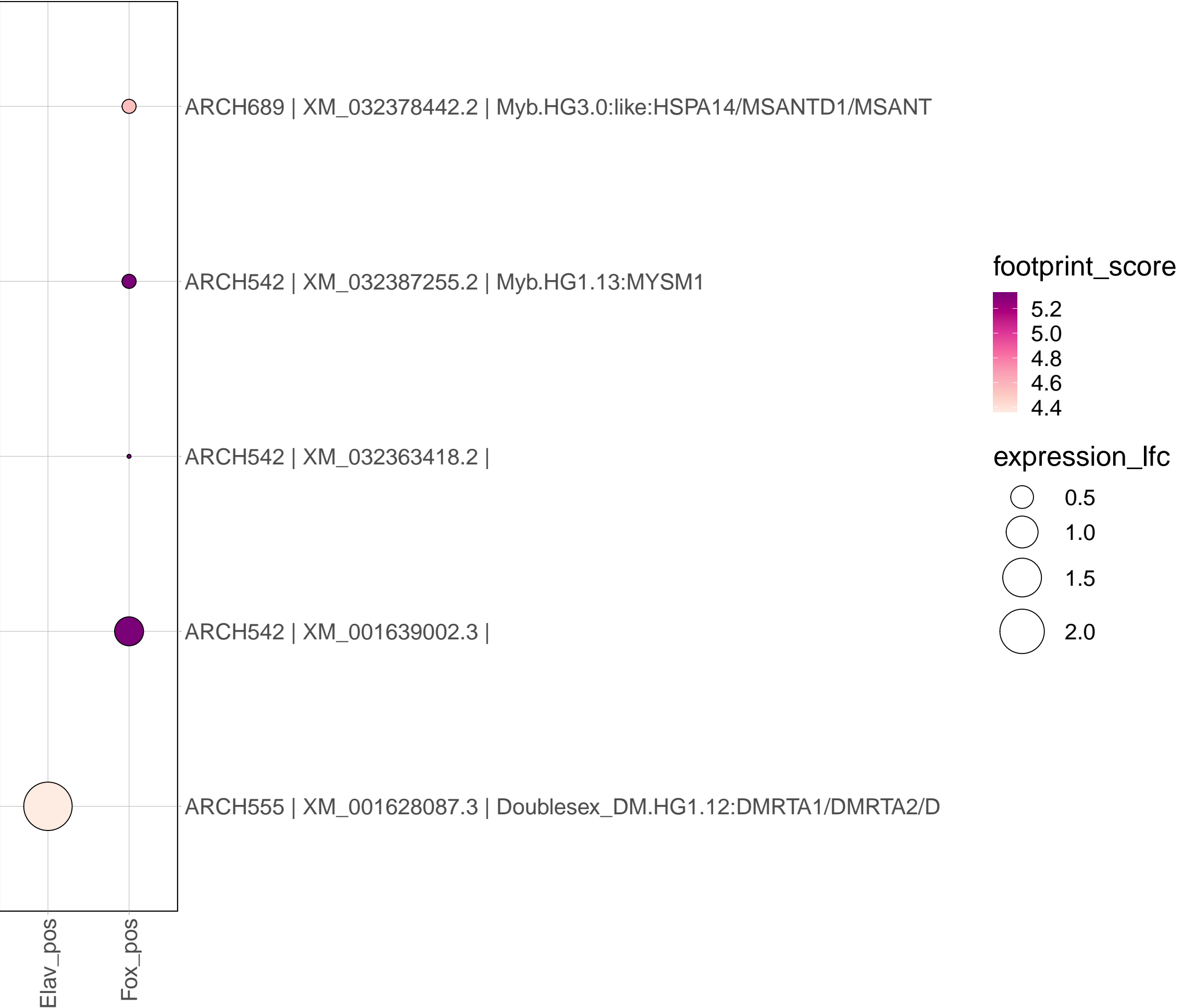


2.0

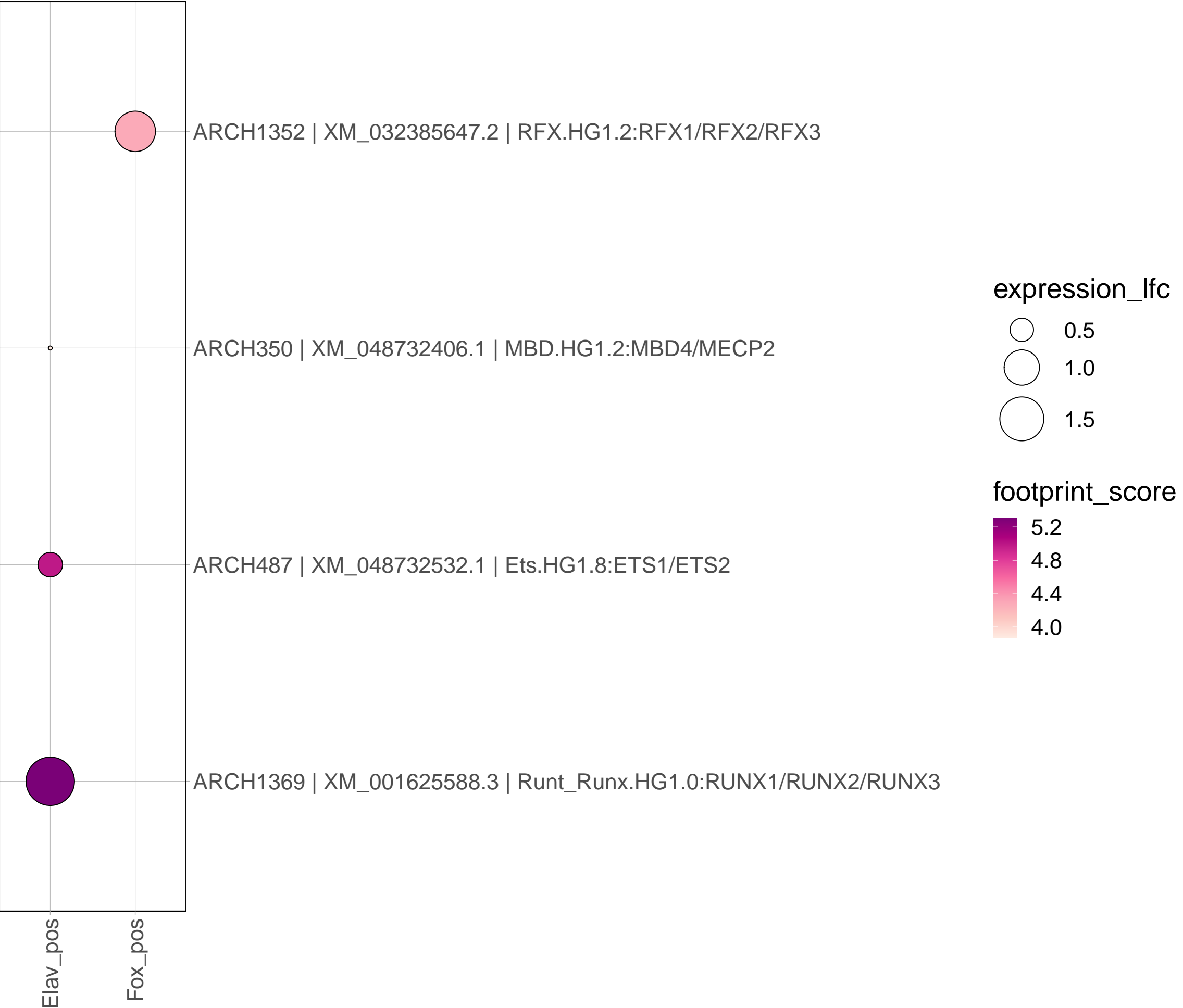


2.4

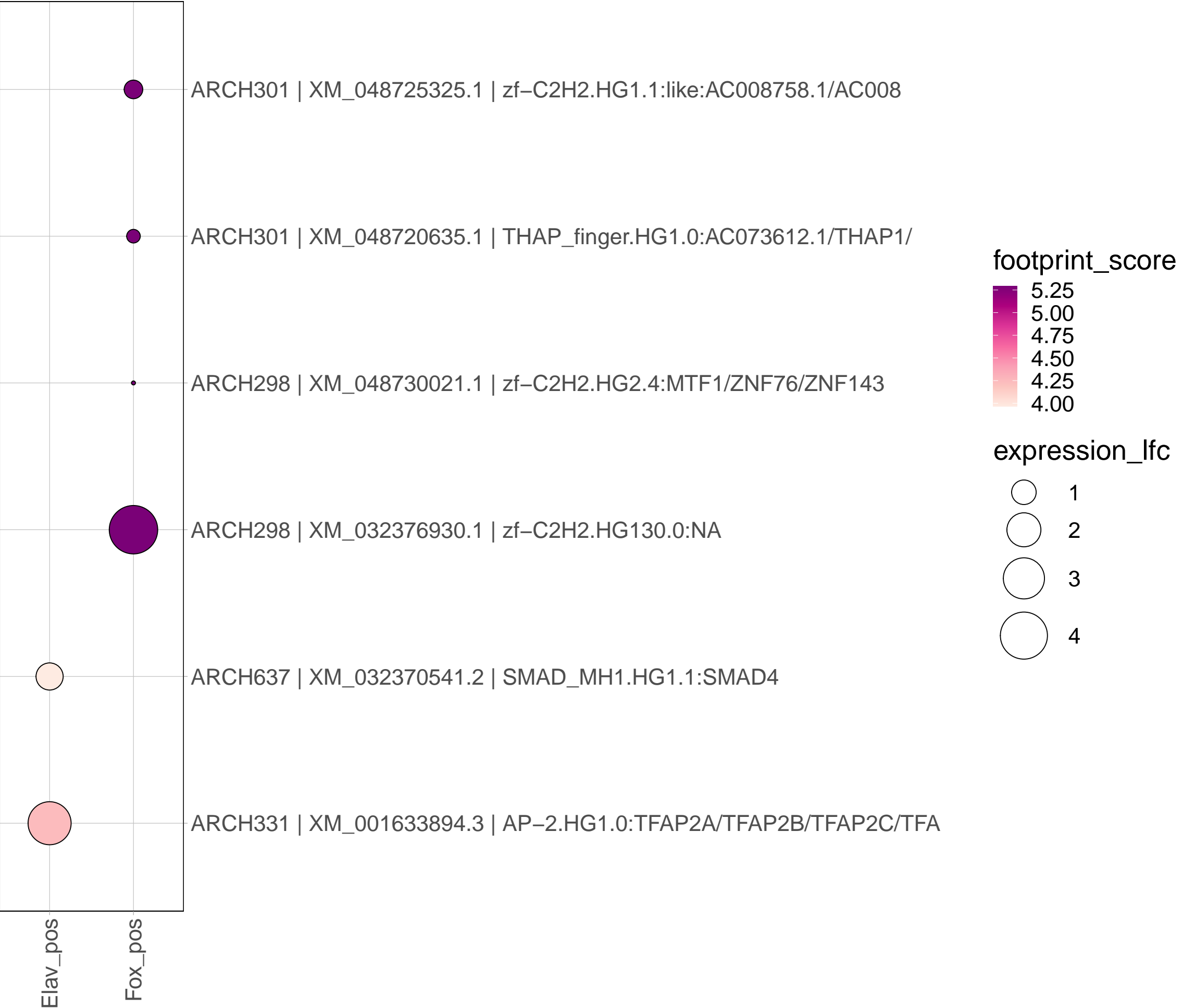
Nvec_vc1.1_XM_001629129.3
Probable aminopeptidase NPEPL1; Peptidase_M17/



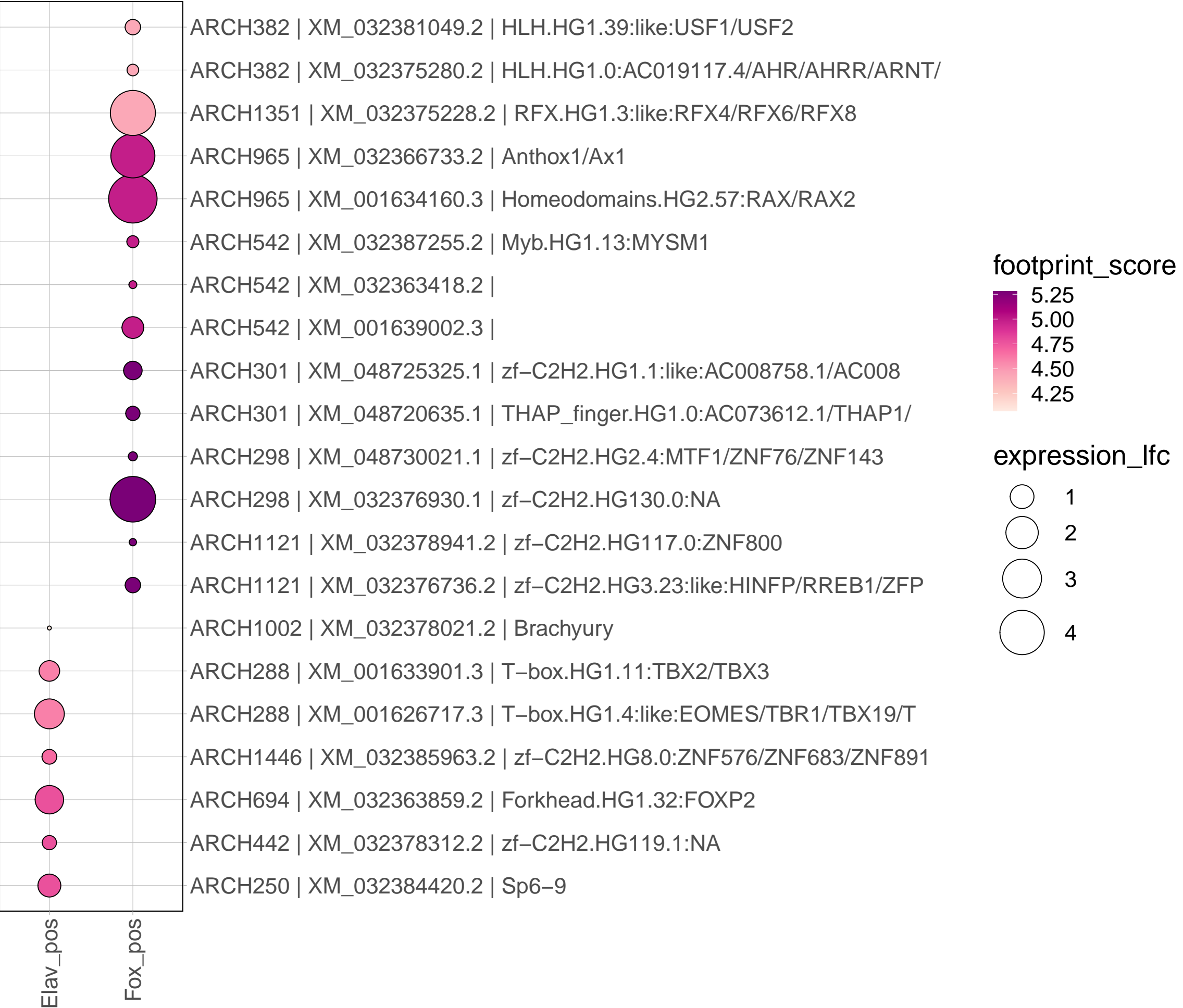
Nvec_vc1.1_XM_032364225.2
HMGbox_Sox.HG1.64:like:HMG20B;

