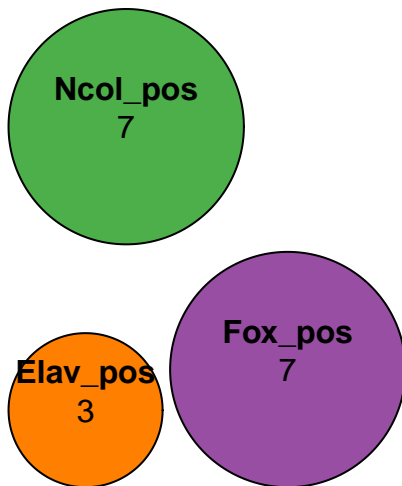
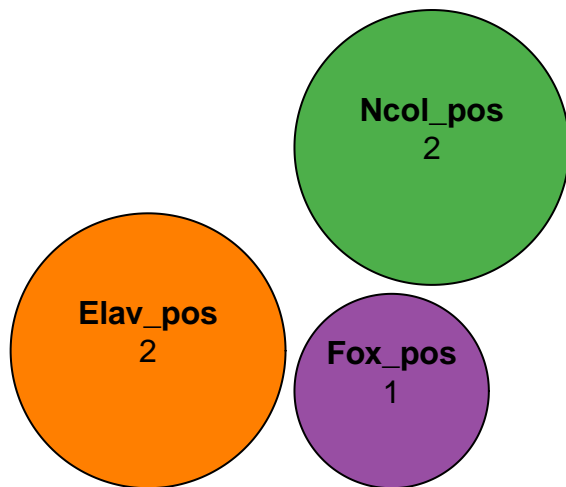


Nvec_vc1.1_XM_032385685.2
GPCR-rhodopsin:like:GPR17; 7tm_1/

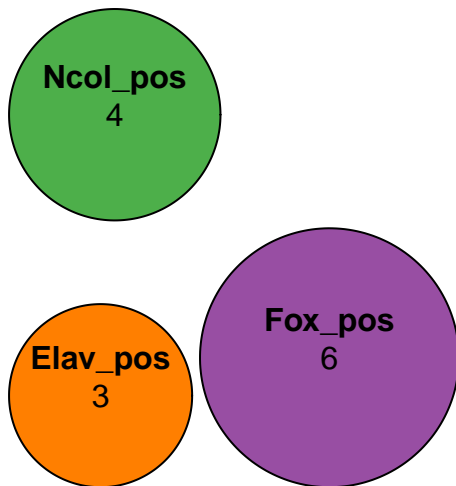


Nvec_vc1.1_XM_032367127.2

bZIP.HG2.10:DBP/HLF/TEF; DBP/HLF/TEF

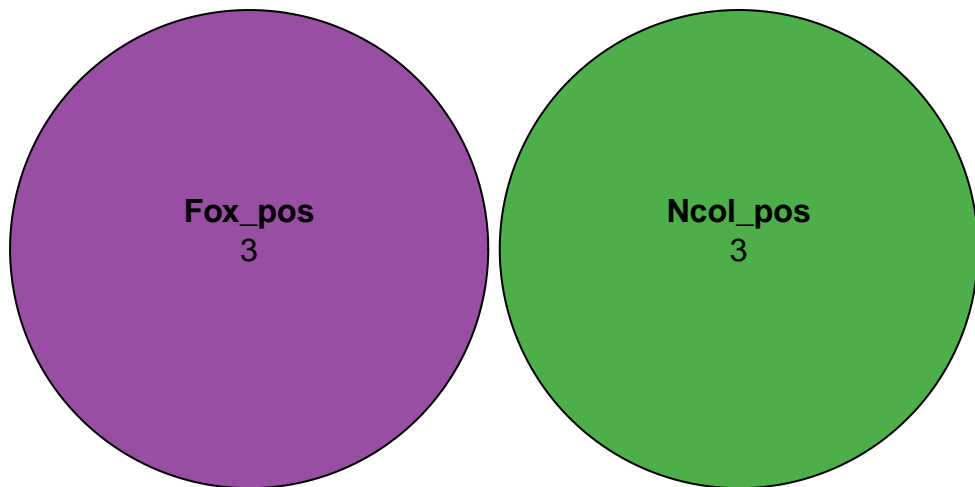


Nvec_vc1.1_XM_048720228.1
Sushi, nidogen and EGF-li; NIDO/



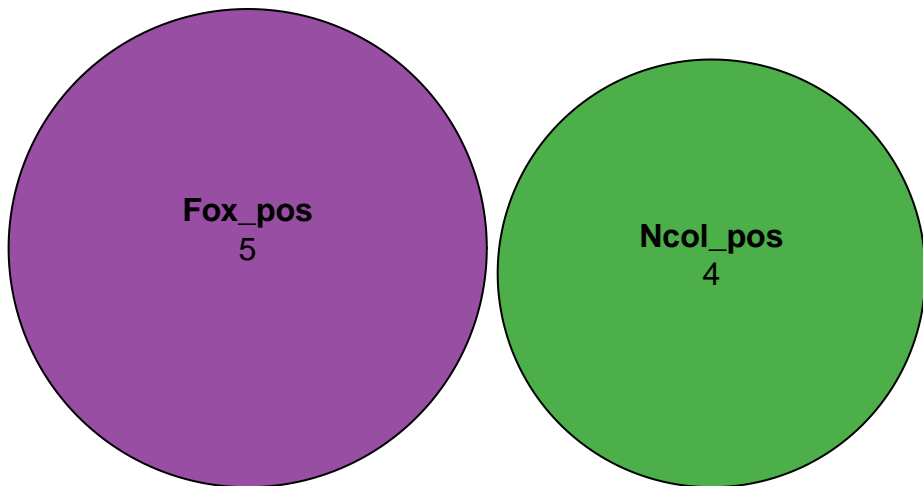
Nvec_vc1.1_XM_048734432.1

Folliculin-interacting pr; FNIP_N/FNIP_M/FNIP_C/



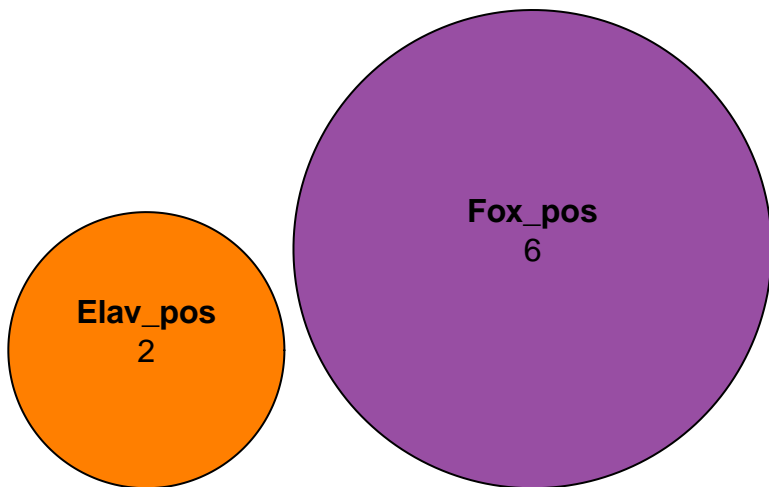
Nvec_vc1.1_XM_001627750.3

;



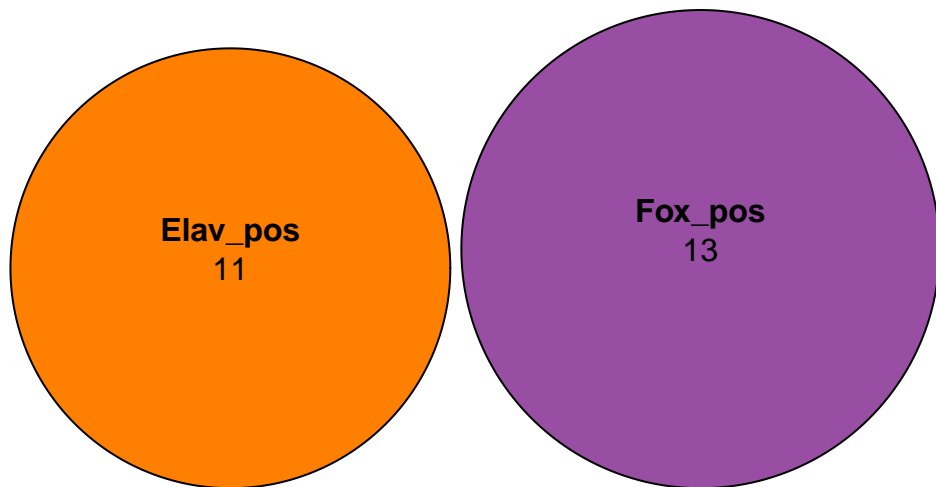
Nvec_vc1.1_XM_048722463.1

GPCR-rhodopsin:NA; 7tm_1/



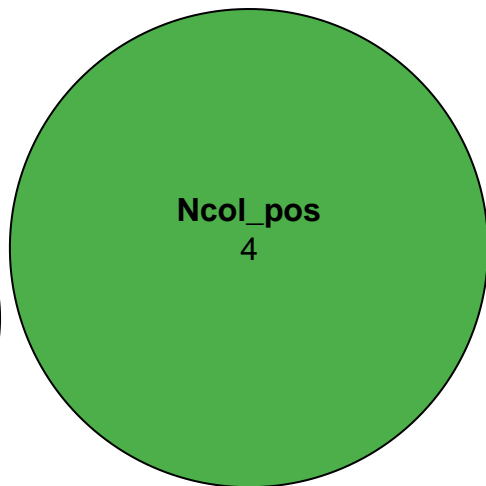
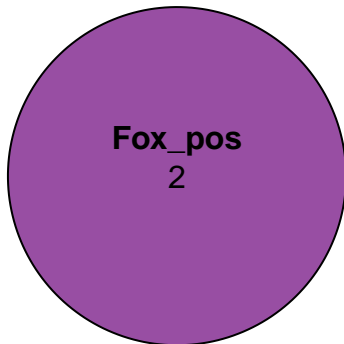
Nvec_vc1.1_XM_001627964.3

n-NaKCa:KCNA1/KCNA10/KC; ion-NaKCa.HG1.6:KCNA1/KC



Nvec_vc1.1_XM_048729250.1

;



Nvec_vc1.1_XM_001641405.3

;