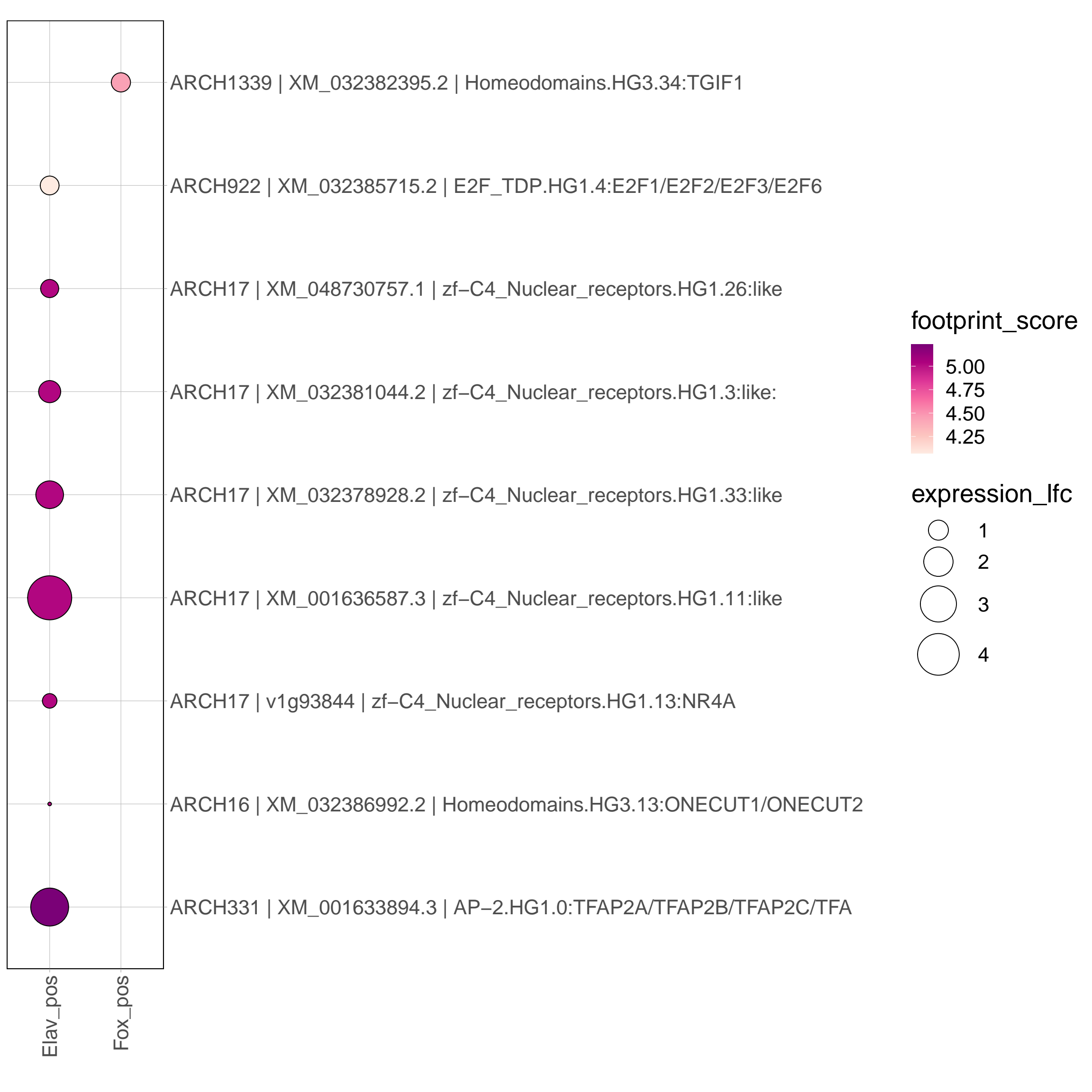


Nvec_vc1.1_XM_032373148.2
S1:SUPT6H; S1.HG5.0:SUPT6H



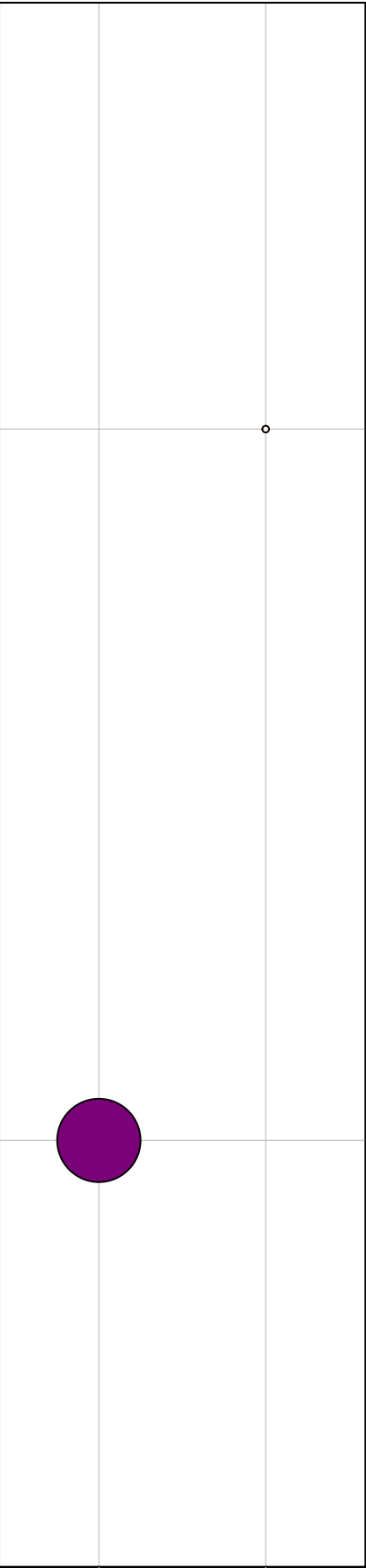
Nvec_vc1.1_XM_032381433.2
Guanylate cyclase soluble subunit b; HNOB/HNOBA/Guanylate_cyc/





Nvec_NVE7893

;



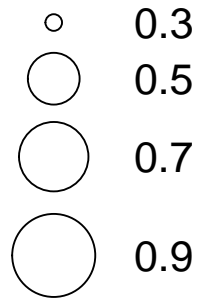
Fox_pos

Ncol_pos

ARCH59 | XM_032384433.2 | HLH.HG1.86:TWIST1/TWIST2

ARCH700 | XM_032366790.2 | THAP_finger.HG1.0:AC073612.1/THAP1/

expression_lfc

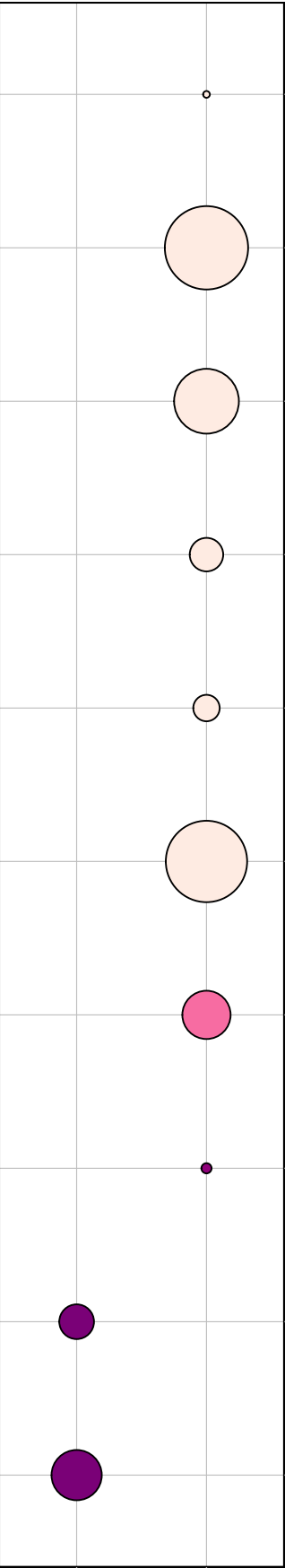


footprint_score



Nvec_vc1.1_XM_032372214.2

;



Elav_pos

Ncol_pos

ARCH671 | XM_048723945.1 | Forkhead.HG1.16:FOXN2/FOXN3

ARCH671 | XM_032387167.2 | FoxA/D

ARCH671 | XM_001627792.3 | Forkhead.HG1.12:FOXN1/FOXN4/FOXR1/F

ARCH442 | XM_032378312.2 | zf-C2H2.HG119.1:NA

ARCH1460 | XM_032373099.2 | Homeodomains.HG4.10:POU3F1/POU3F2/P

ARCH1151 | XM_032363992.2 | Pou4

ARCH1425 | XM_001639701.3 | NvSox2

ARCH729 | XM_032387164.2 | FOXD3

ARCH382 | XM_032381049.2 | HLH.HG1.39:like:USF1/USF2

ARCH382 | XM_032375280.2 | HLH.HG1.0:AC019117.4/AHR/AHRR/ARNT/

expression_lfc



1



2

footprint_score



5.1

4.9

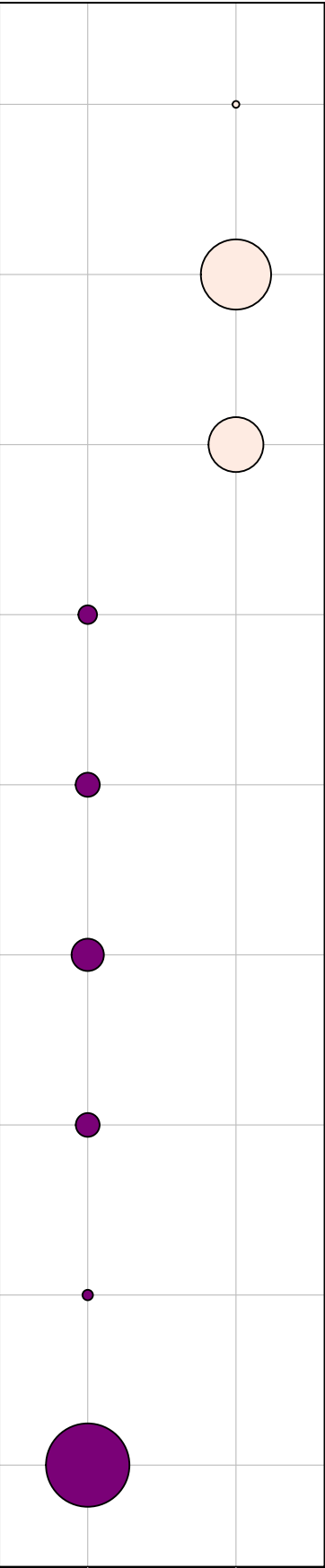
4.7

4.5

4.3

Nvec_vc1.1_XM_048730962.1

;
;



ARCH671 | XM_048723945.1 | Forkhead.HG1.16:FOXN2/FOXN3

ARCH671 | XM_032387167.2 | FoxA/D

ARCH671 | XM_001627792.3 | Forkhead.HG1.12:FOXN1/FOXN4/FOXR1/F

ARCH723 | XM_032371996.2 | HMGbox_Sox.HG1.53:LEF1/TCF7/TCF7L1/

ARCH570 | XM_001623084.3 | Six4/5

ARCH301 | XM_048725325.1 | zf-C2H2.HG1.1:like:AC008758.1/AC008

ARCH301 | XM_048720635.1 | THAP_finger.HG1.0:AC073612.1/THAP1/

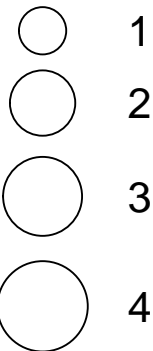
ARCH298 | XM_048730021.1 | zf-C2H2.HG2.4:MTF1/ZNF76/ZNF143

ARCH298 | XM_032376930.1 | zf-C2H2.HG130.0:NA

footprint_score



expression_lfc



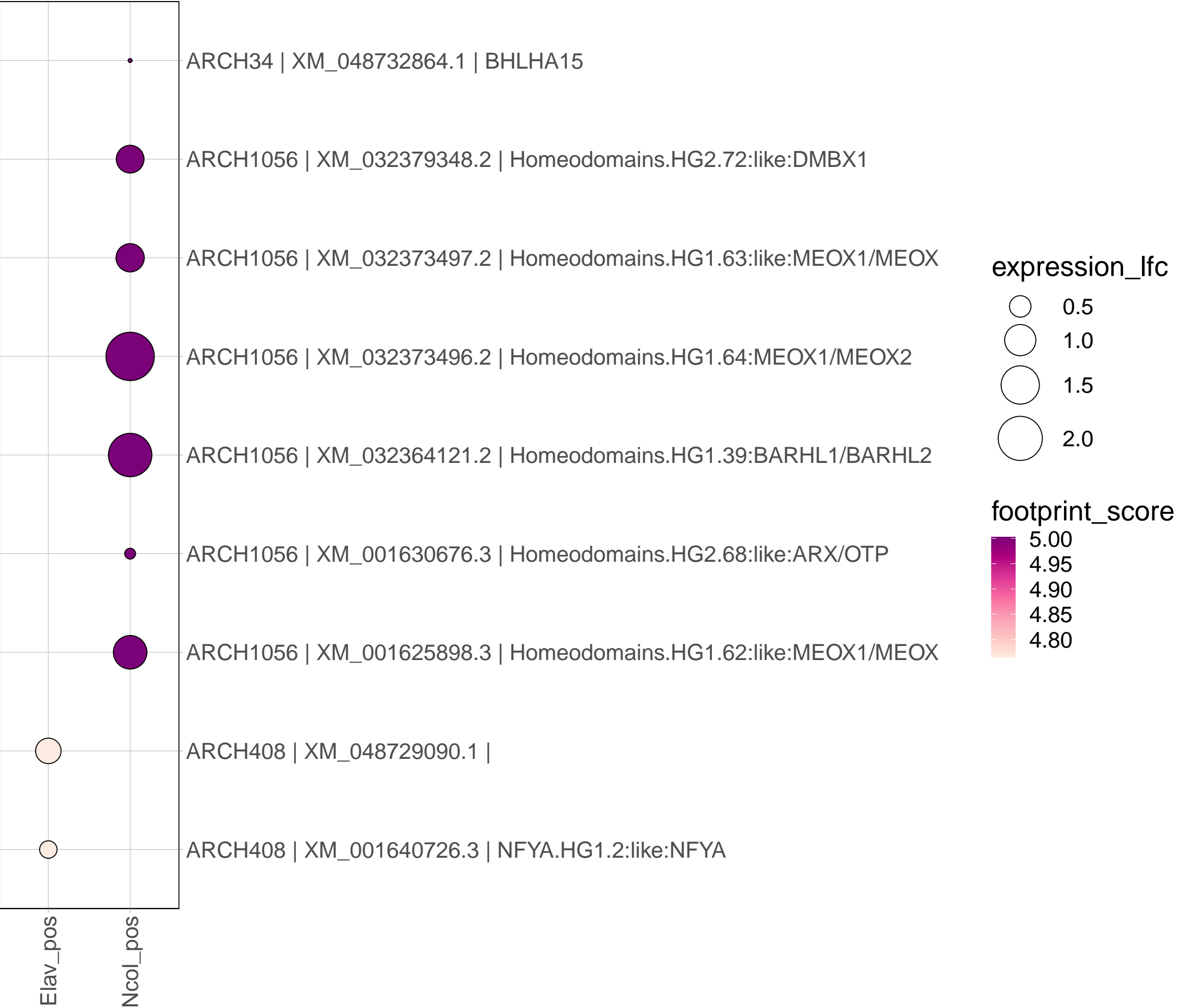
Nvec_vc1.1_XM_001637616.3
Fukutin related protein; LicD/



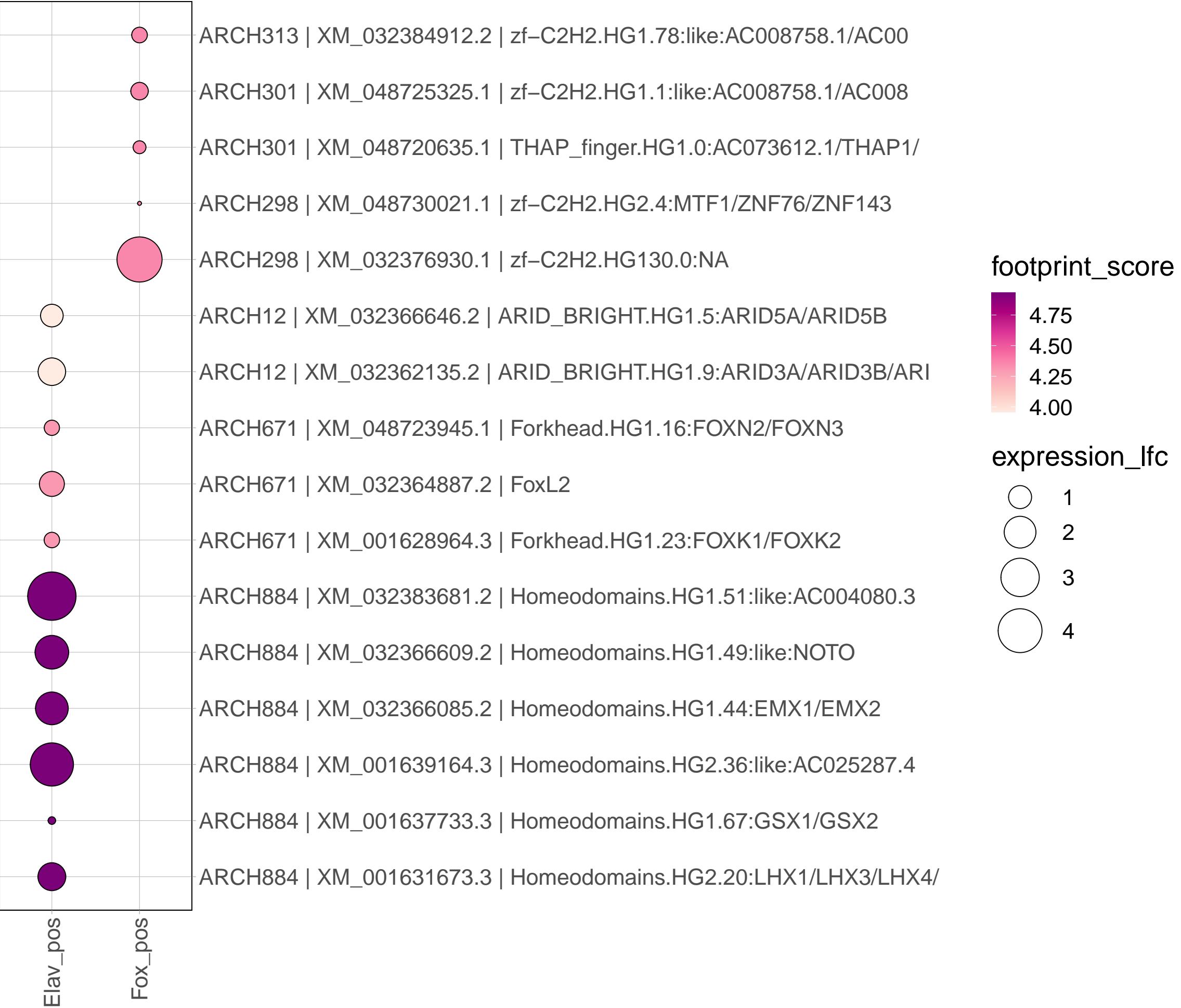
Nvec_vc1.1_XM_001628059.3
Fatty acid amide hydrolase 2; Amidase/



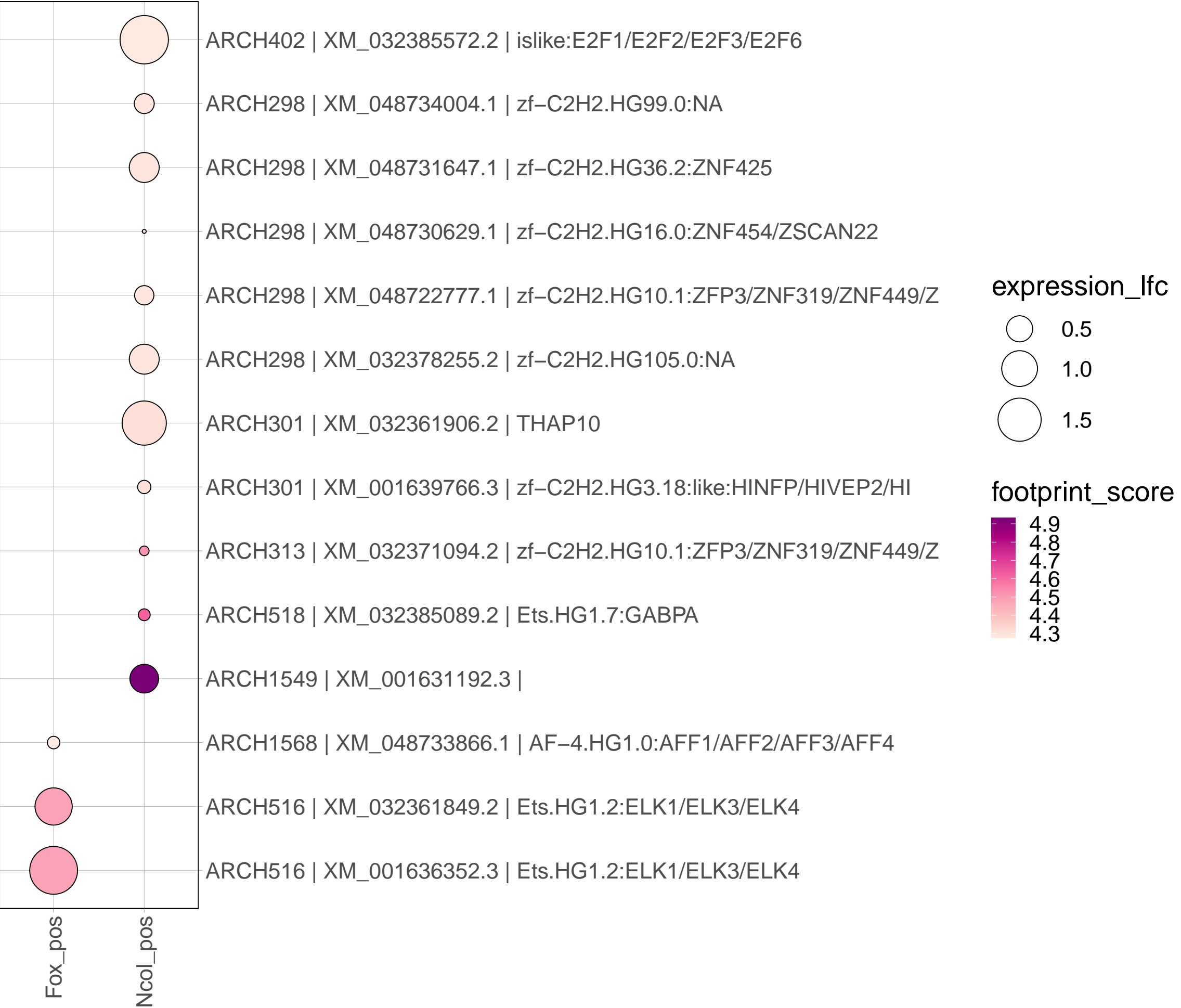
Nvec_vc1.1_XM_048730184.1
Thyroid receptor–interacting protei;