Fox_pos

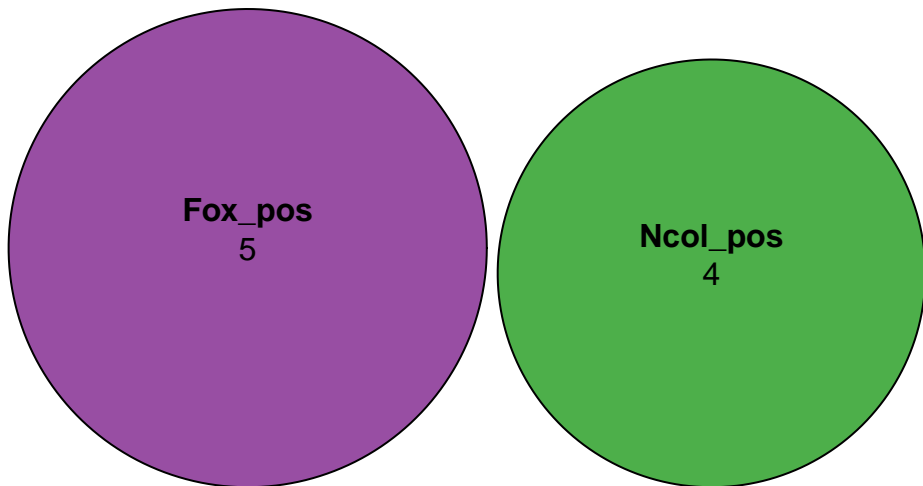
2

Ncol_pos

2

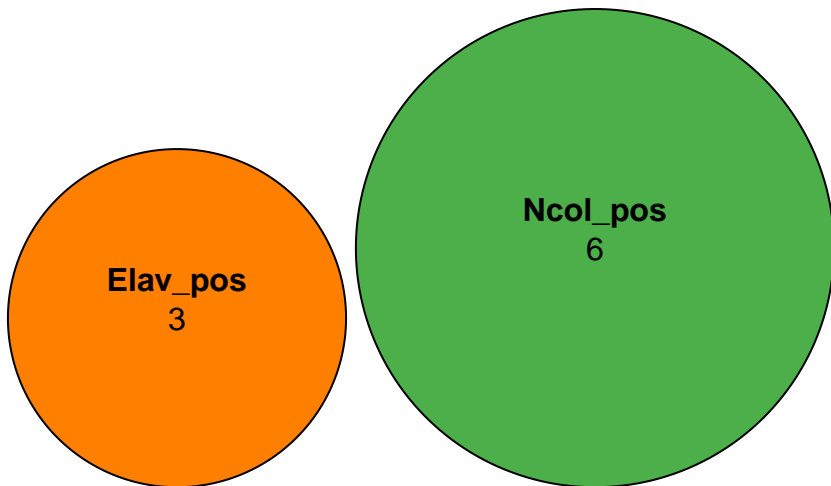
Nvec_vc1.1_XM_032375896.2

;



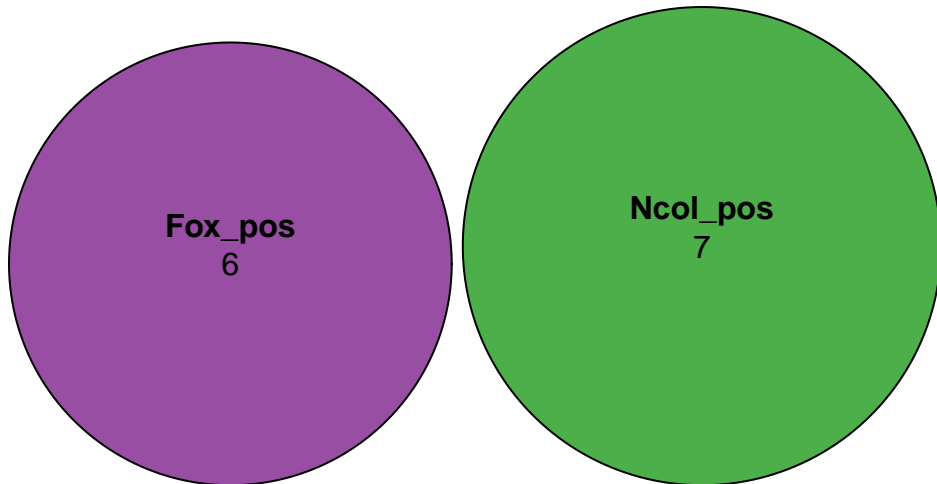
Nvec_vc1.1_XM_032376323.2

Pkinase:like:GUCY2C/GUCY2; Pkinase_Tyr/HNOBA/Guanyla



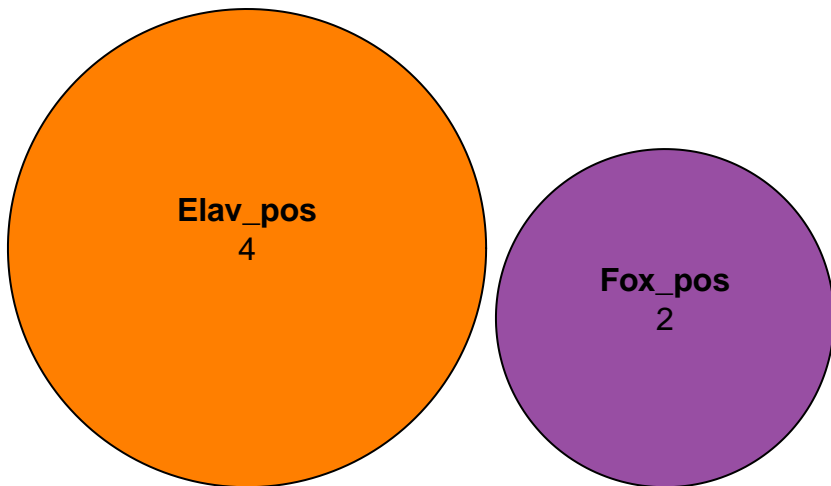
Nvec_vc1.1_XM_032381641.2

b(0,+)-type amino acid tr; AA_permease_2/



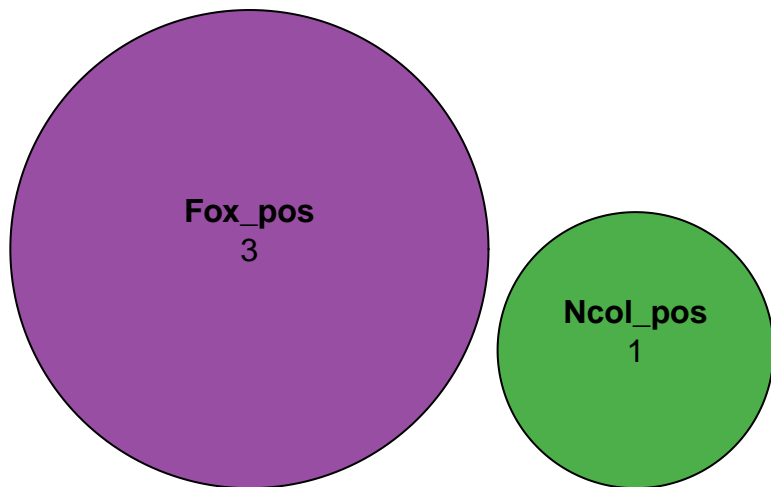
Nvec_vc1.1_XM_001632745.3

Protein NipSnap homolog 3; NIPSNAP/NIPSNAP/

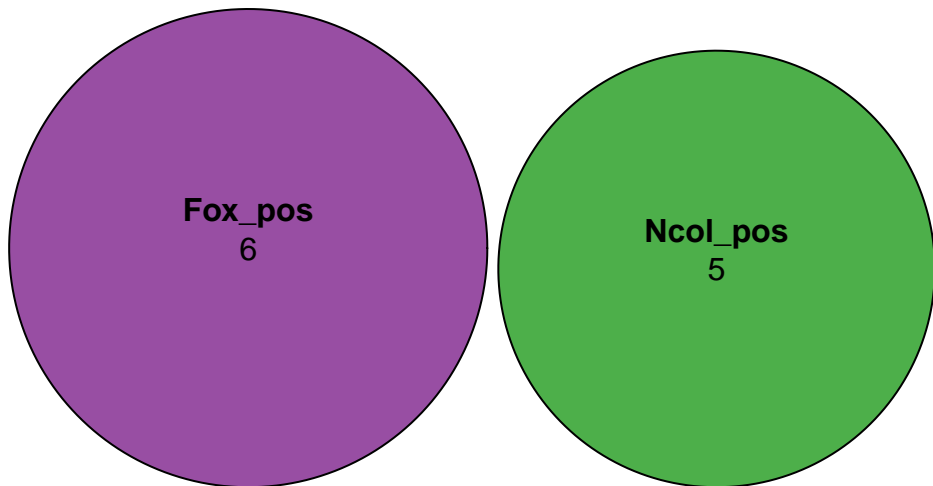


Nvec_vc1.1_XM_048724152.1

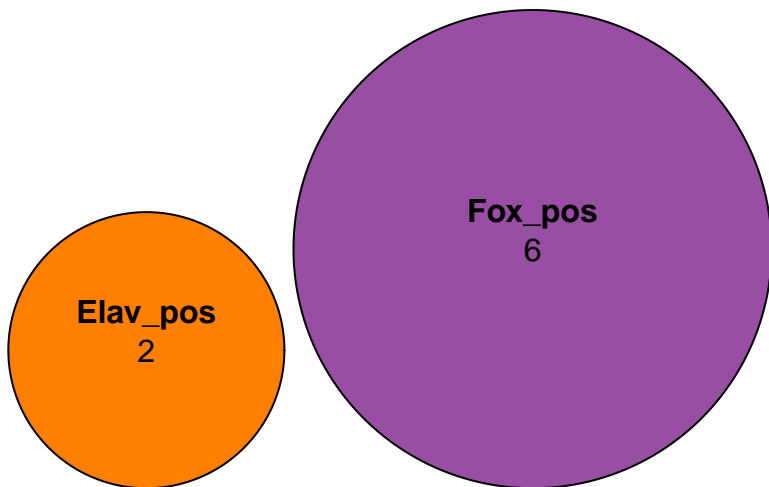
GDP-fucose protein O-fuco; O-FucT/



Nvec_vc1.1_XM_048731865.1
GPCR-rhodopsin:like:GPR17; 7tm_1/

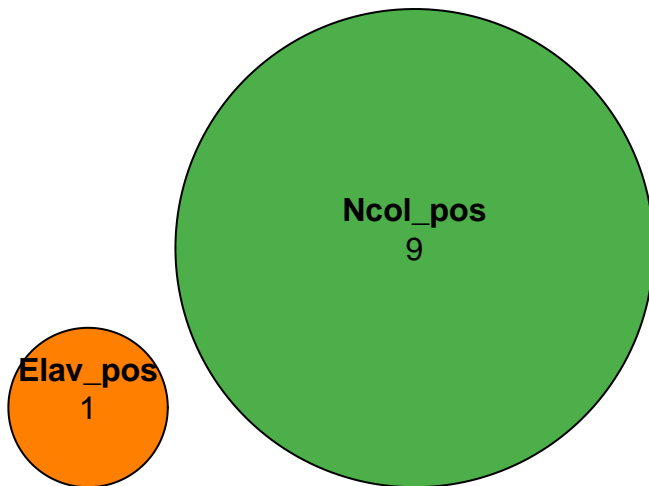


Nvec_vc1.1_XM_048731890.1
GPCR-rhodopsin:like:GPR17; 7tm_1/



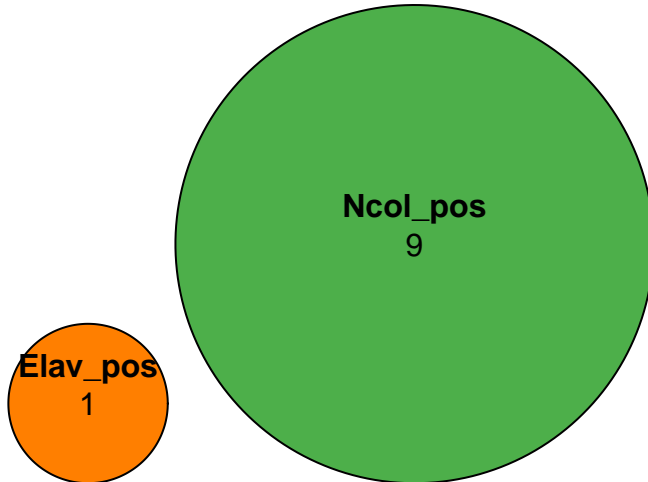
Nvec_NVE13556

;



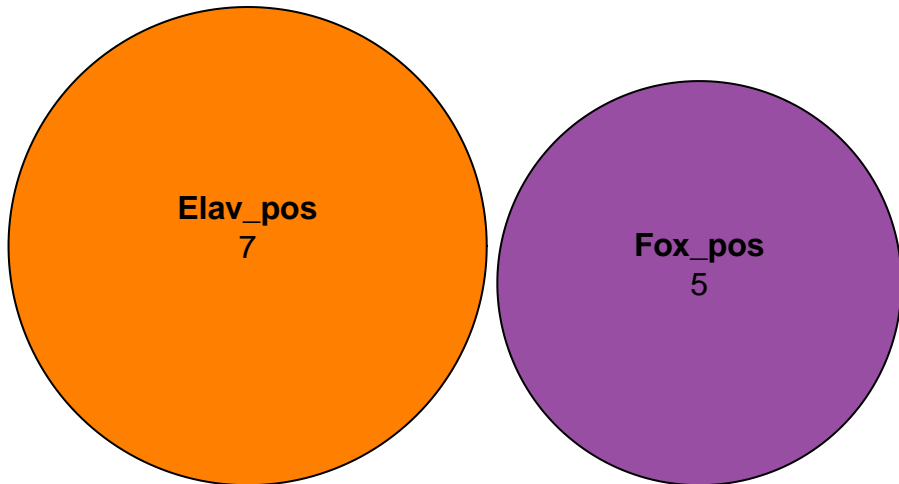
Nvec_vc1.1_XM_032387399.2

;



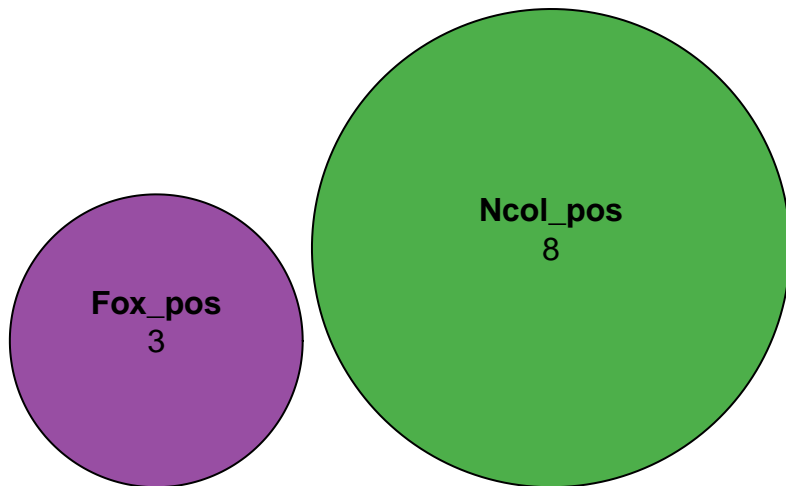
Nvec_vc1.1_XM_048729747.1

Kinesin-like protein KIF1; Kinesin/FHA/

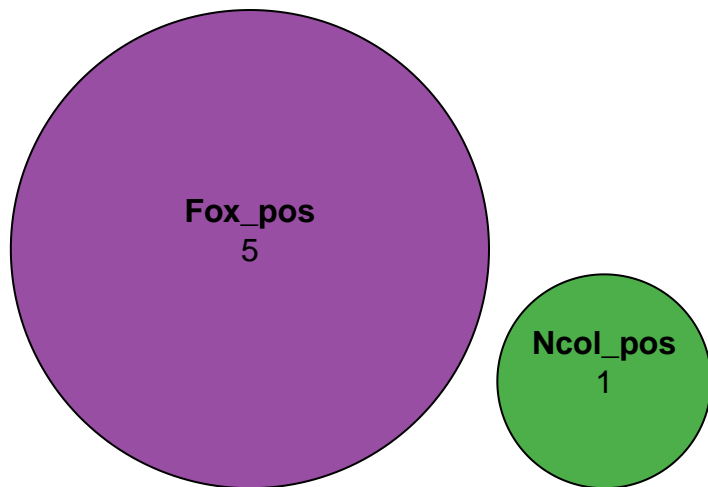


Nvec_vc1.1_XM_032382956.2

Galectin; VHL/

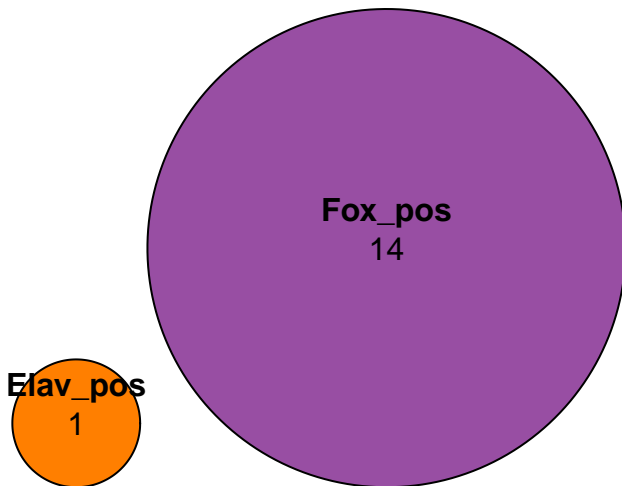


Nvec_vc1.1_XM_001627245.3
cDNA FLJ56852, highly sim; I_LWEQ/



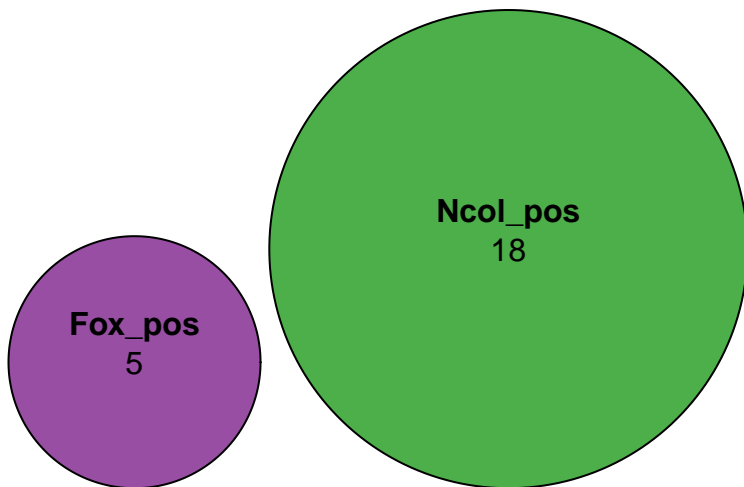
Nvec_vc1.1_XM_001640356.3

;