Nvec_vc1.1_XM_048724823.1
cDNA FLJ14638 fis, clone NT2RP20013; Lon_C/

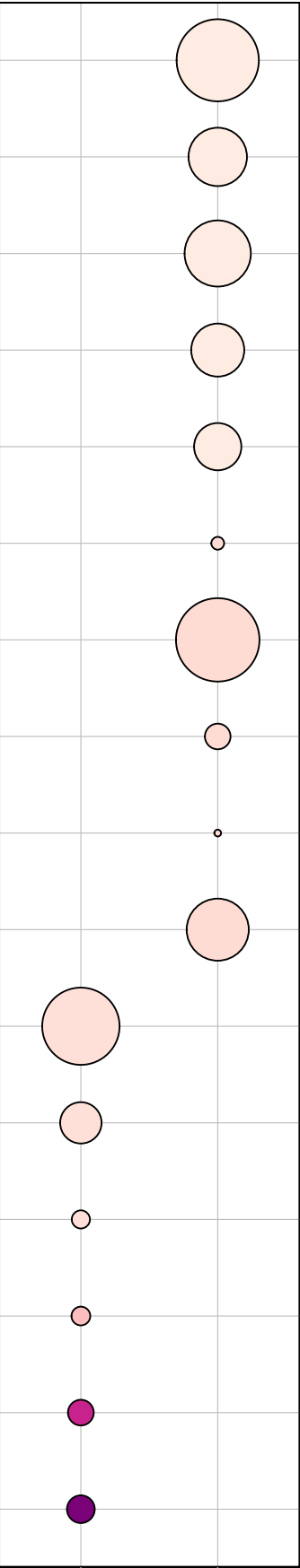


Nvec_vc1.1_XM_001641783.3
mTERF.HG1.0:MTERF4; MTERF4



Nvec_vc1.1_XM_032374380.2

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,



ARCH292 | XM_032364851.2 | zf-C2H2.HG1.32:like:AC008758.1/AC00

ARCH292 | XM_032362875.2 | zf-C2H2.HG14.4:PRDM6

ARCH292 | XM_032361814.2 | zf-C2H2.HG1.35:FEZF1/FEZF2/GFI1/GFI

ARCH292 | XM_001641129.3 | ZNF845

ARCH292 | v1g19278 | Klf13

ARCH288 | XM_001633901.3 | T-box.HG1.11:TBX2/TBX3

ARCH1003 | XM_048731064.1 | zf-C2H2.HG87.0:NA

ARCH1003 | XM_048726174.1 | zf-C2H2.HG6.2:MYT1/MYT1L/ST18

ARCH1003 | XM_032384134.2 | zf-C2H2.HG3.3:ZNF641

ARCH1003 | XM_032364814.2 | zf-C2H2.HG1.35:FEZF1/FEZF2/GFI1/GFI

ARCH673 | XM_032385706.2 | FoxQ2d

ARCH652 | XM_032384126.2 | Forkhead.HG1.17:like:FOXI1/FOXI2/FO

ARCH652 | XM_001627072.3 | Forkhead.HG1.17:like:FOXI1/FOXI2/FO

ARCH372 | XM_048732912.1 | TTF1

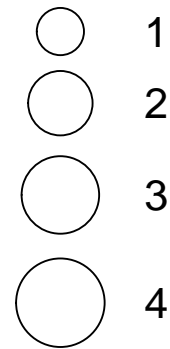
ARCH997 | XM_032364773.2 | NOTO

ARCH1457 | XM_001635374.3 | Homeodomains.HG3.7:PBX1/PBX2/PBX3/P

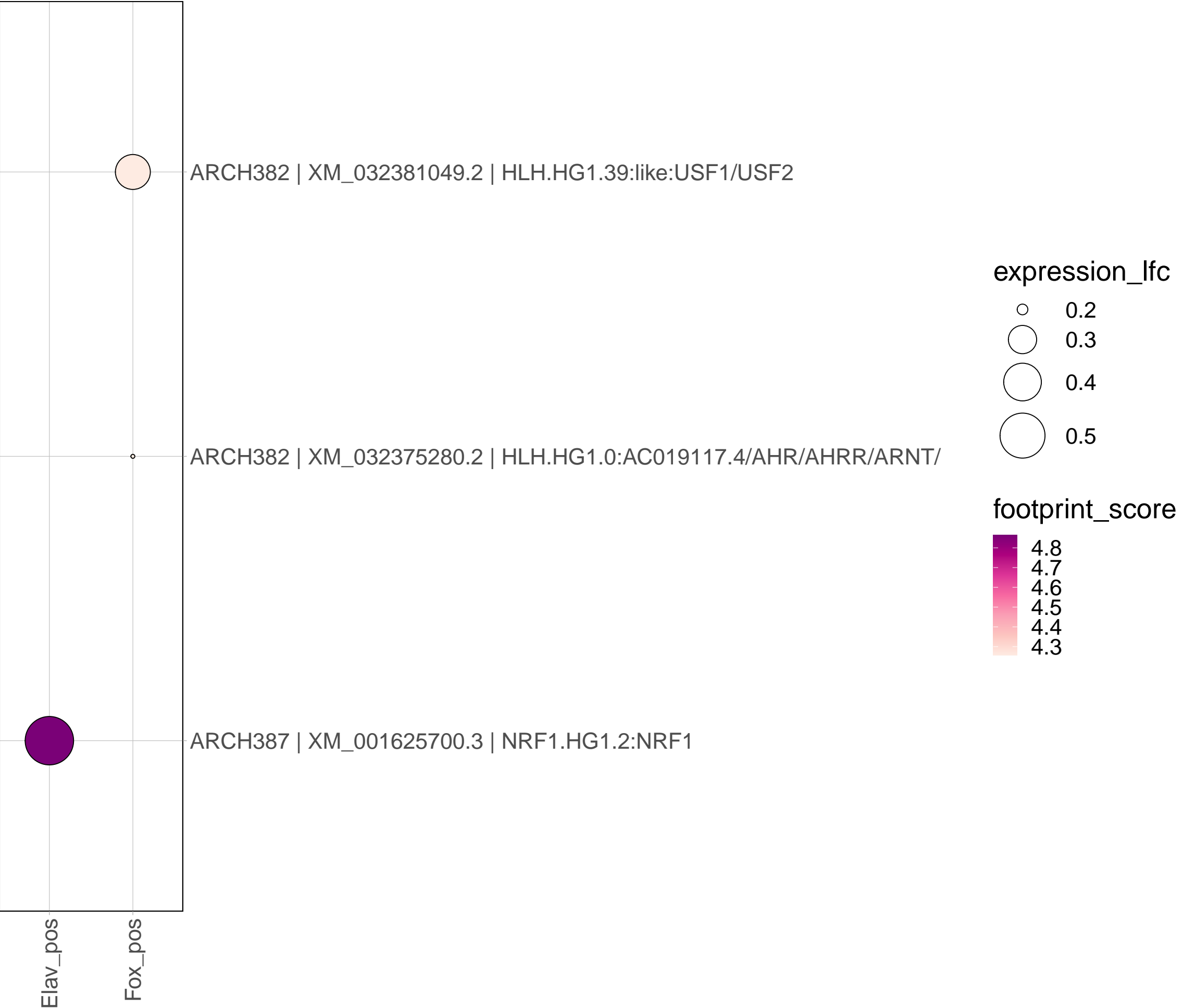
footprint_score



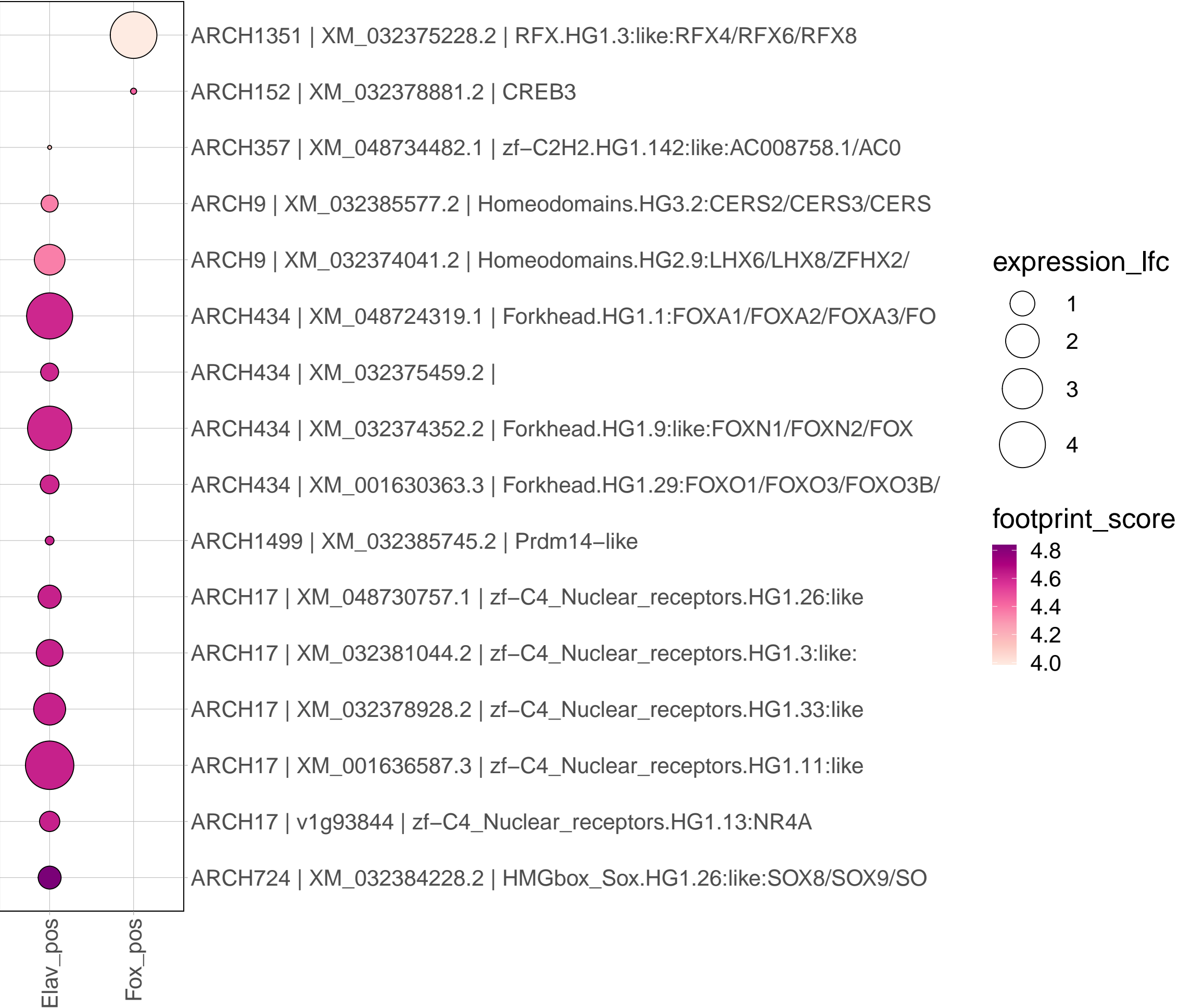
expression_lfc



Nvec_vc1.1_XM_032384990.2
Protein odr-4 homolog; ODR4-like/ODR4-like/



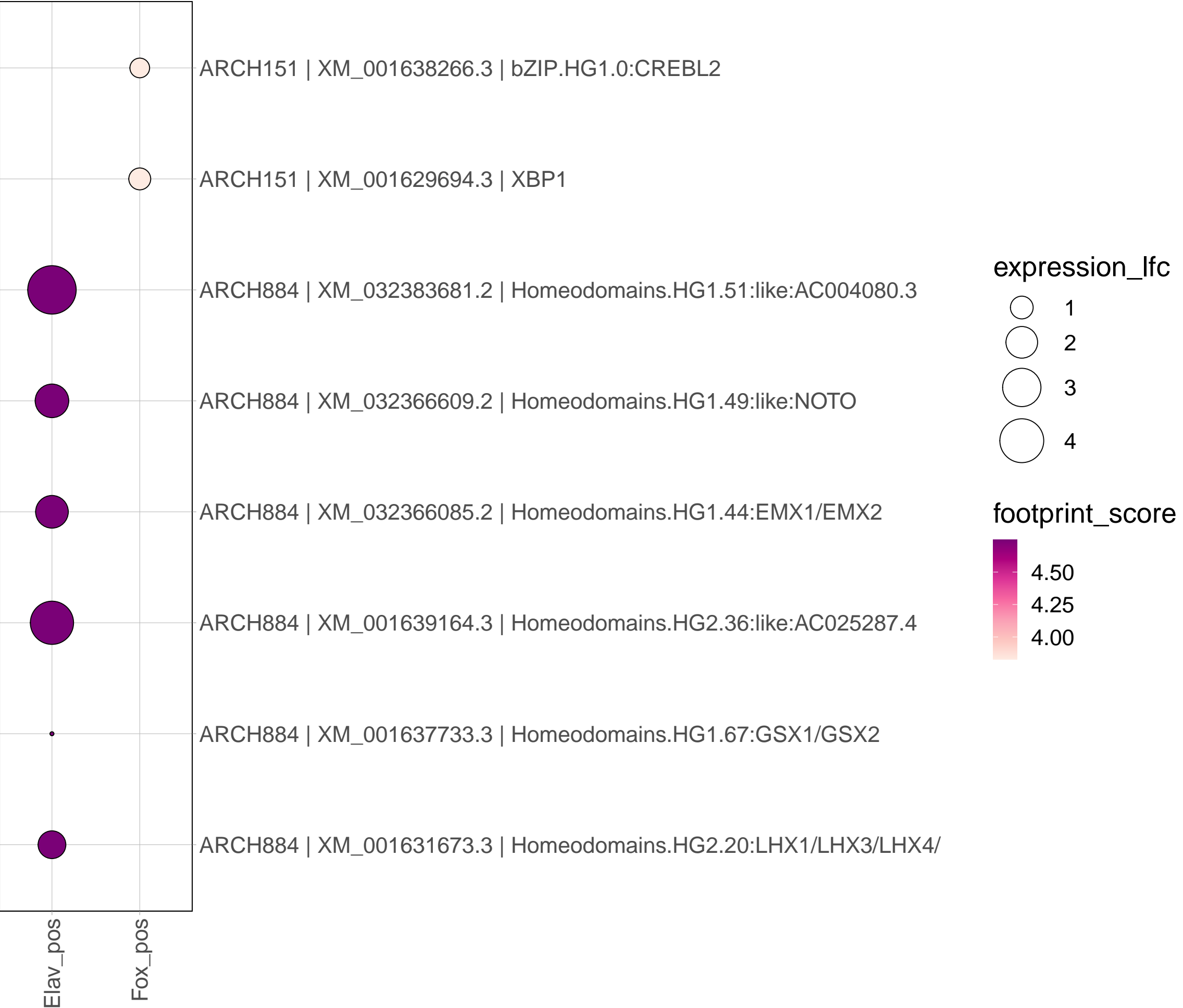
Nvec_vc1.1_XM_048729479.1
TAF5-like RNA polymerase II p300/CB; TFIID_90kDa/WD40/WD40/WD40/WD40



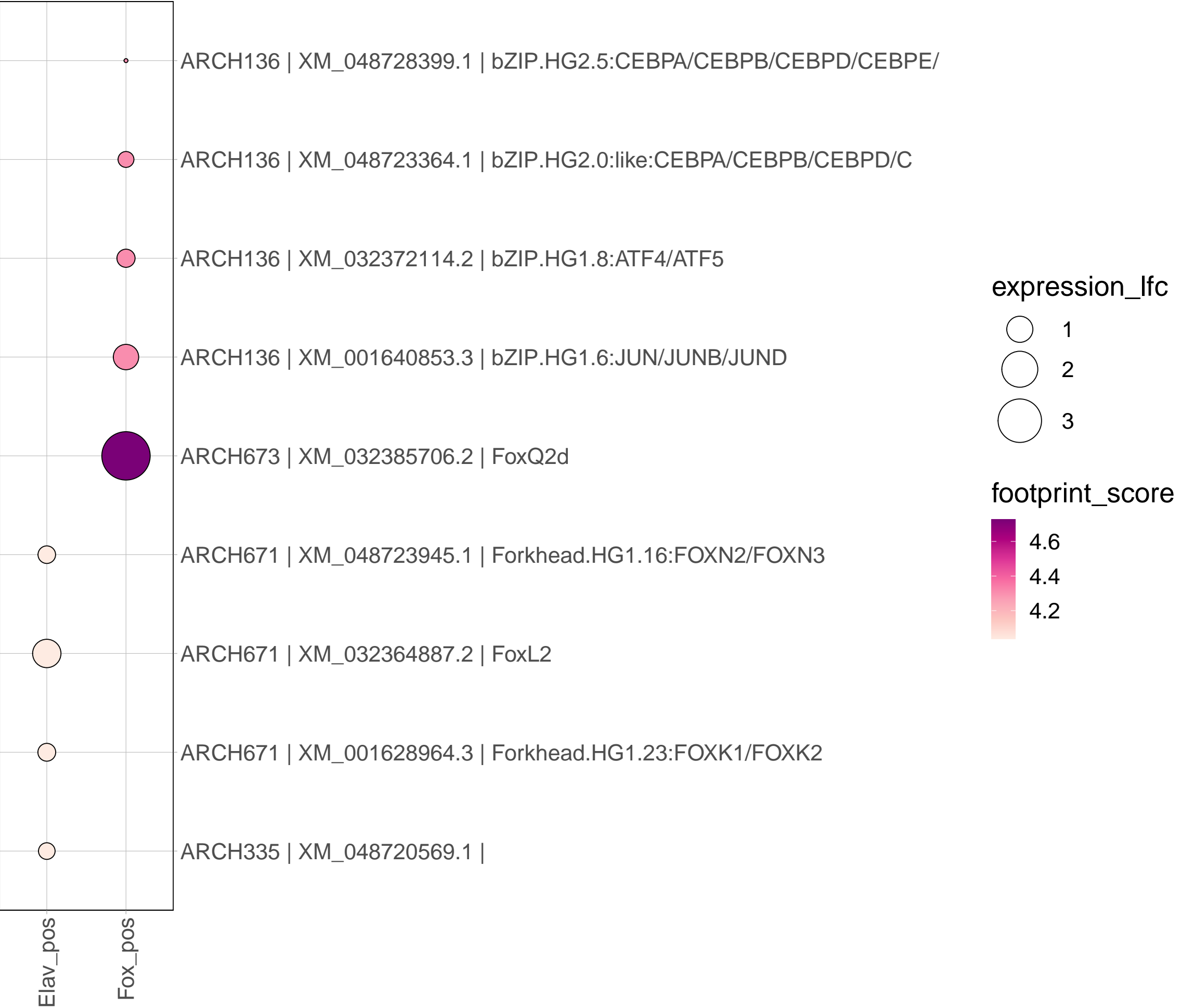
Nvec_vc1.1_XM_001639817.3
Ephrin:AL691442/EFNA1/EFNA2/EFNA3/E; Ephrin.HG1.0:AL691442/EFNA1/EF



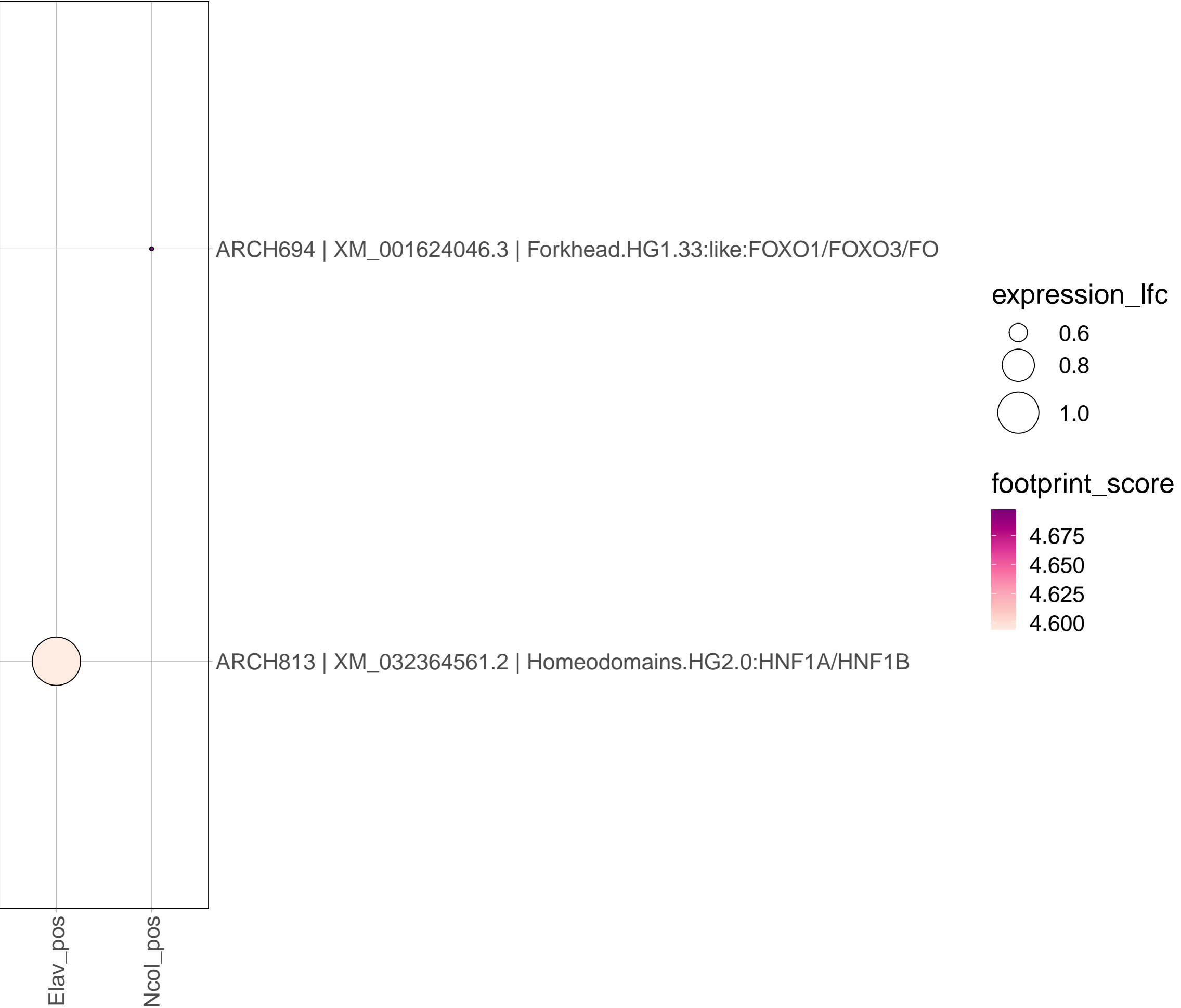
Nvec_vc1.1_XM_032366492.2
N-acetylated-alpha-linked acidic di; TFR_dimer/



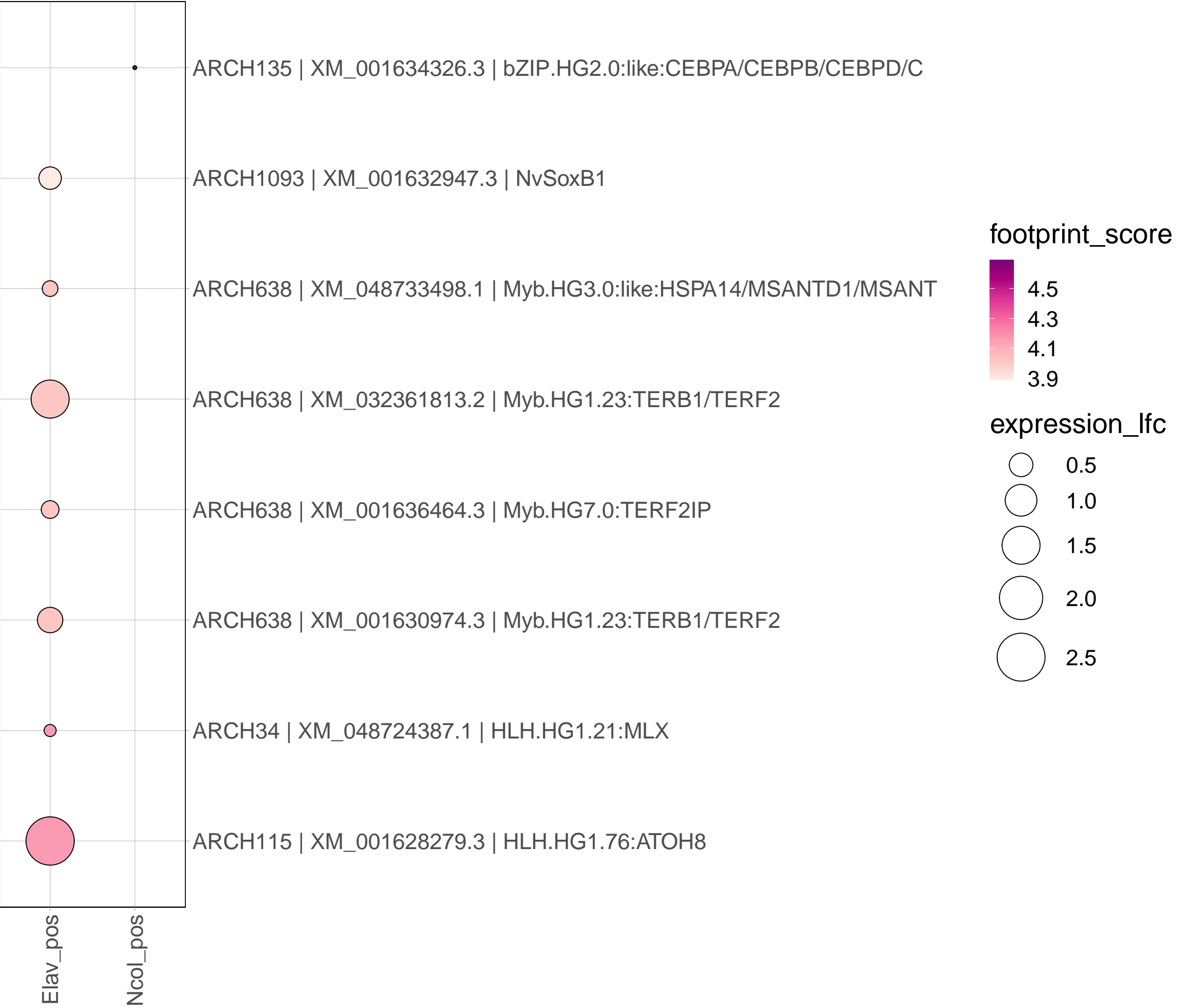
Nvec_vc1.1_XM_048729868.1
E3 ubiquitin–protein ligase UBR2;



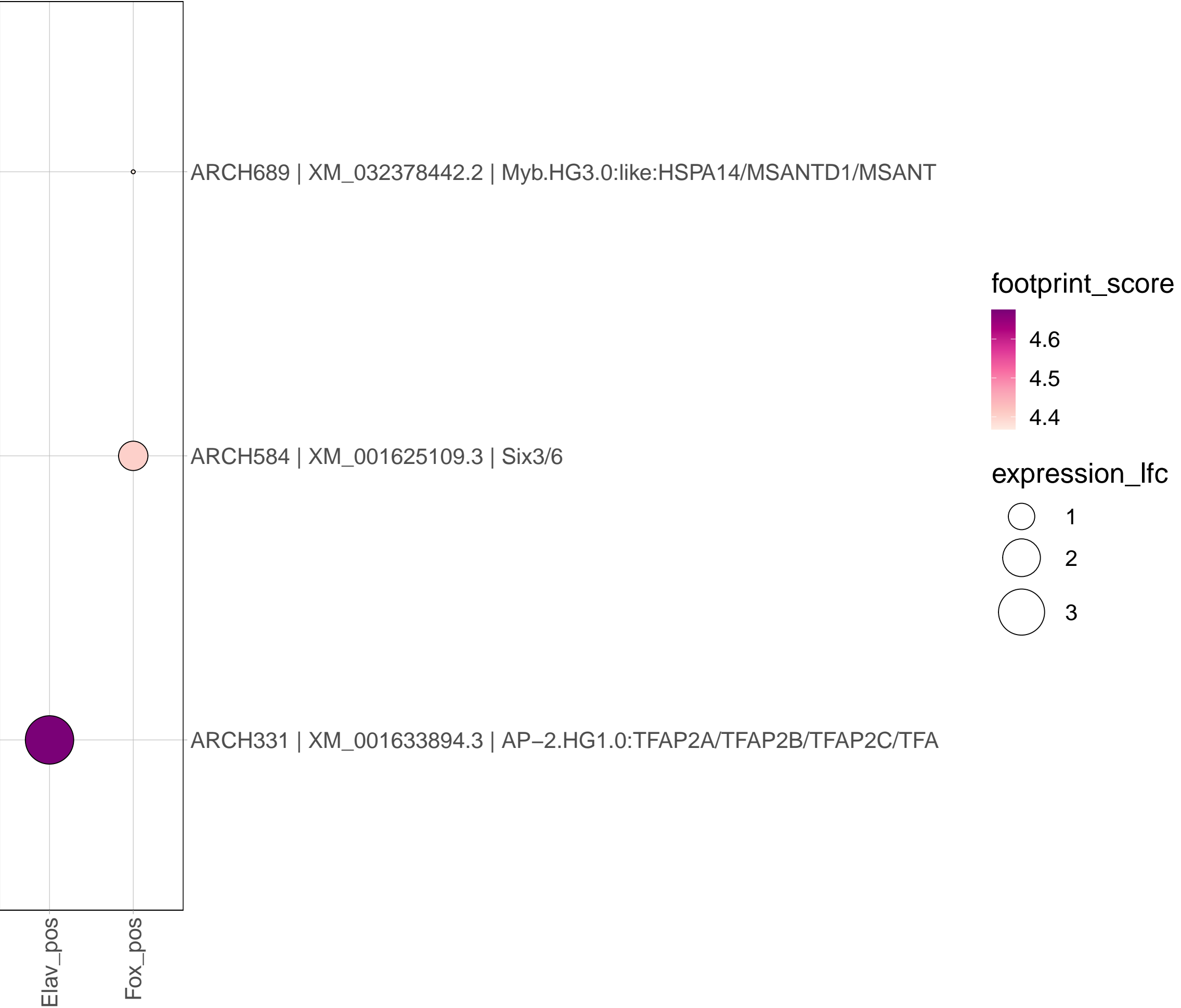
Nvec_vc1.1_XM_048734712.1
Very long-chain-specific acyl-CoA d;



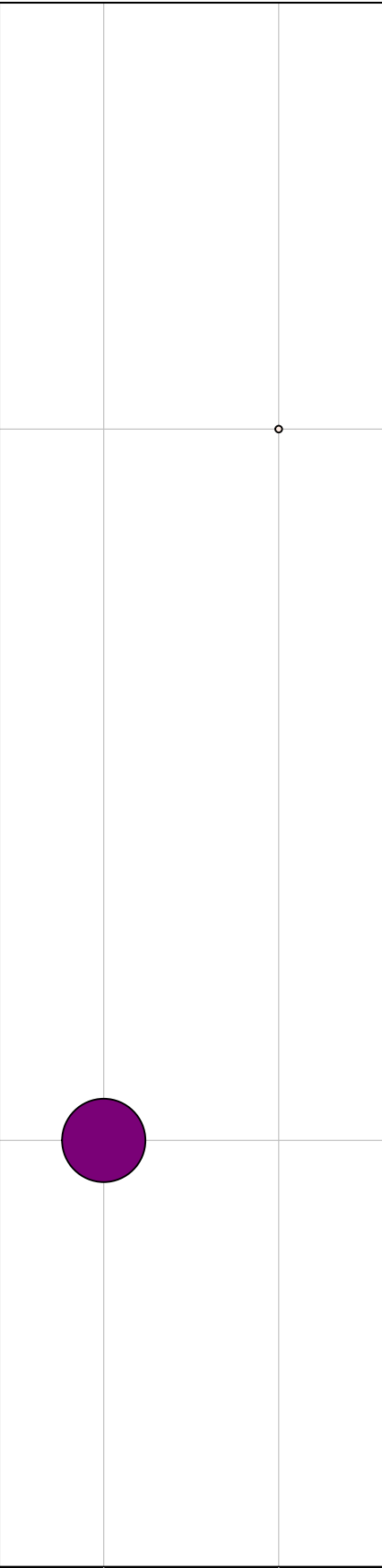
Nvec_vc1.1_XM_001628363.3
cDNA, FLJ94698, highly similar to H; Sybindin/



Nvec_vc1.1_XM_032387176.2
LMBR1 domain-containing protein 2; LMBR1/



Nvec_vc1.1_XM_001631292.3
ADP-ribosylation factor related pro;



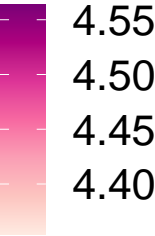
Elav_pos

Ncol_pos

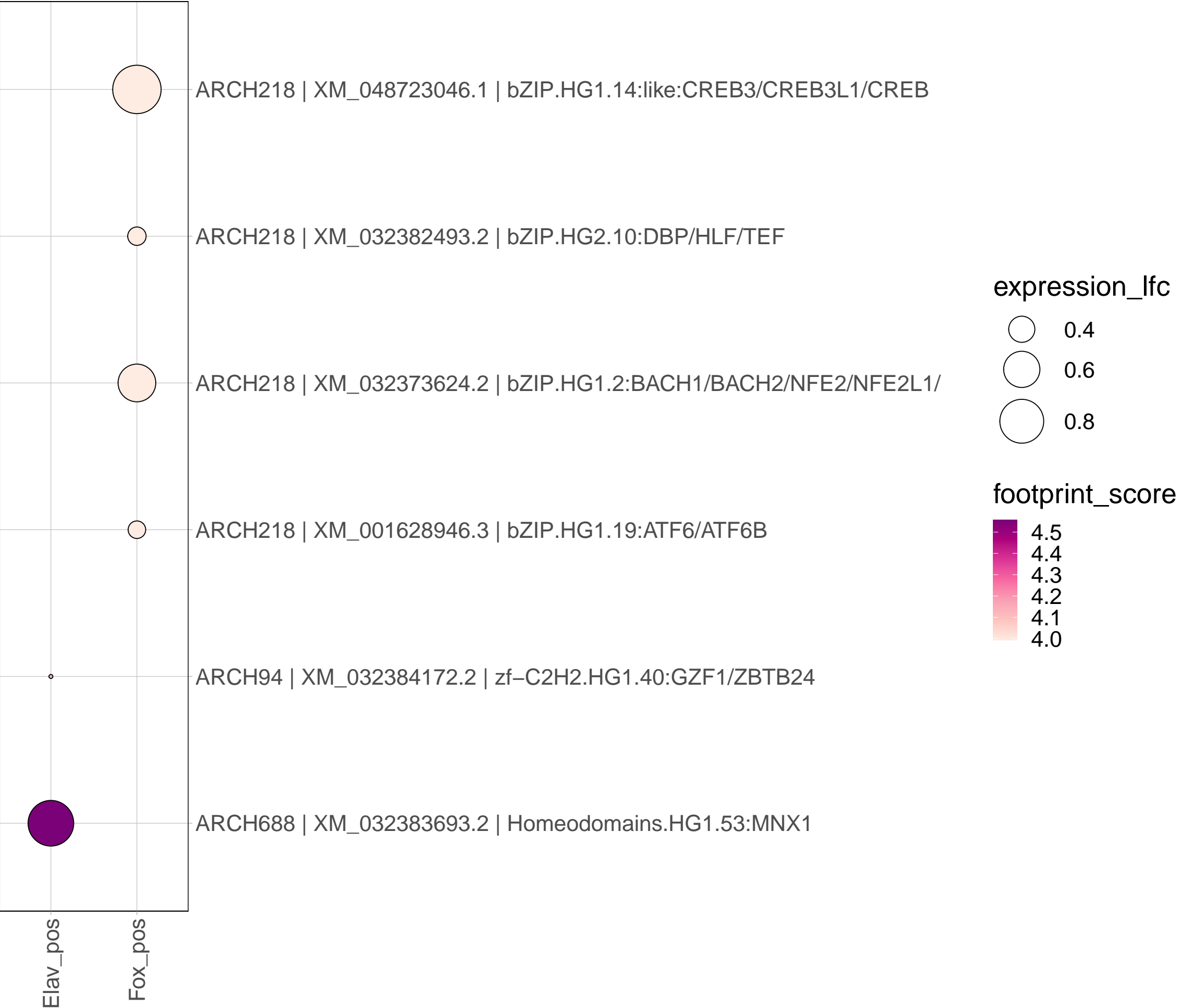
expression_lfc



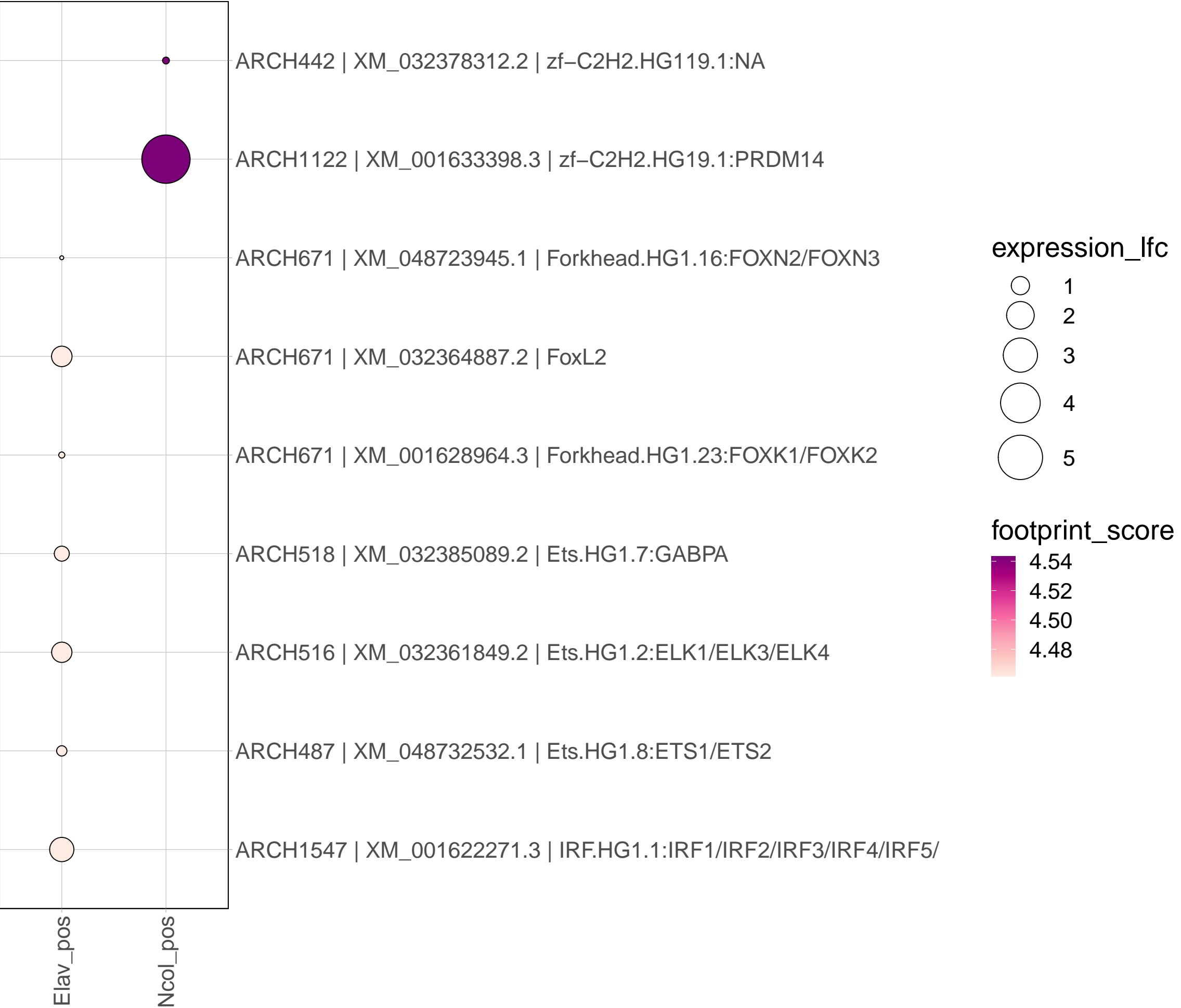
footprint_score



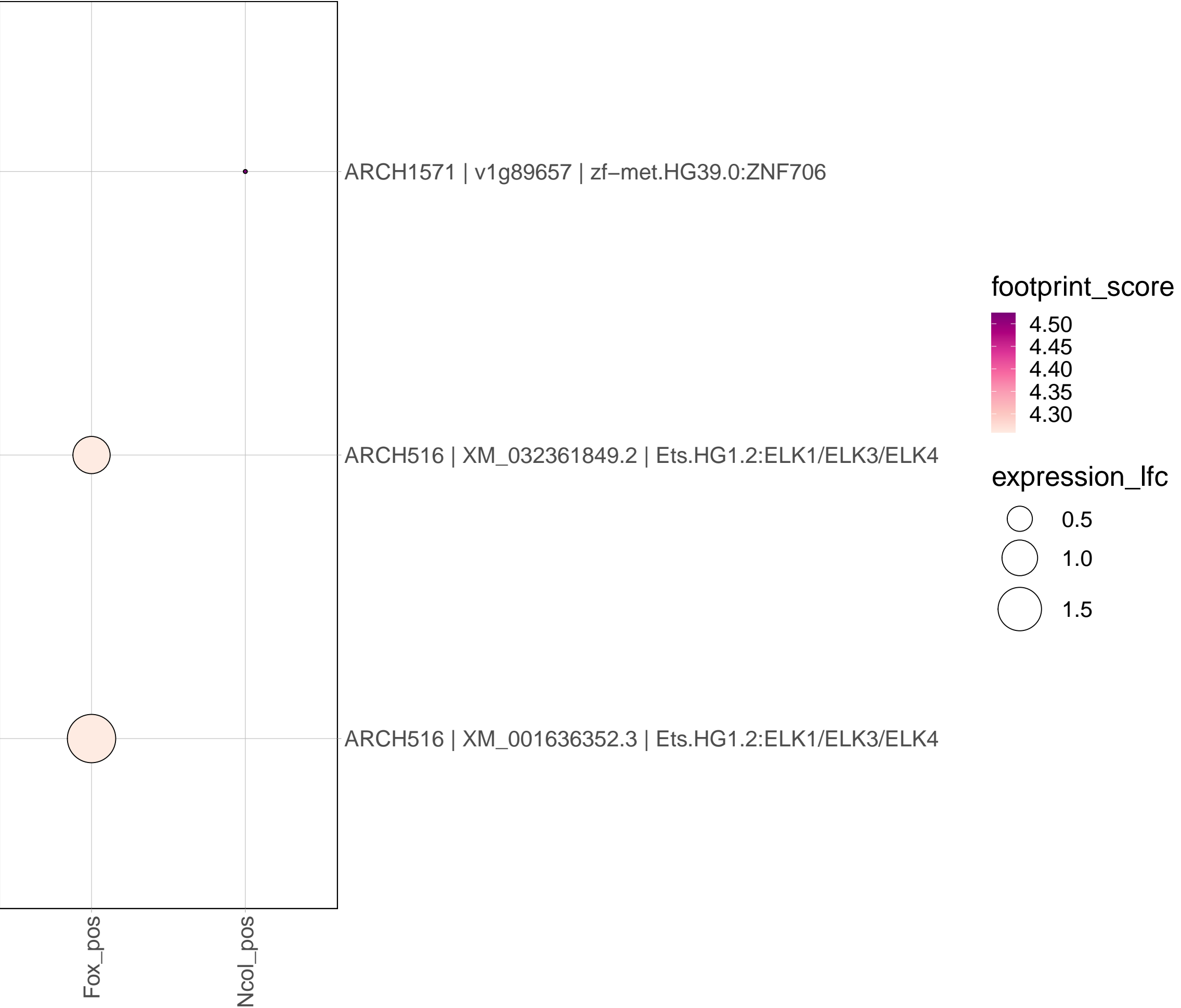
Nvec_vc1.1_XM_001631286.2
NNMT protein; NNMT_PNMT_TEMT/