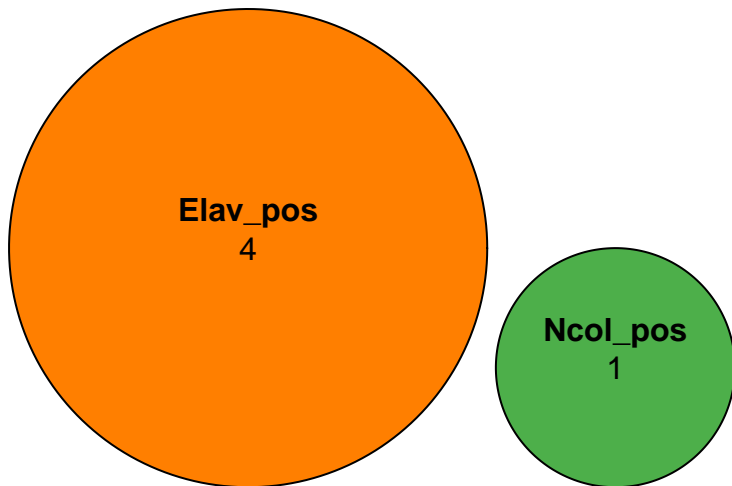


Nvec_vc1.1_XM_032377937.2

ion-neuro-gated:like:GABR; Neur_chan_LBD/Neur_chan_m

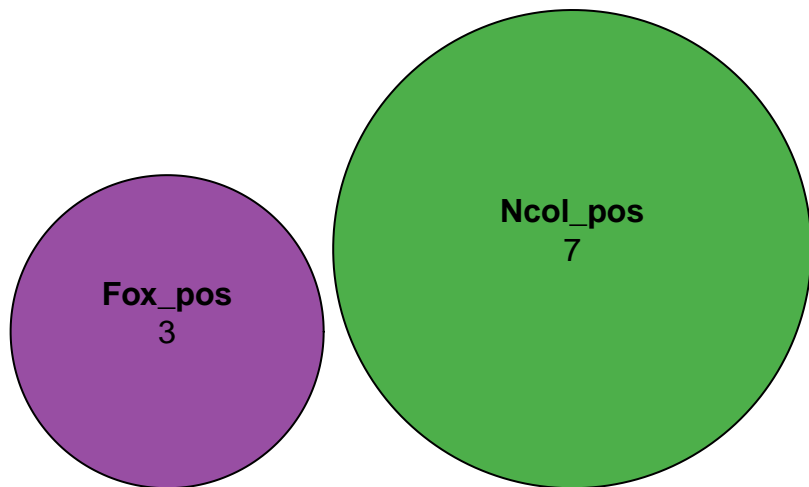


Nvec_vc1.1_XM_001636964.3
cDNA FLJ77769, highly sim; NAD_kinase/

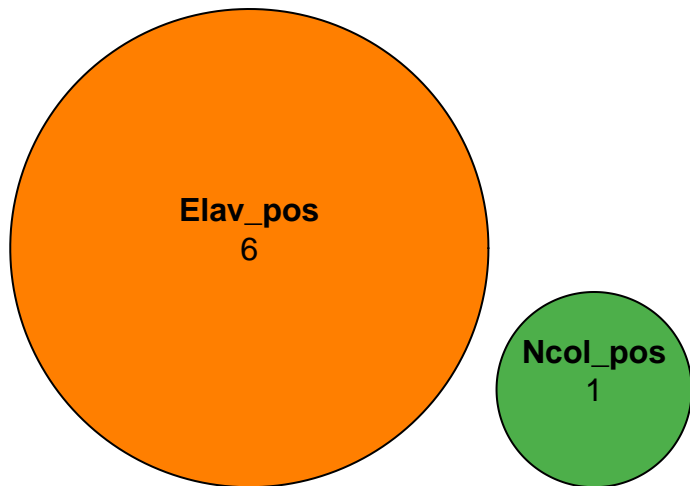


Nvec_vc1.1_XM_001623818.3

ion-NaKCa:like:KCNA1/KCNA; BTB_2/Ion_trans/

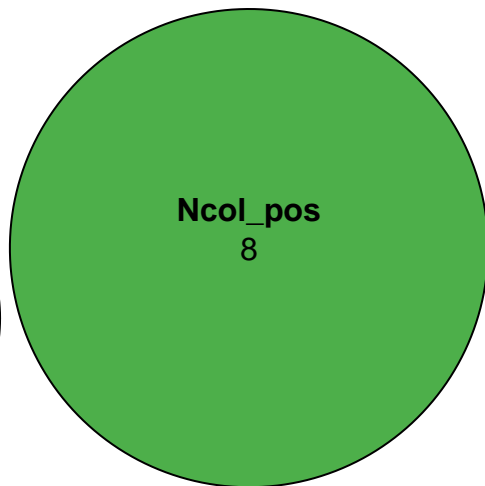


Nvec_vc1.1_XM_001637022.3
GPCR-rhodopsin:NA; 7tm_1/

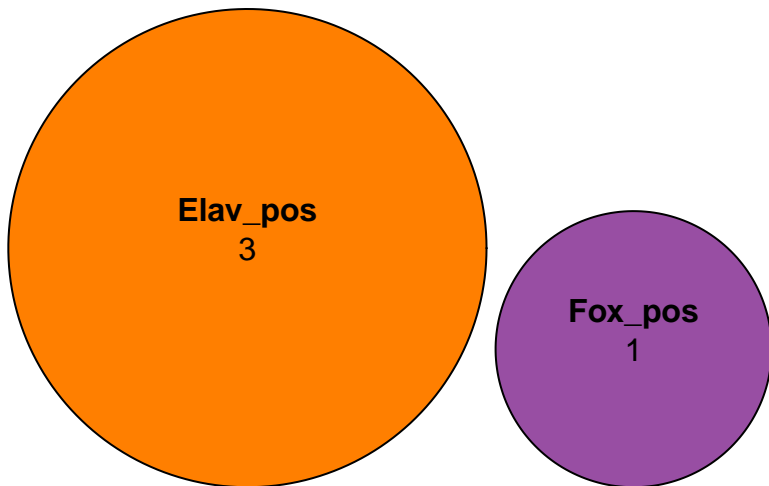


Nvec_vc1.1_XM_048731921.1

;