Nvec_vc1.1_XM_001624997.3
; F-box-like/



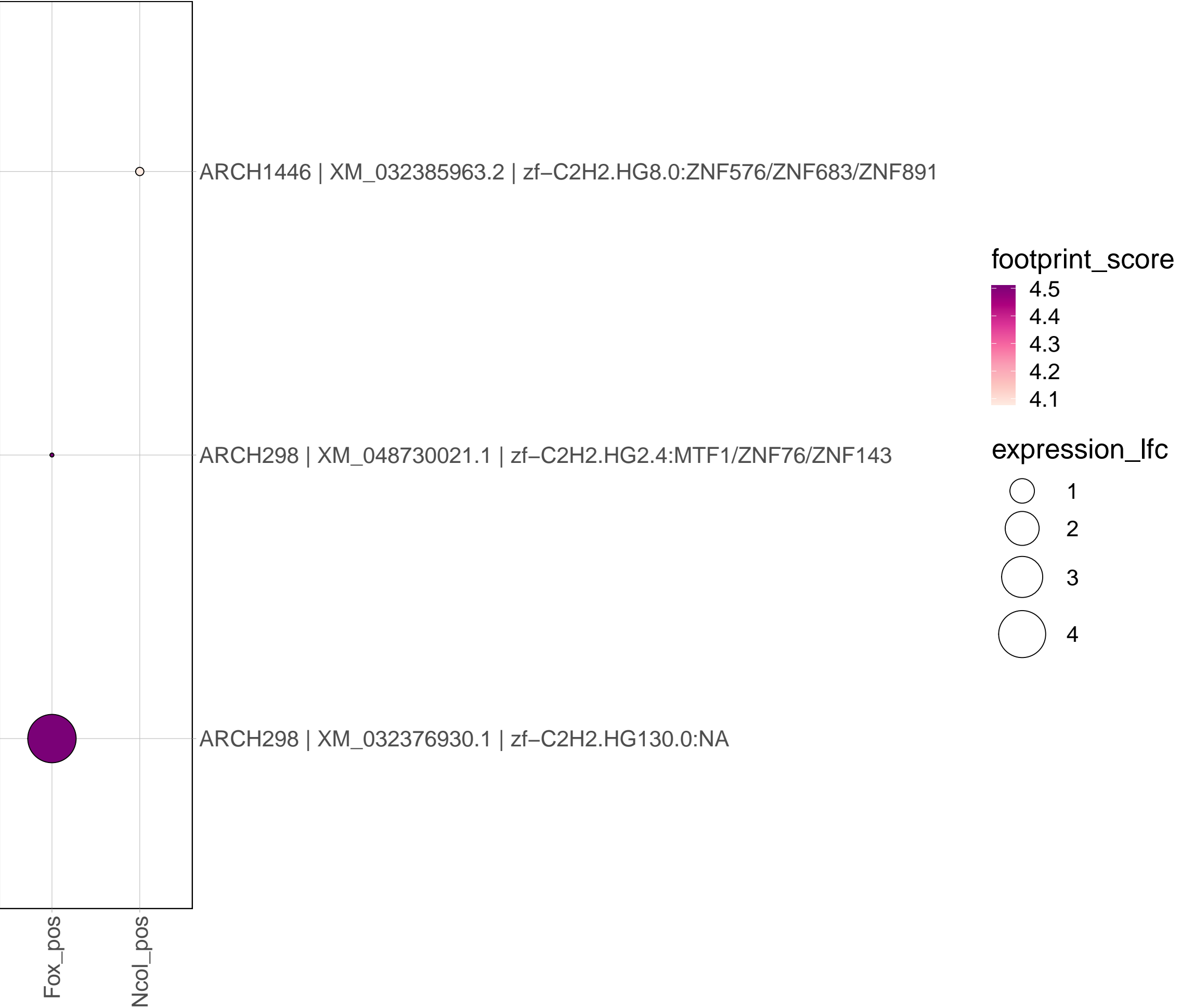
Nvec_vc1.1_XM_001641286.3
cDNA FLJ51917, highly similar to Me; TIMP/



Nvec_vc1.1_XM_032368559.2
Protein prenyltransferase alpha sub; PPTA/



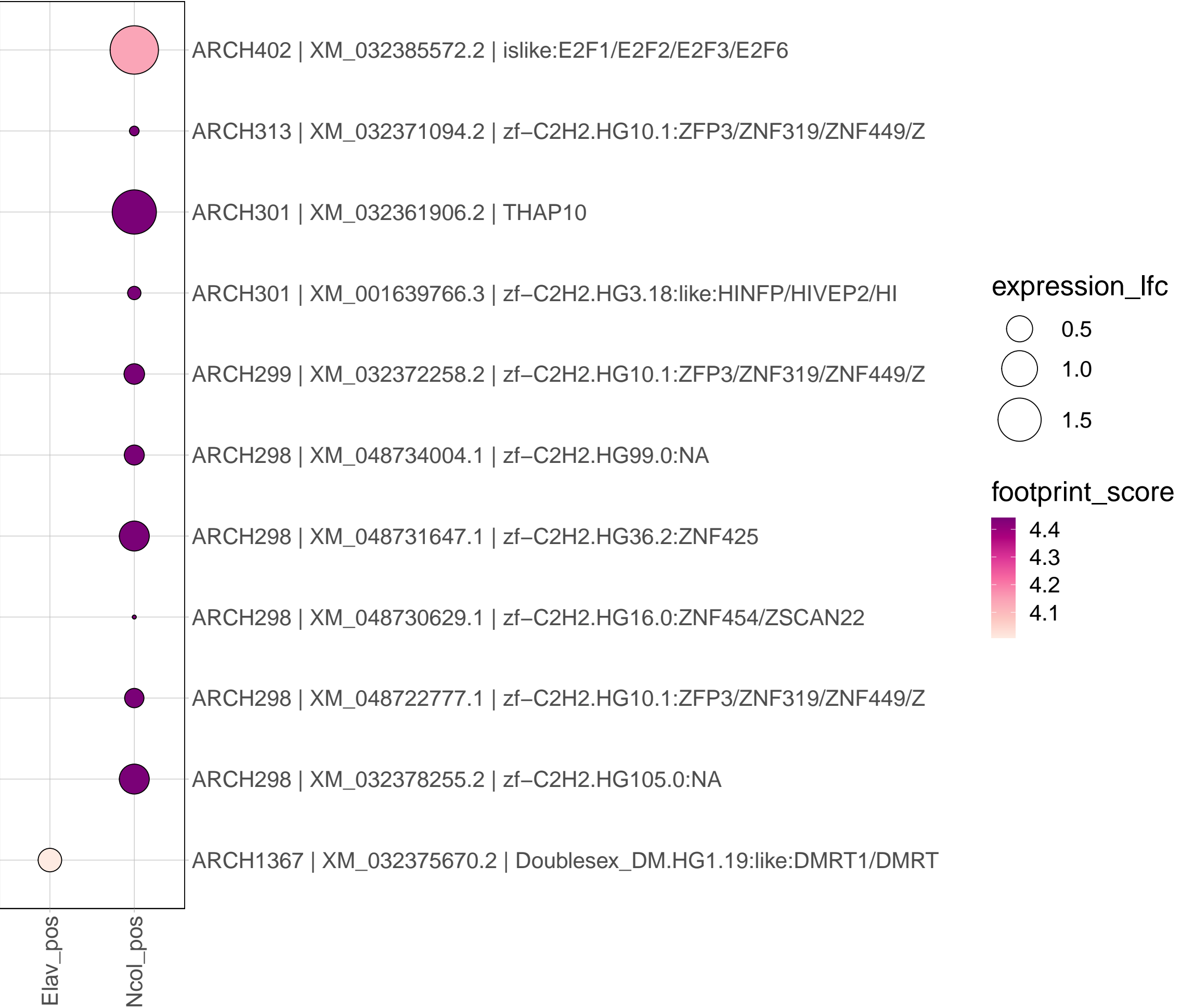
Nvec_vc1.1_XM_048727242.1
Zinc transporter 7; Cation_efflux/



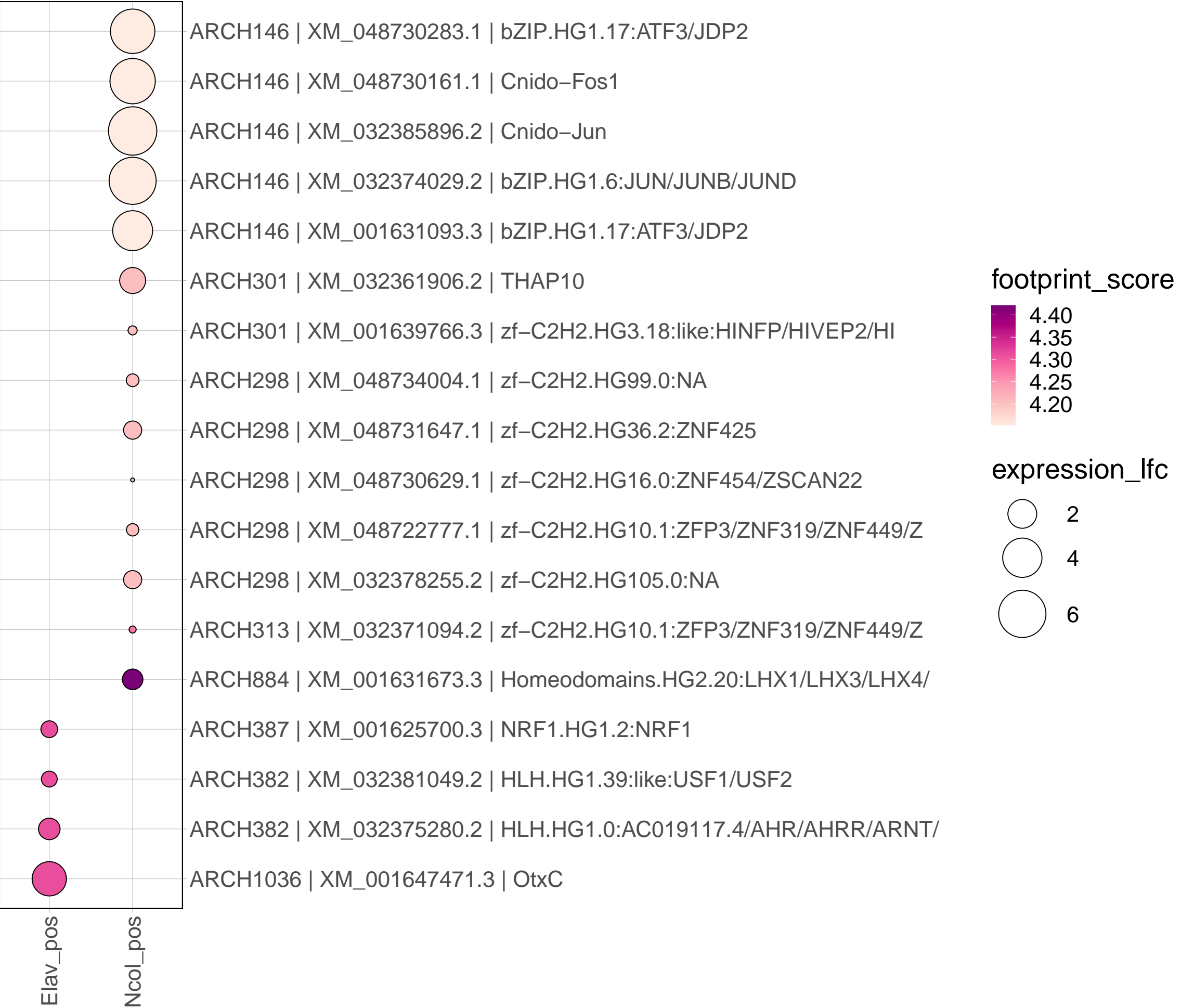
Nvec_vc1.1_XM_032372191.2
Abhydrolase domain containing 6; Abhydrolase_6/



Nvec_vc1.1_XM_048730264.1
Component of oligomeric golgi compl; COG2/DUF3510/

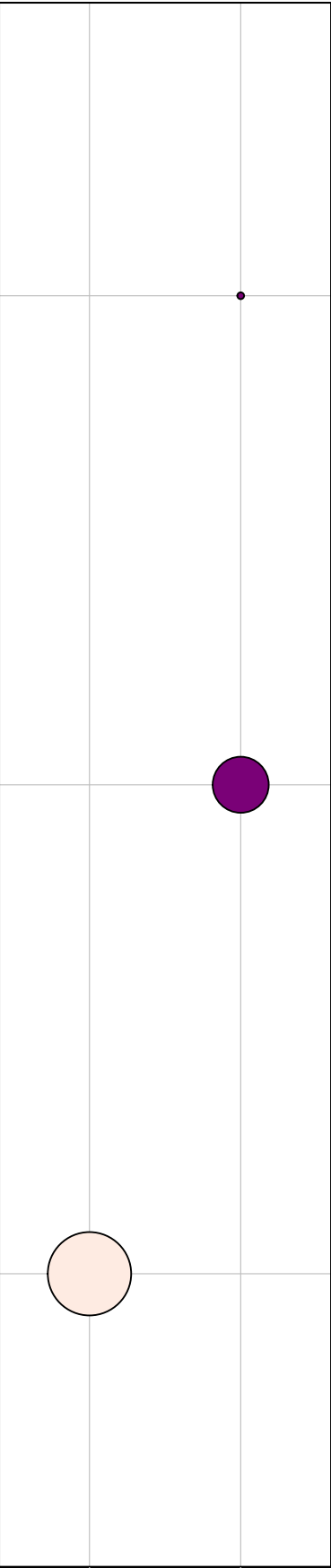


Nvec_vc1.1_XM_032375751.2
Transmembrane protein 115; DUF1751/



Nvec_vc1.1_XM_048725016.1

;