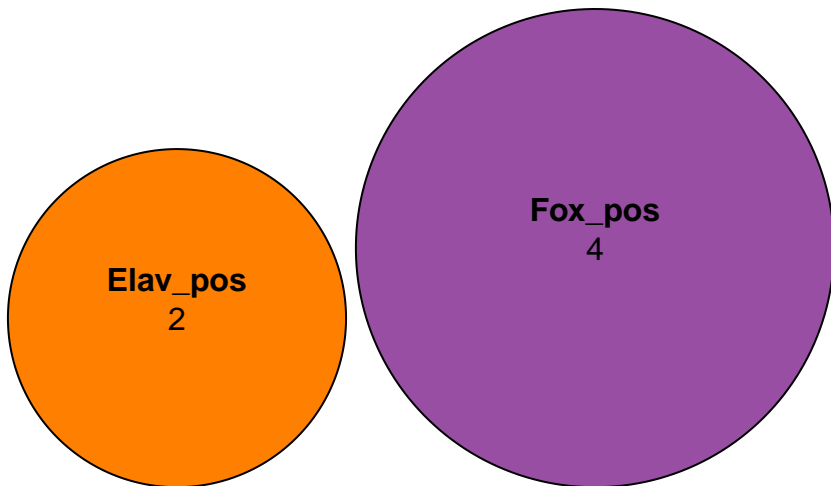


Nvec_vc1.1_XM_001631508.3
YTH:YTHDC1; YTH.HG1.0:YTHDC1



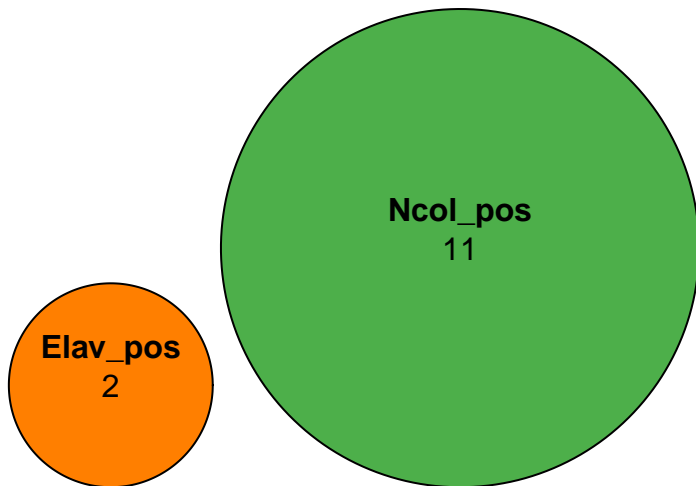
Nvec_vc1.1_XM_032370981.2

Synaptotagmin:SYT12; Synaptotagmin.HG1.38:SYT1

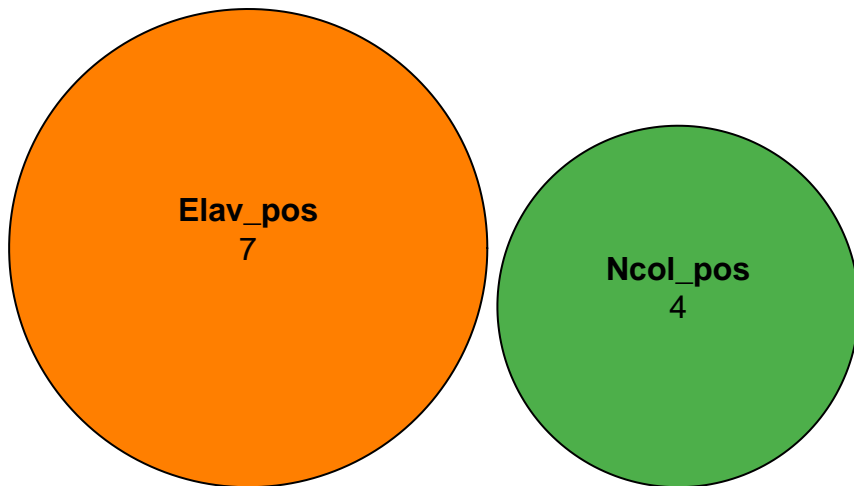


Nvec_vc1.1_XM_032384420.2

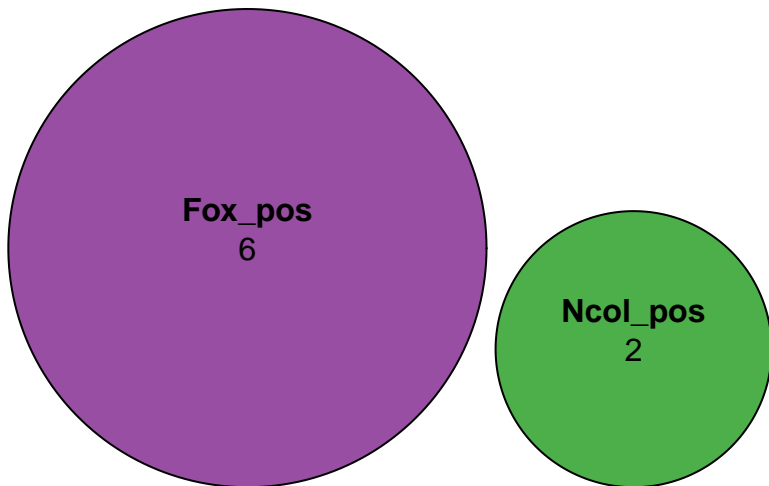
Sp6-9; SP6/SP7/SP8/SP9



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

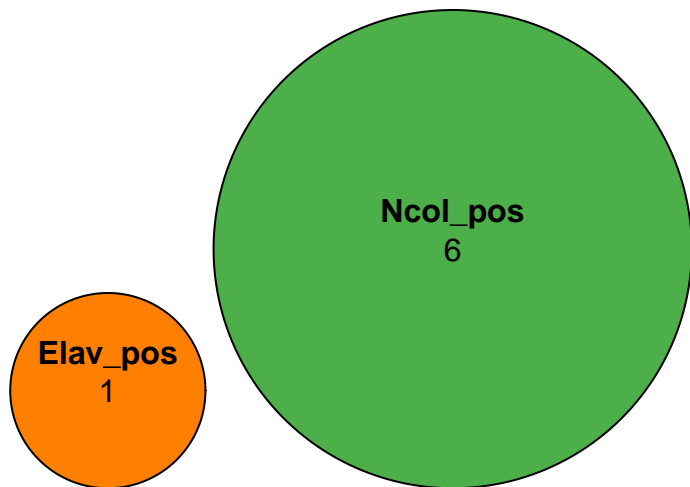


Nvec_vc1.1_XM_001637149.3
RAP1A, member of RAS onco; Ras/

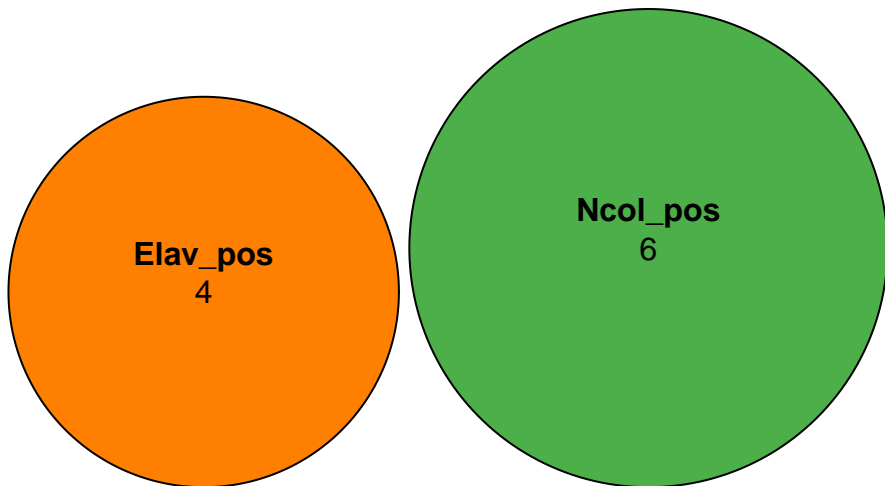


Nvec_vc1.1_XM_001628711.3

:HNRNPC/HNRNPCL1/HNRNP; RRM.HG17.3:HNRNPC/HNRNP

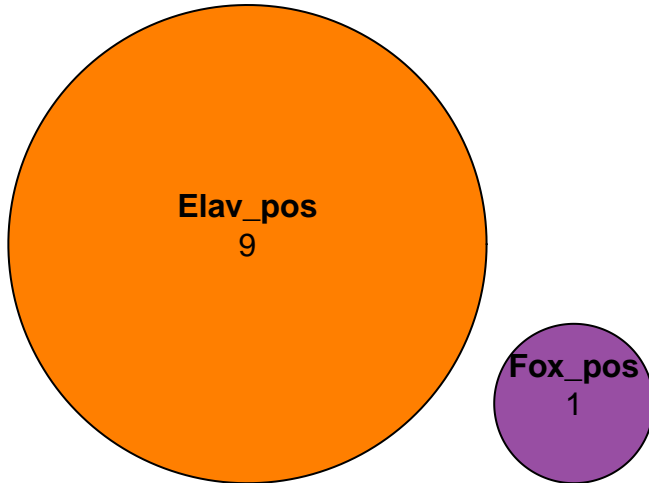


Nvec_vc1.1_XM_001630845.3
ion-glutamate:NA; Ion_trans_2/



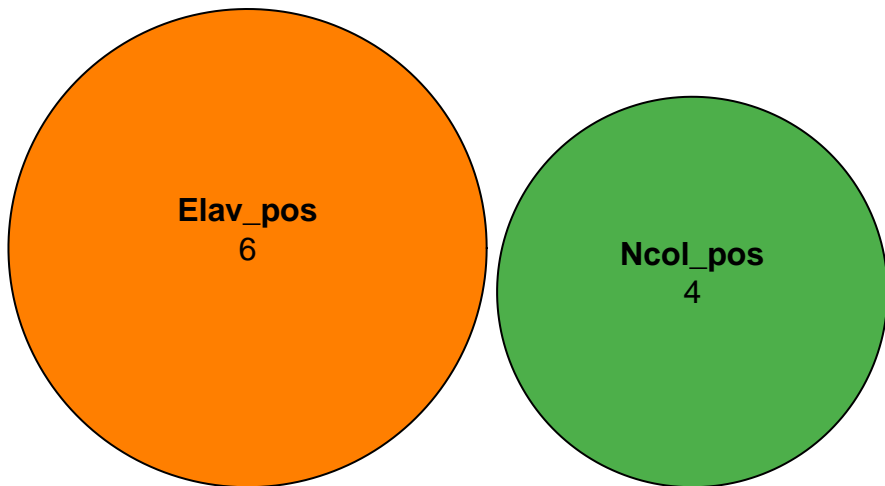
Nvec_vc1.1_XM_032379049.2

;



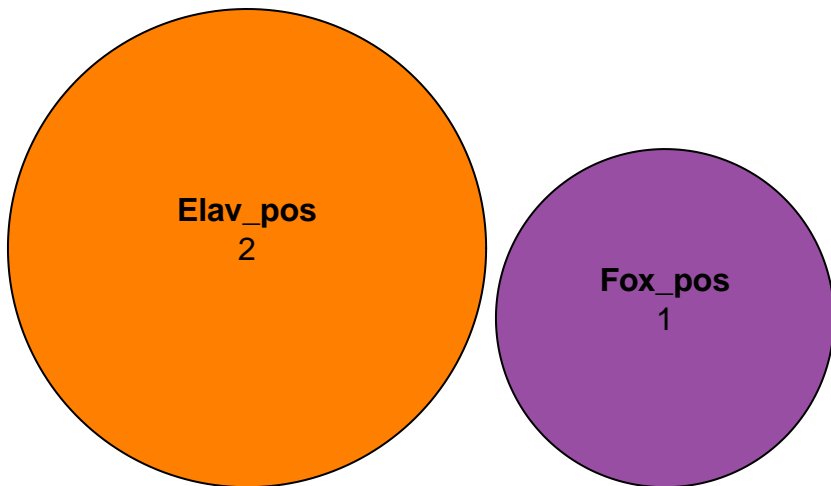
Nvec_vc1.1_XM_032376506.2

Zinc finger protein 862; Dimer_Tnp_hAT/



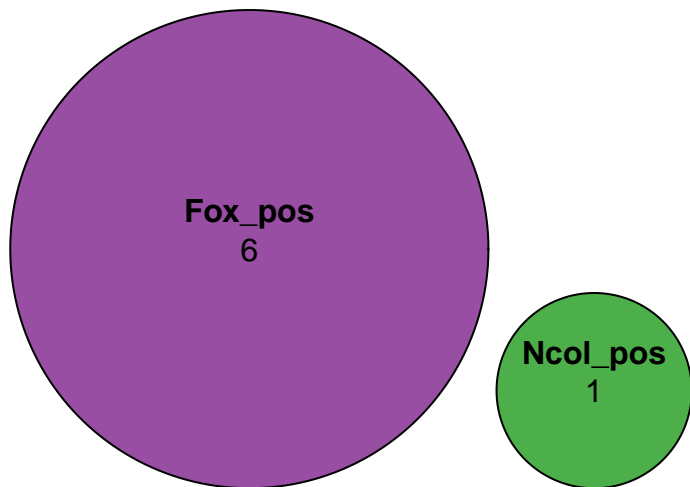
Nvec_vc1.1_XM_032370048.2

CCCH:MKRN1/MKRN2/MKRN3; zf-CCCH.HG1.17:MKRN1/MK

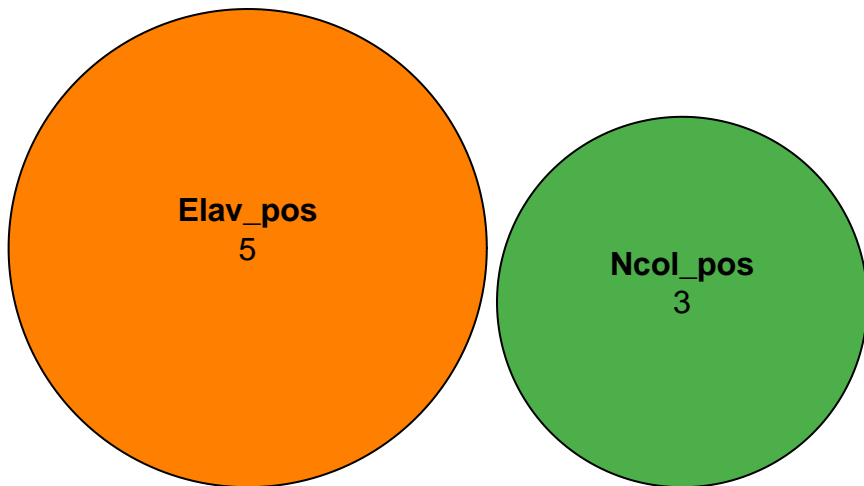


Nvec_vc1.1_XM_048731597.1

Allograft inflammatory fa; EF-hand_7/

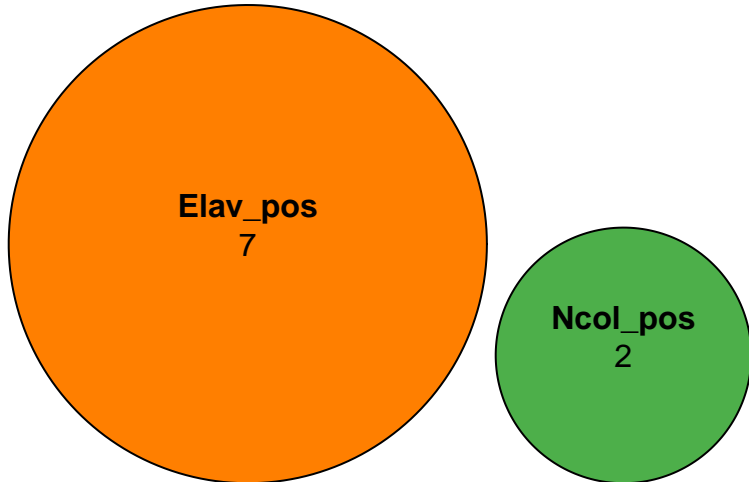


Nvec_vc1.1_XM_048731252.1
OTU domain-containing pro; OTU/



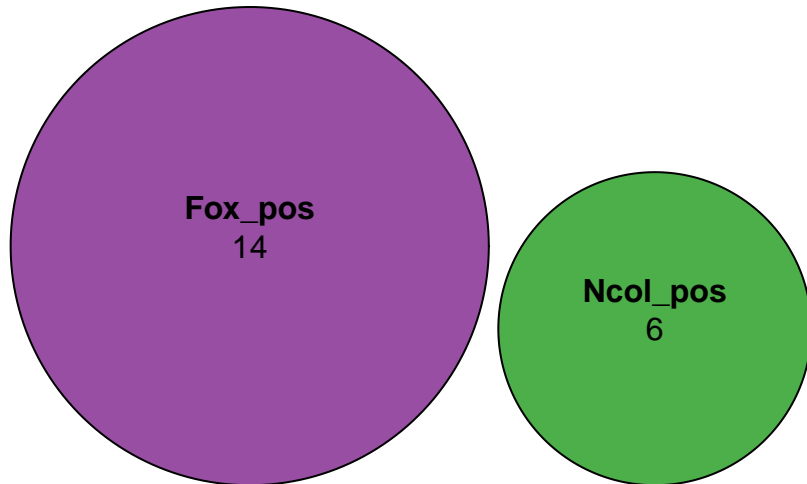
Nvec_vc1.1_XM_001633821.3

;



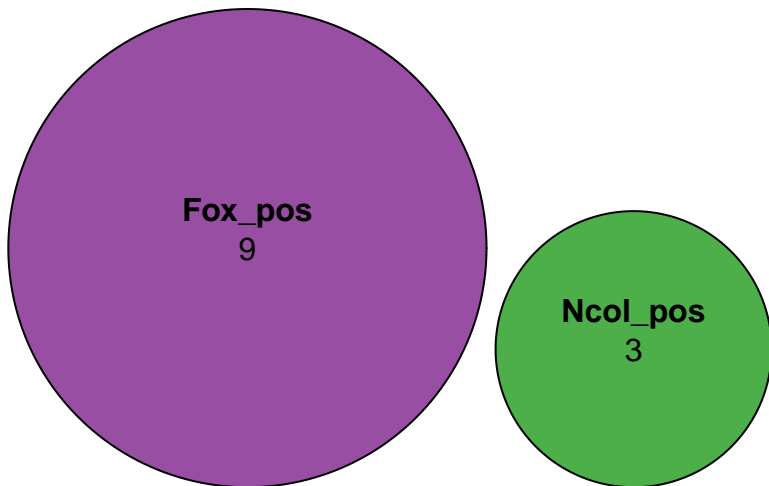
Nvec_vc1.1_XM_032379911.2

Sparc/osteonectin, cwcv a; Thyroglobulin_1/



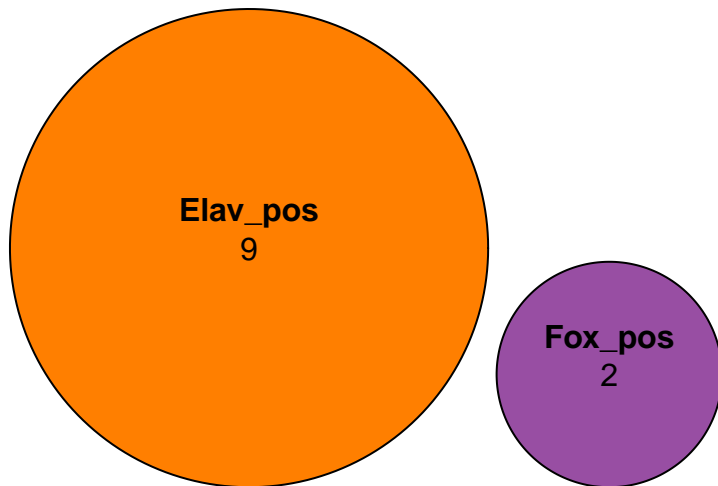
Nvec_vc1.1_XM_032370084.2

;



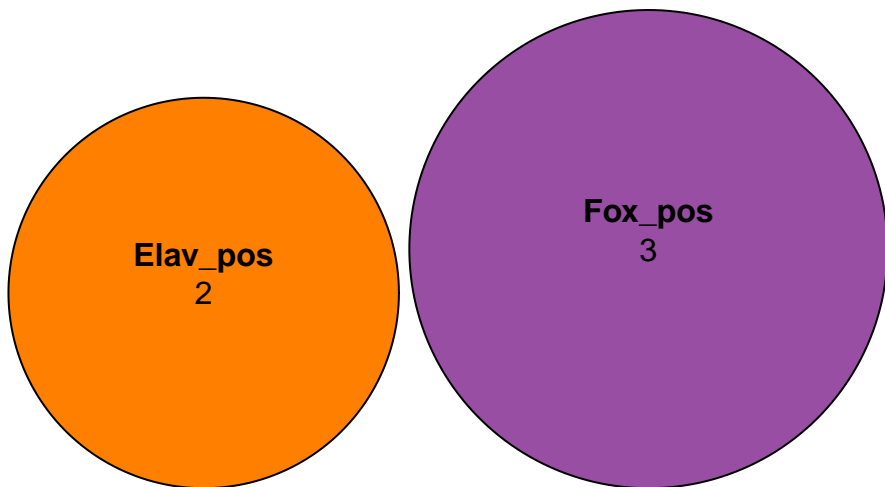
Nvec_vc1.1_XM_048723783.1

Cilia- and flagella-assoc;



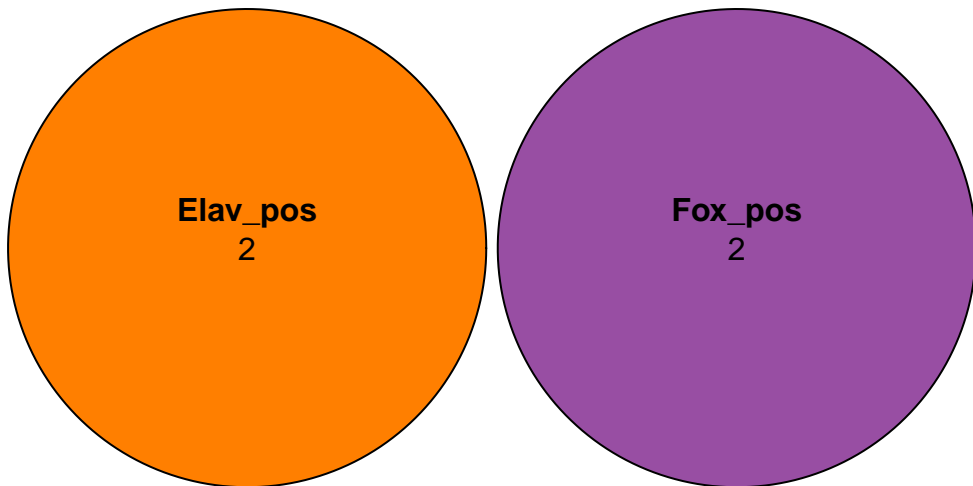
Nvec_vc1.1_XM_048725946.1

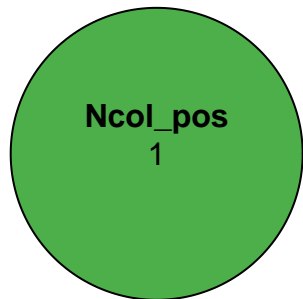
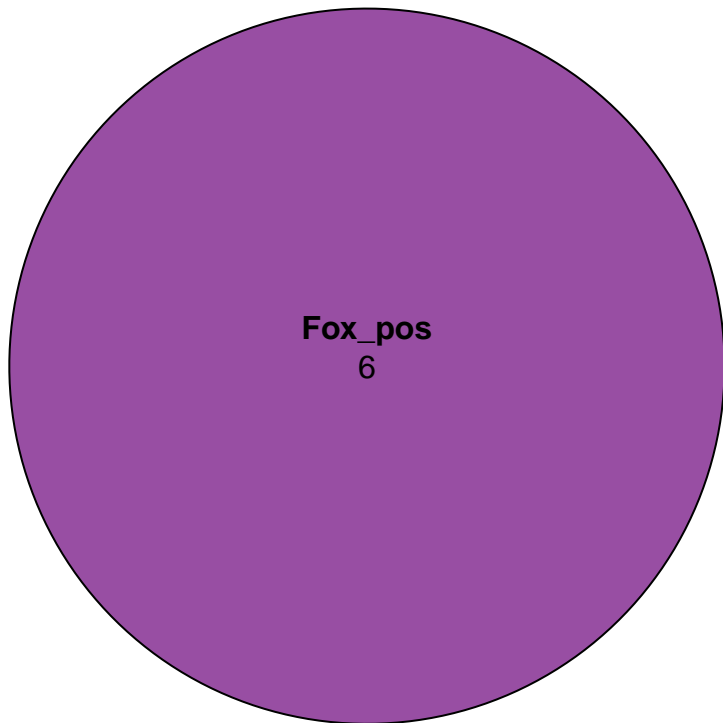
WD repeat-containing prot;



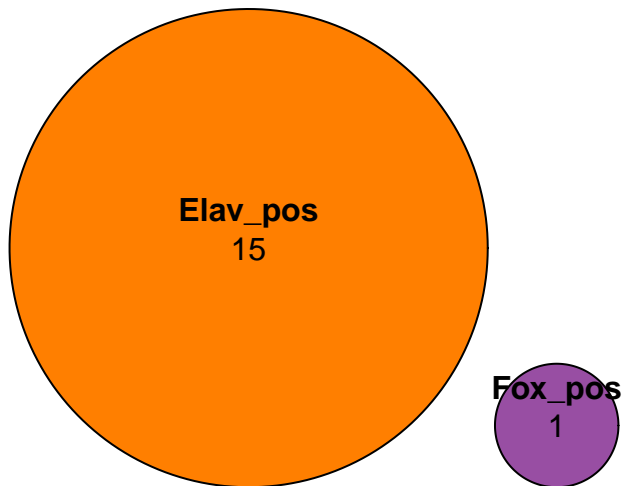
Nvec_vc1.1_XM_001636610.3

;