Fox_pos

Ncol_pos

expression_lfc



0.4



0.6



0.8

footprint_score



4.35

4.30

4.25

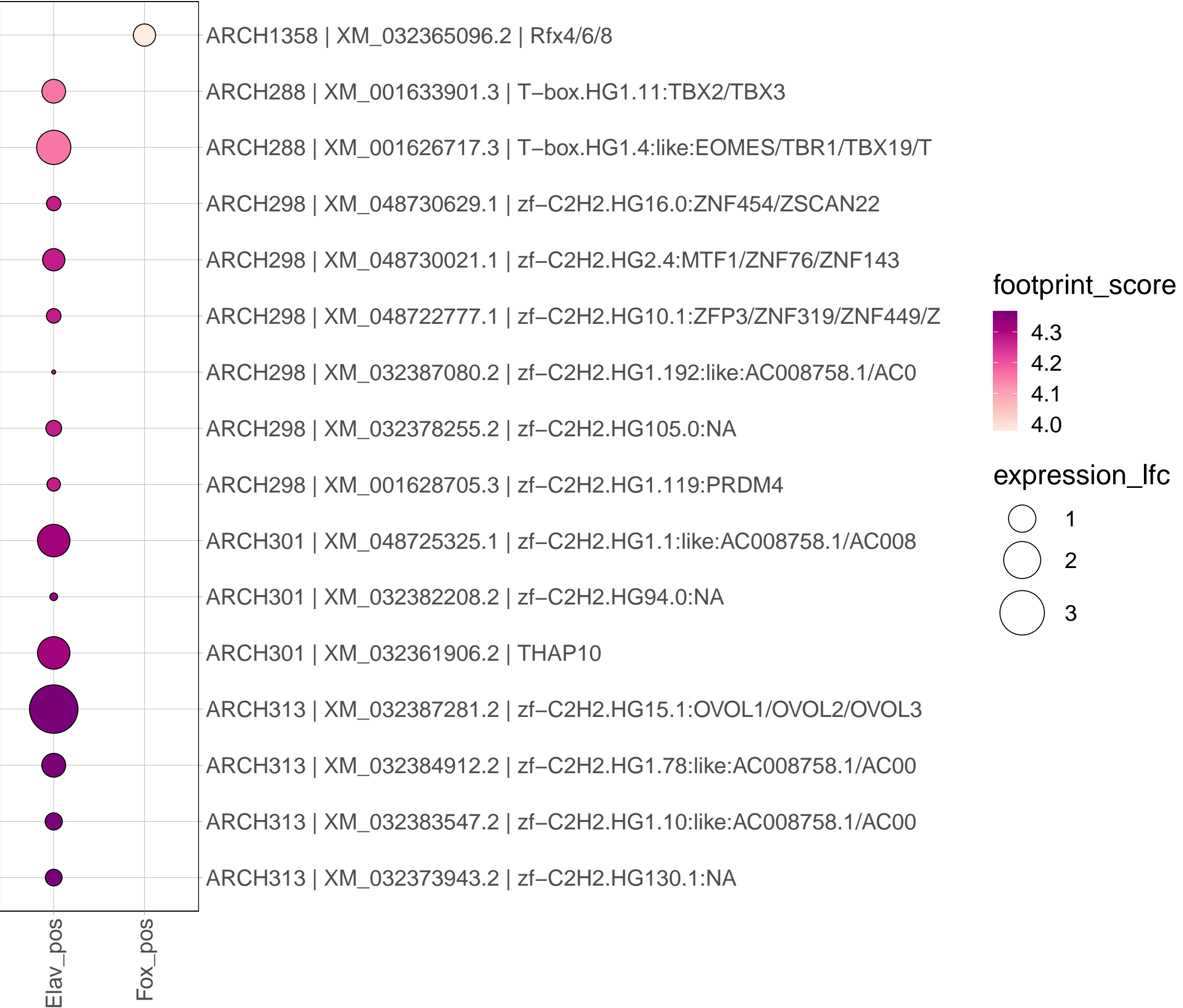
4.20

ARCH372 | XM_048732912.1 | TTF1

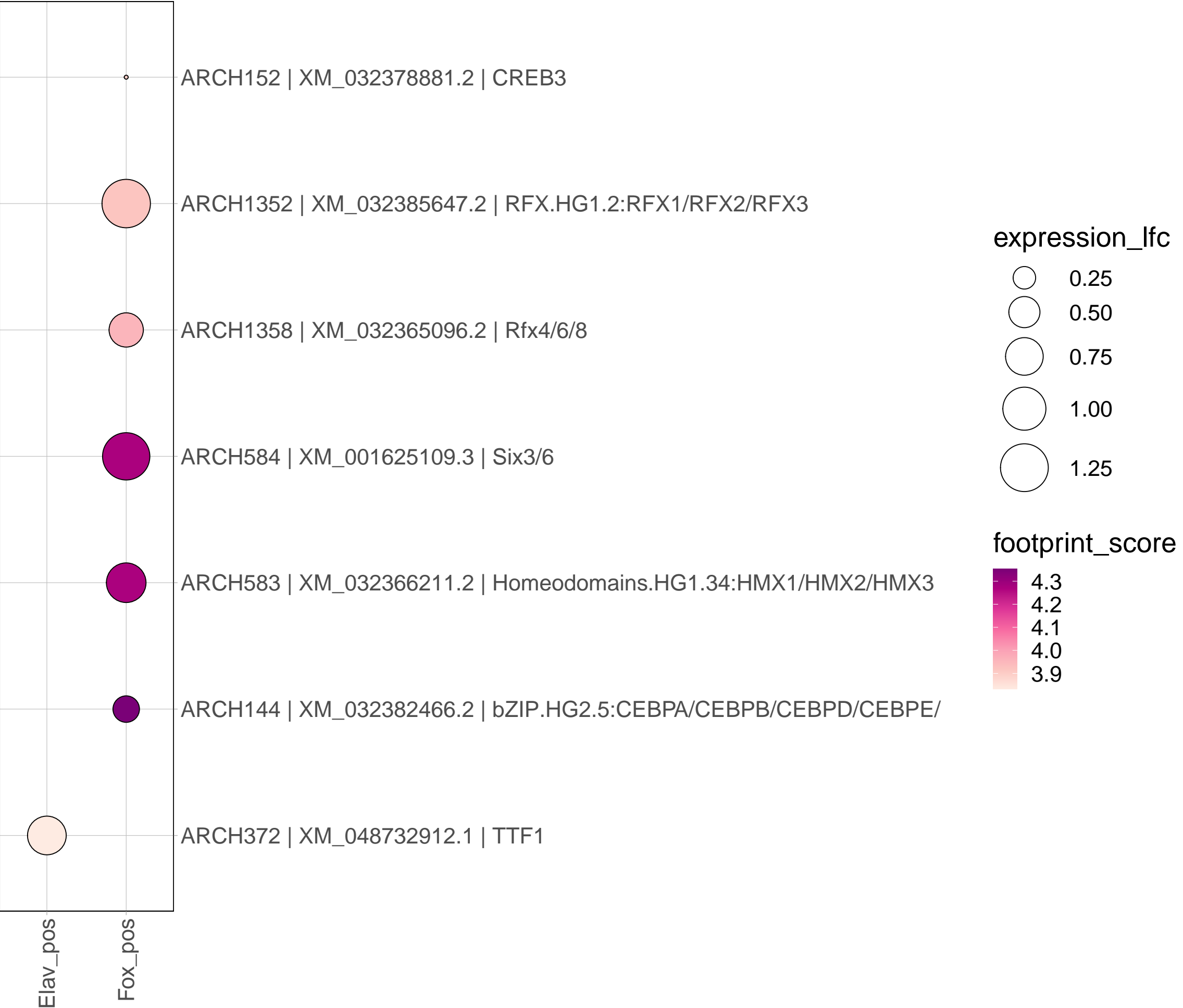
ARCH372 | XM_001625437.3 | Myb.HG2.3:NA

ARCH998 | XM_032376608.2 | Homeodomains.HG3.20:MEIS1/MEIS2/MEI

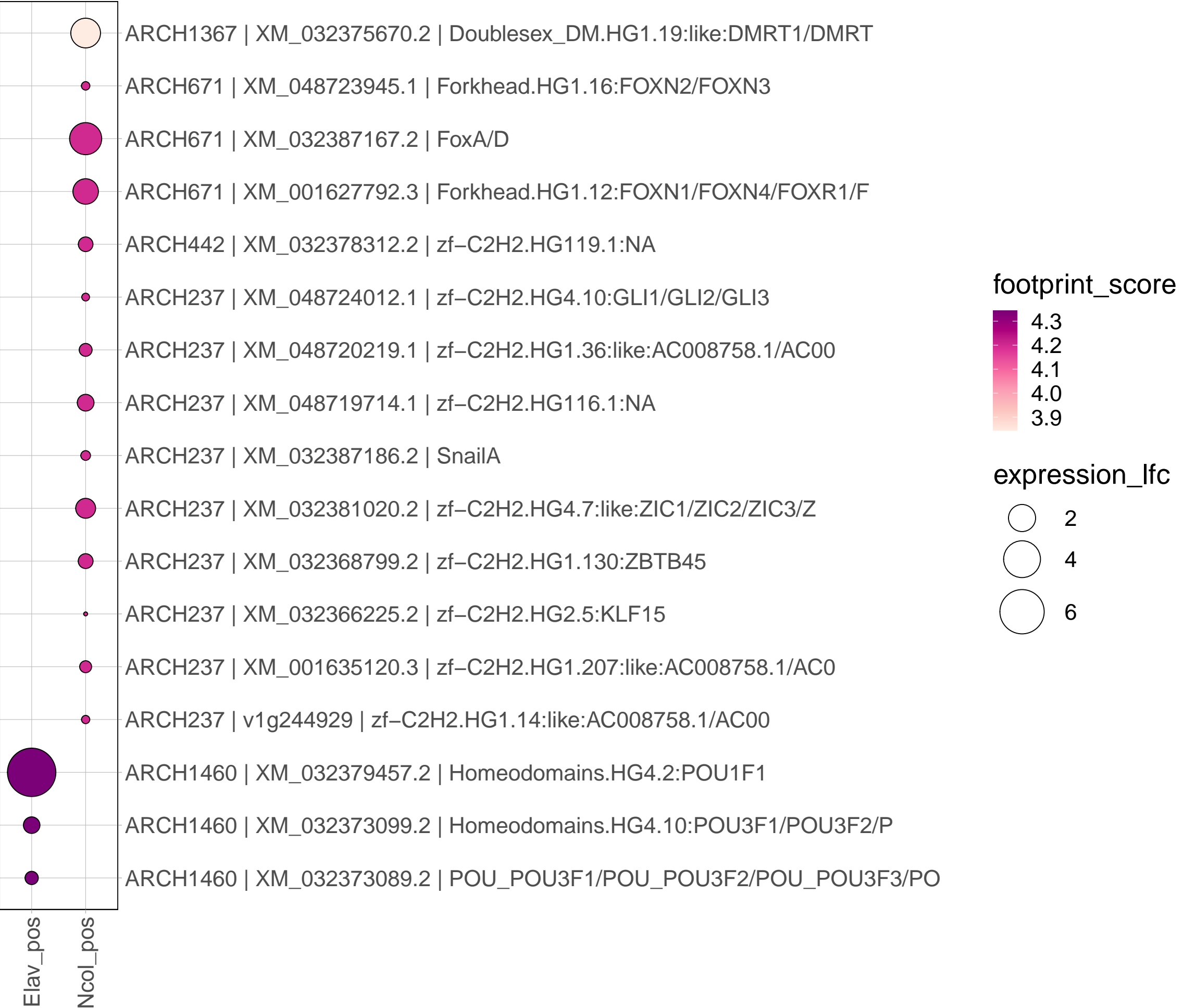
Nvec_vc1.1_XM_001638793.3
Zinc finger CCHC domain-containing ; PSP/



Nvec_vc1.1_XM_032363581.2
GPCR–rhodopsin:NA; 7tm_1/

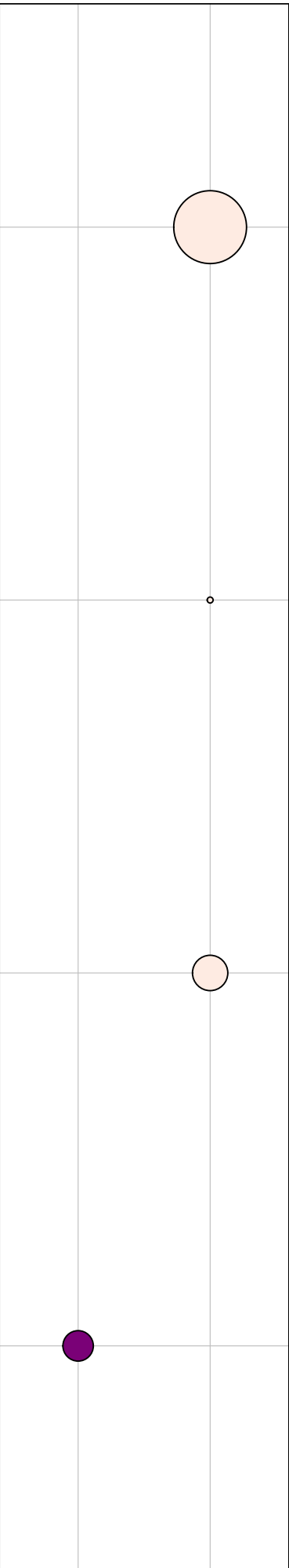


Nvec_vc1.1_XM_032364770.2
GPCR–rhodopsin:NA; 7tm_1/



Nvec_vc1.1_XM_032367592.2
cDNA, FLJ79298, highly similar to P;





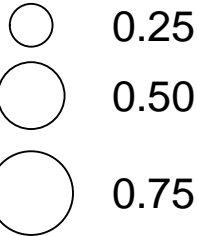
ARCH884 | XM_001631673.3 | Homeodomains.HG2.20:LHX1/LHX3/LHX4/

ARCH12 | XM_032366646.2 | ARID_BRIGHT.HG1.5:ARID5A/ARID5B

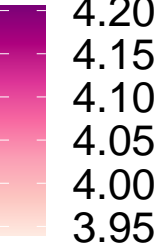
ARCH12 | XM_032362135.2 | ARID_BRIGHT.HG1.9:ARID3A/ARID3B/ARI

ARCH94 | XM_032384172.2 | zf-C2H2.HG1.40:GZF1/ZBTB24

expression_lfc

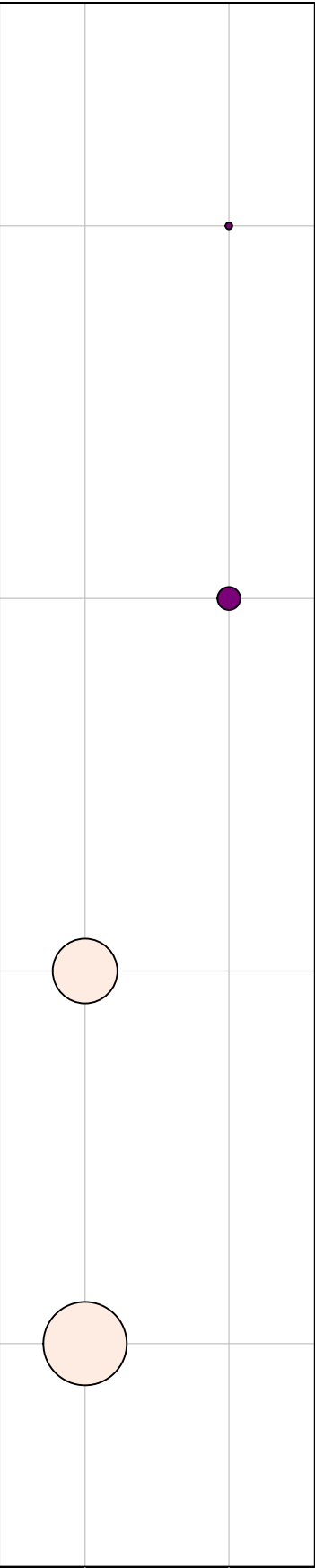


footprint_score



Nvec_v1g243589

;