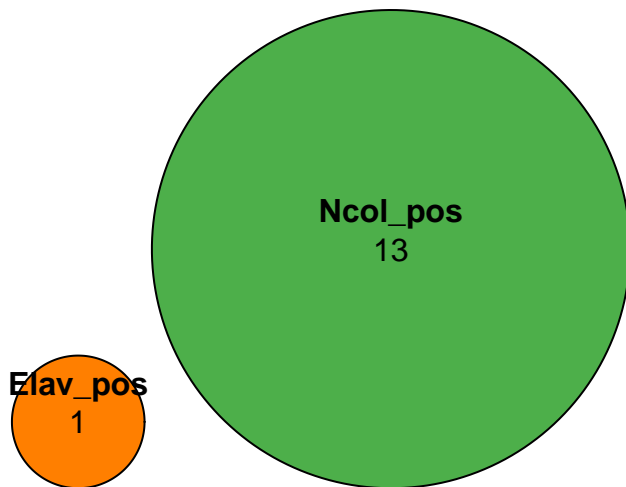




Nvec_vc1.1_XM_001628279.3
HLH.HG1.76:ATOH8; ATOH8

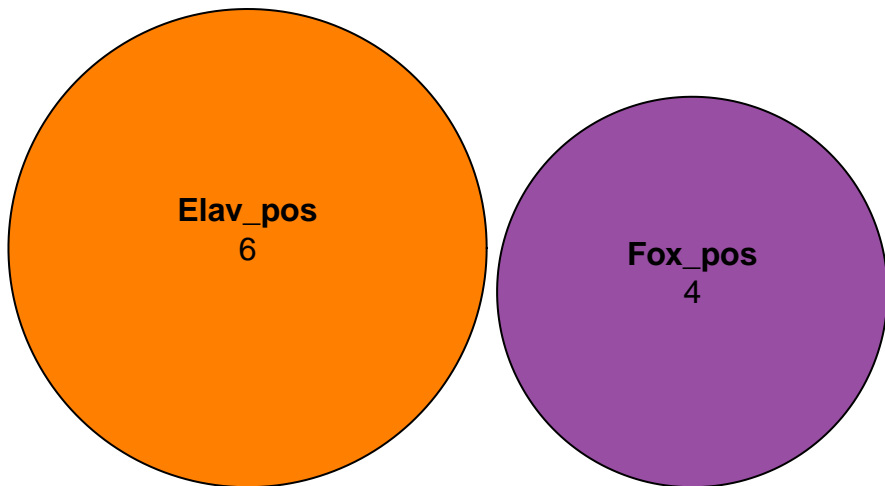


Nvec_vc1.1_XM_032377736.2
Protein FAM173B; Methyltransf_23/

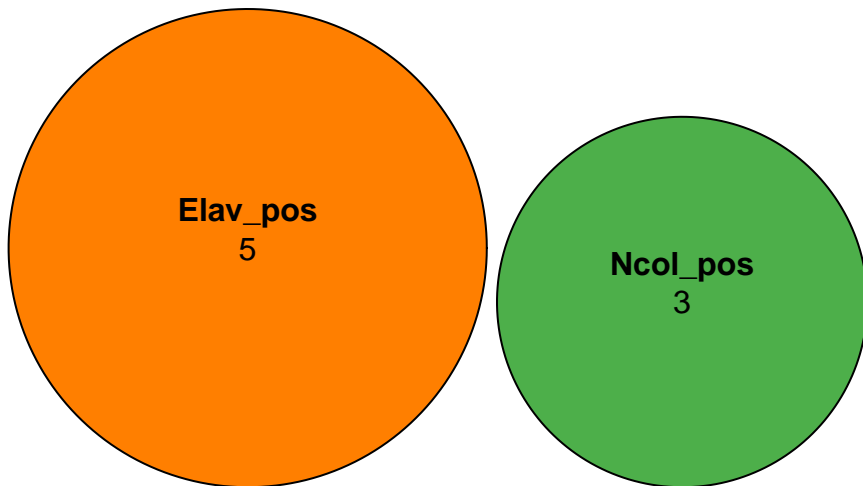


Nvec_vc1.1_XM_032361849.2

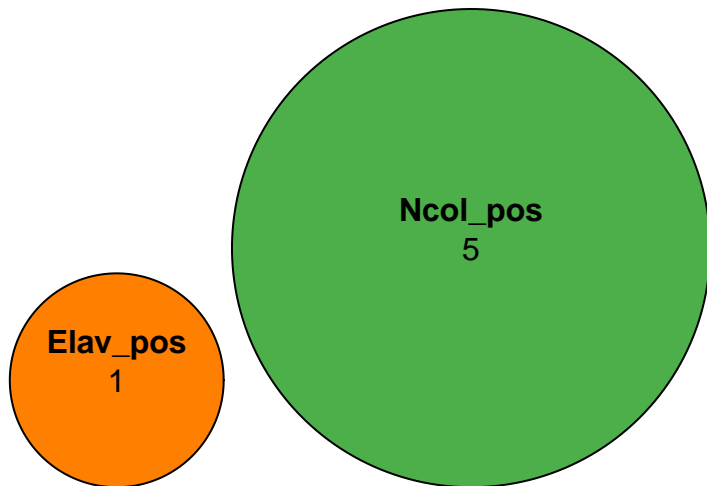
Ets.HG1.2:ELK1/ELK3/ELK4; ELK1/ELK3/ELK4