Fox_pos

Ncol_pos

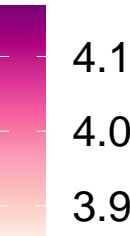
expression_lfc

0.5

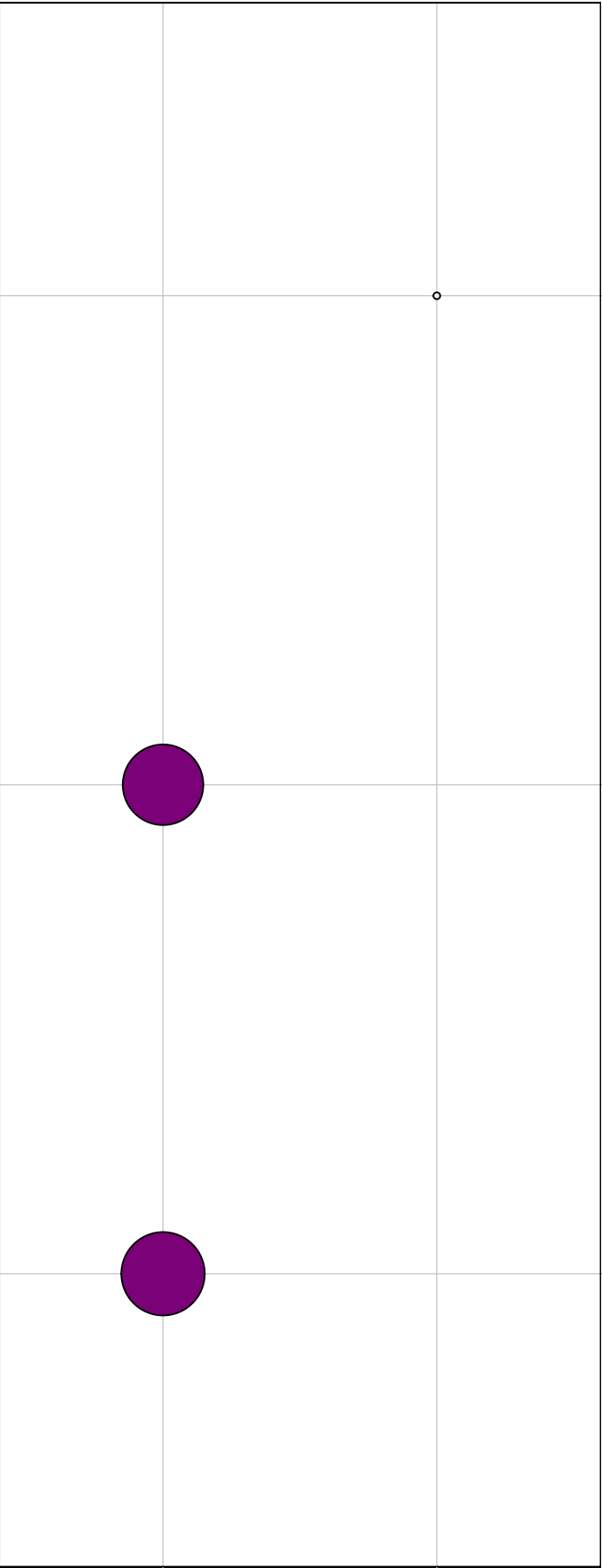
1.0

1.5

footprint_score



Nvec_vc1.1_XM_032385089.2
Ets.HG1.7:GABPA; GABPA

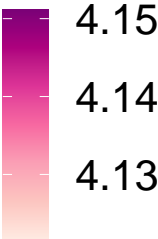


ARCH518 | XM_032385089.2 | Ets.HG1.7:GABPA

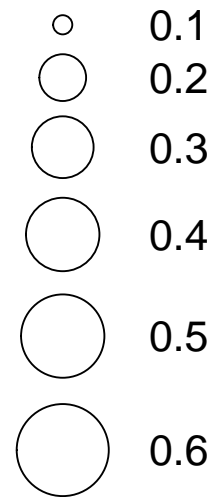
ARCH271 | XM_048732404.1 | MBD.HG1.2:MBD4/MECP2

ARCH271 | XM_048724172.1 | MBD.HG1.1:BAZ2A/BAZ2B

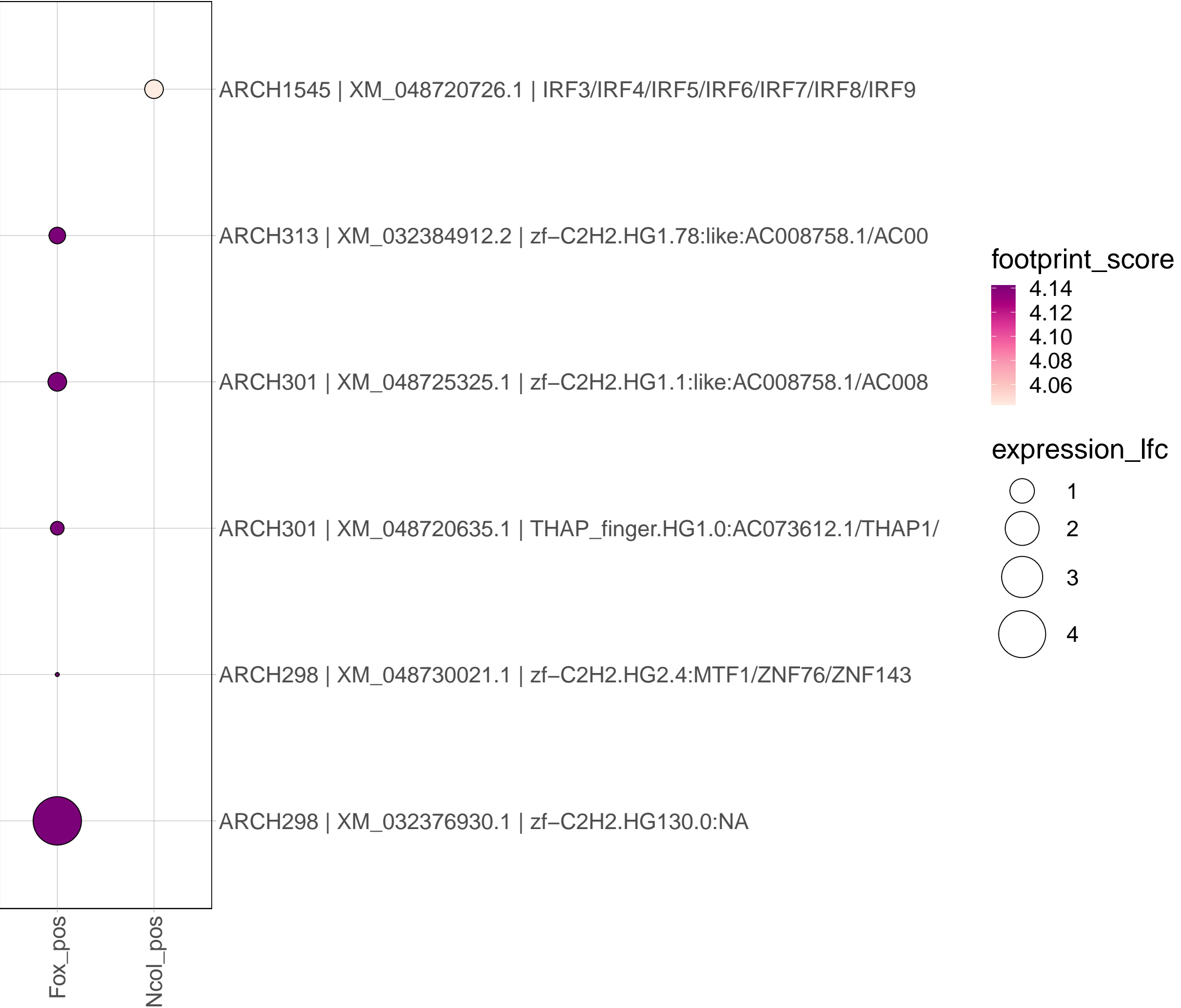
footprint_score



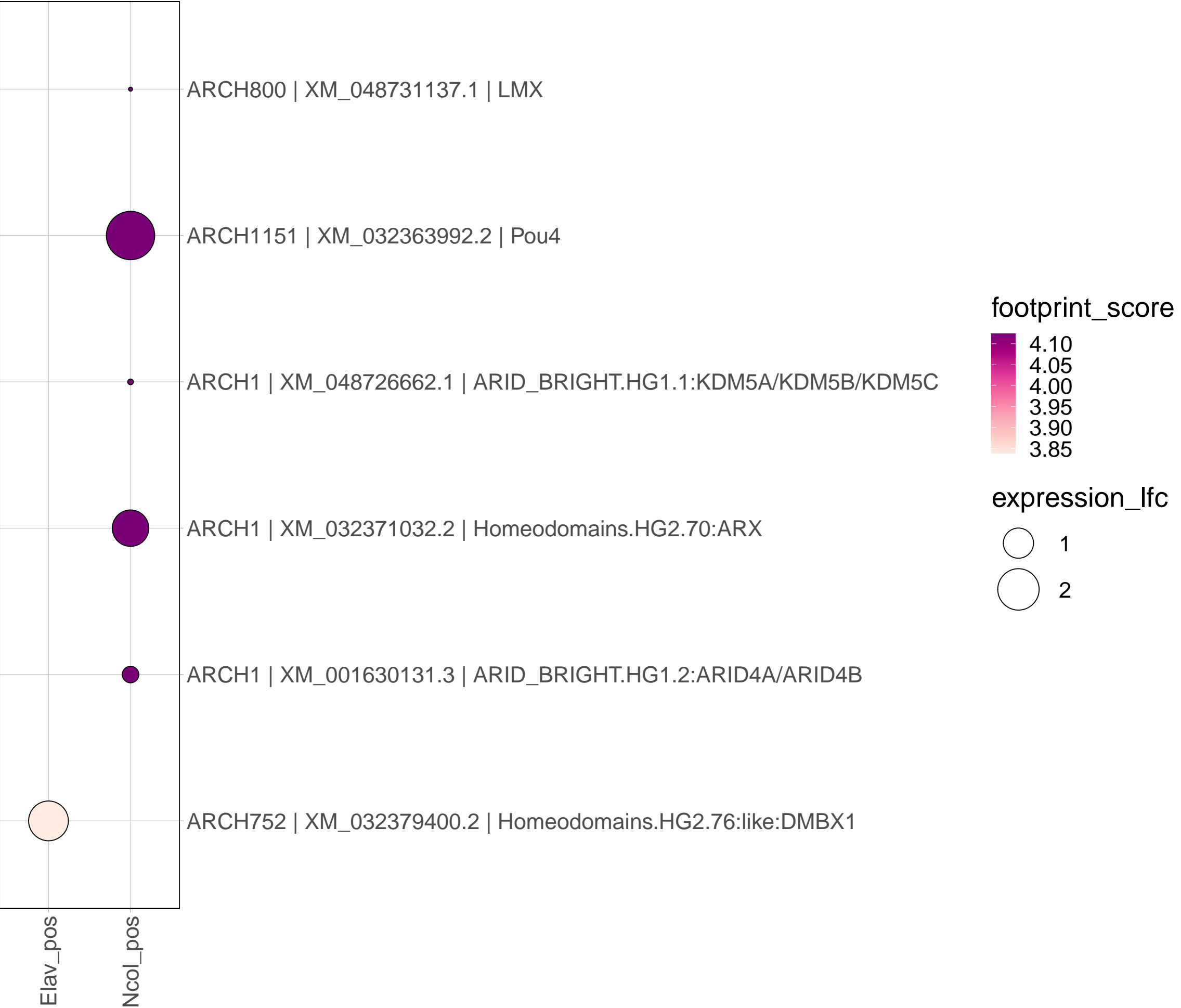
expression_lfc



Nvec_vc1.1_XM_048720705.1
Ubiquitin carboxyl-terminal hydrola;

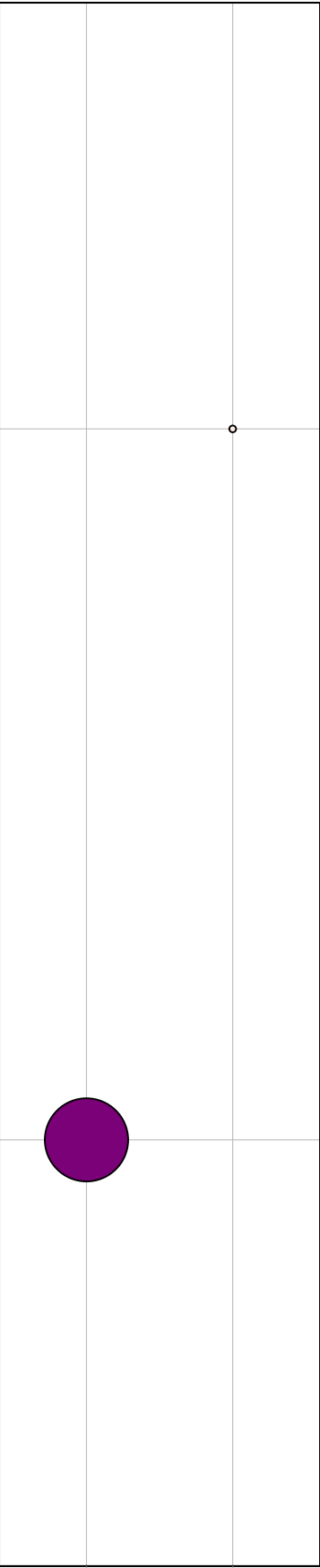


Nvec_vc1.1_XM_032362554.2
Synembryn-A; Ric8/



Nvec_vc1.1_XM_048726963.1

;



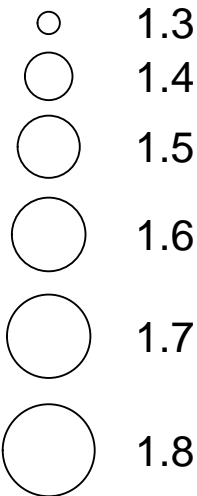
Elav_pos

Fox_pos

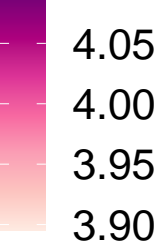
ARCH1352 | XM_032385647.2 | RFX.HG1.2:RFX1/RFX2/RFX3

ARCH1369 | XM_001625588.3 | Runt_Runx.HG1.0:RUNX1/RUNX2/RUNX3

expression_lfc



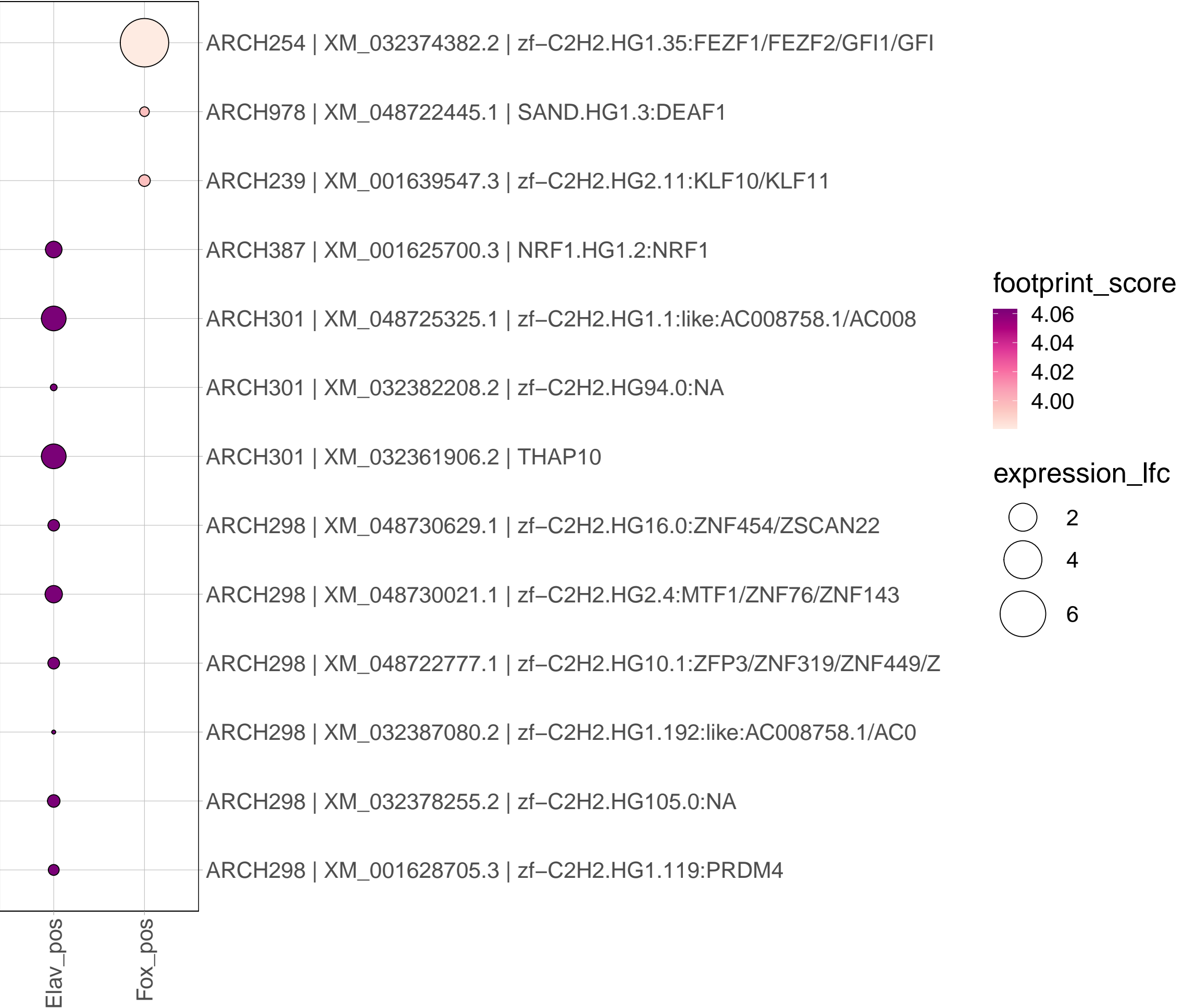
footprint_score



Nvec_vc1.1_XM_032382564.2
Uncharacterized protein; TMEM219/



Nvec_vc1.1_XM_001630938.3
TM2 domain-containing protein 2; TM2/



Nvec_vc1.1_XM_048730070.1
FAM179B protein;