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

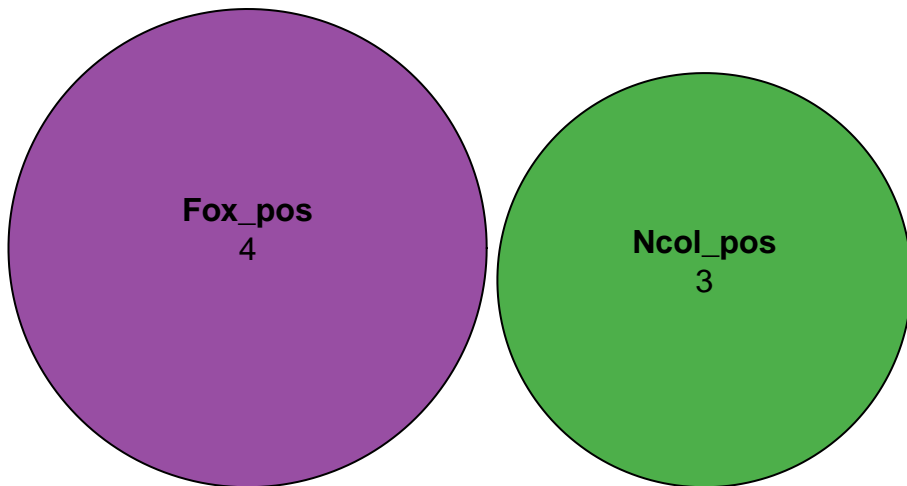


Nvec_vc1.1_XM_001639611.1
; EF-hand_7/



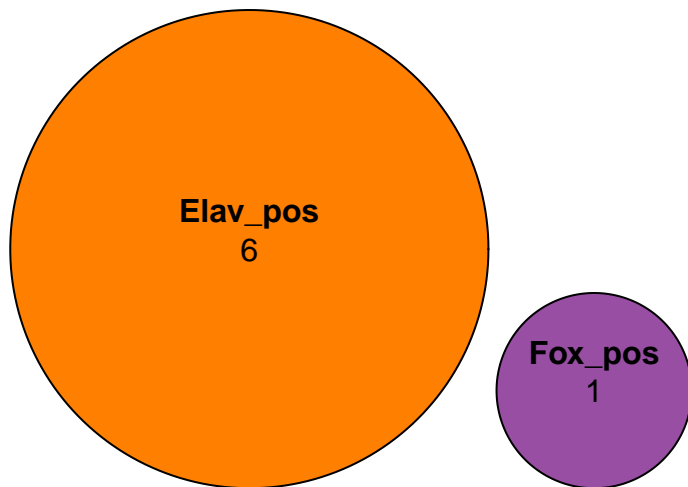
Nvec_vc1.1_XM_032362682.2

CEA protein; V-set/Ig_2/

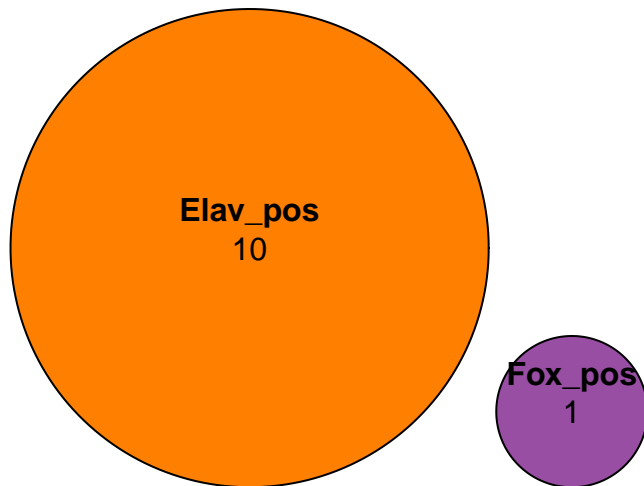


Nvec_vc1.1_XM_032380524.2

Pkinase:PIK3R4; Pkinase.HG180.0:PIK3R4

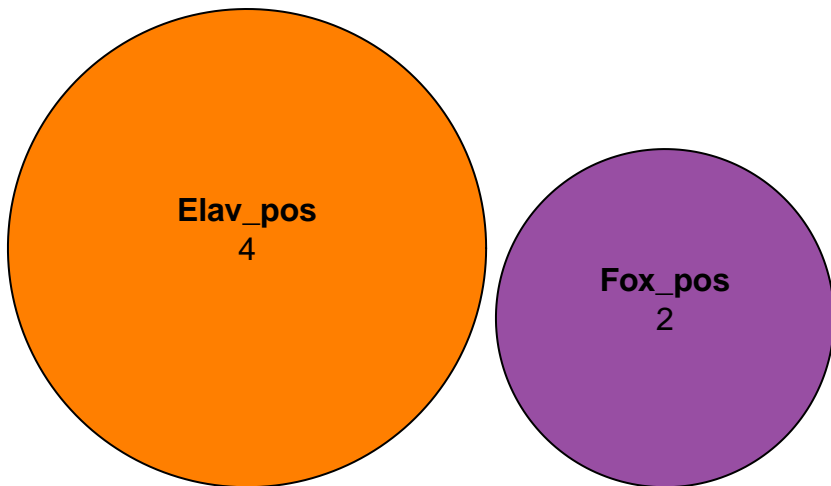


Nvec_vc1.1_XM_032378372.2
GPCR-rhodopsin:like:ADORA; 7tm_1/

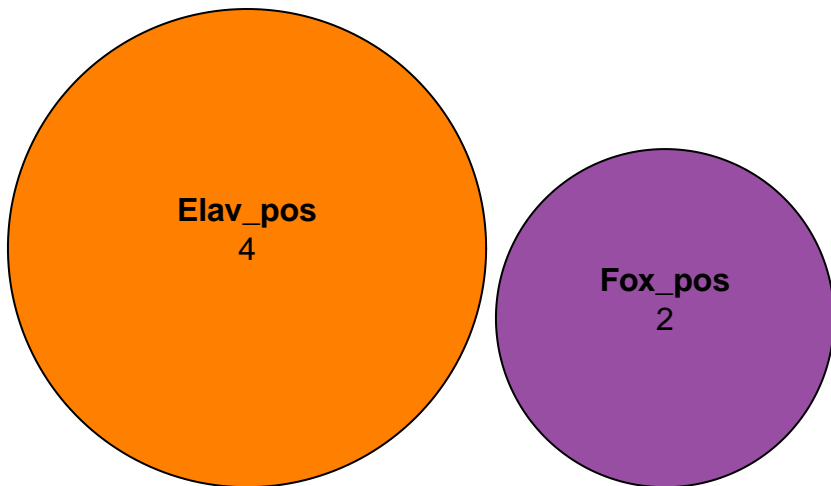


Nvec_vc1.1_XM_032386263.2

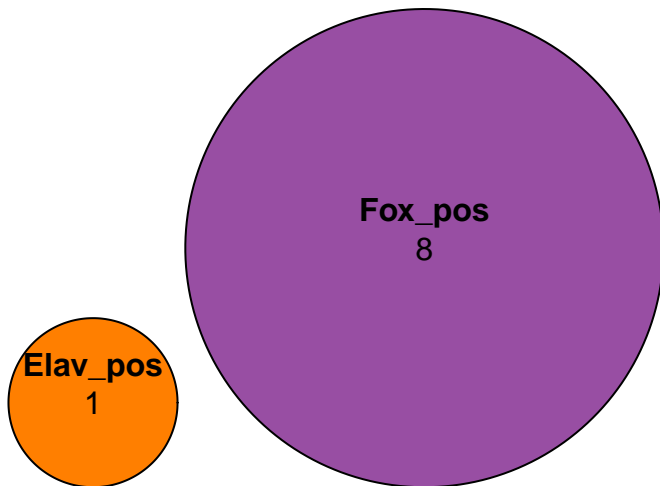
Rhombotin-1; LIM/LIM/



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

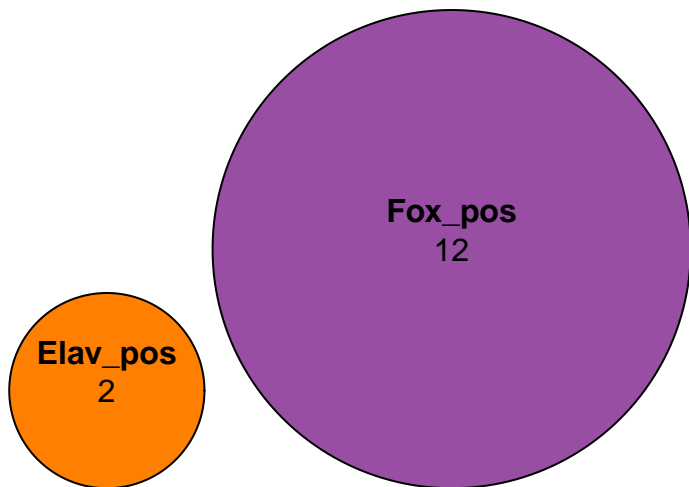


Nvec_vc1.1_XM_032379543.2
; DUF885/



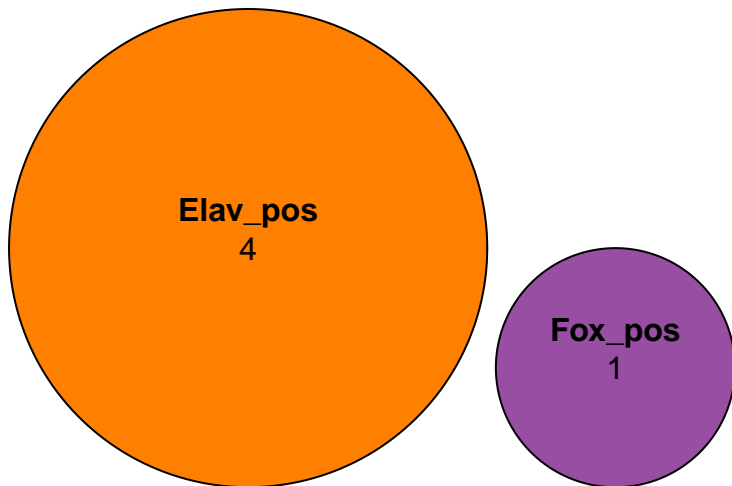
Nvec_vc1.1_XM_001633143.3

Replication factor C subu; AAA/Rep_fac_C/



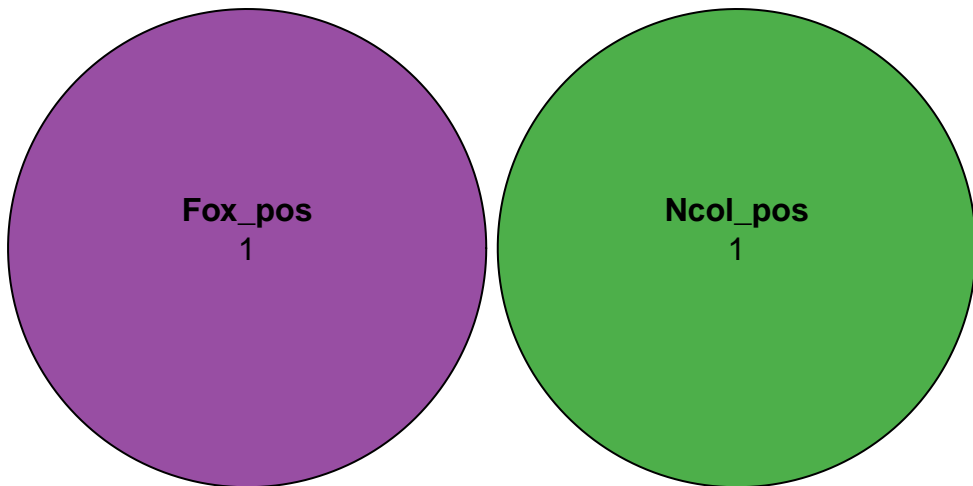
Nvec_vc1.1_XM_048720909.1

;

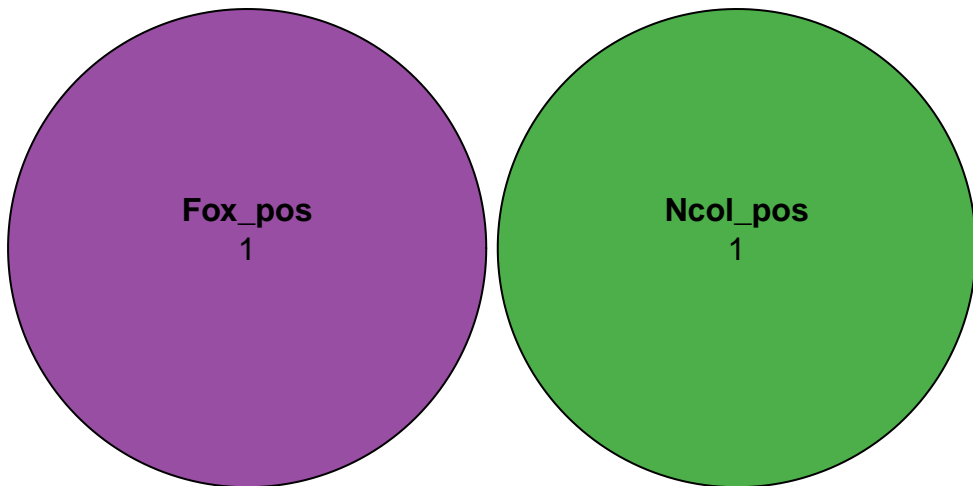


Nvec_vc1.1_XM_048729009.1

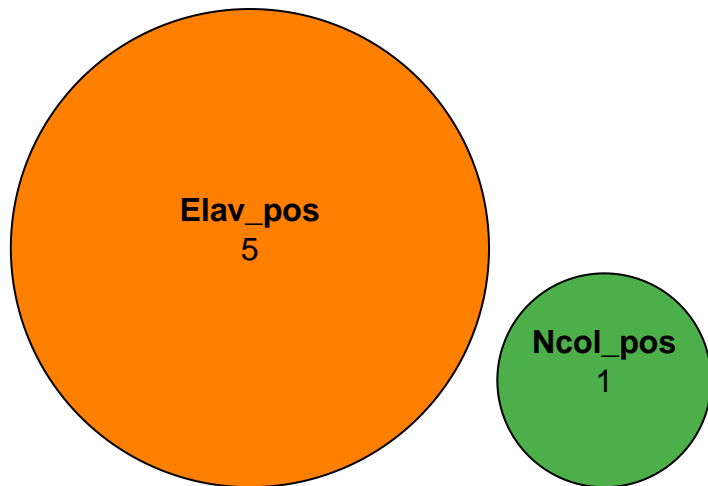
;



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

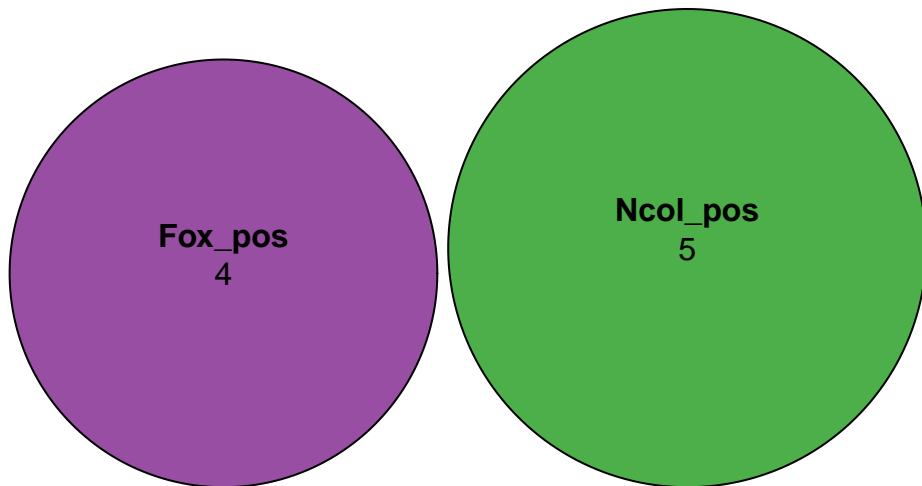


Nvec_vc1.1_XM_001627181.3
Myeloid leukemia factor 1; Mlf1IP/



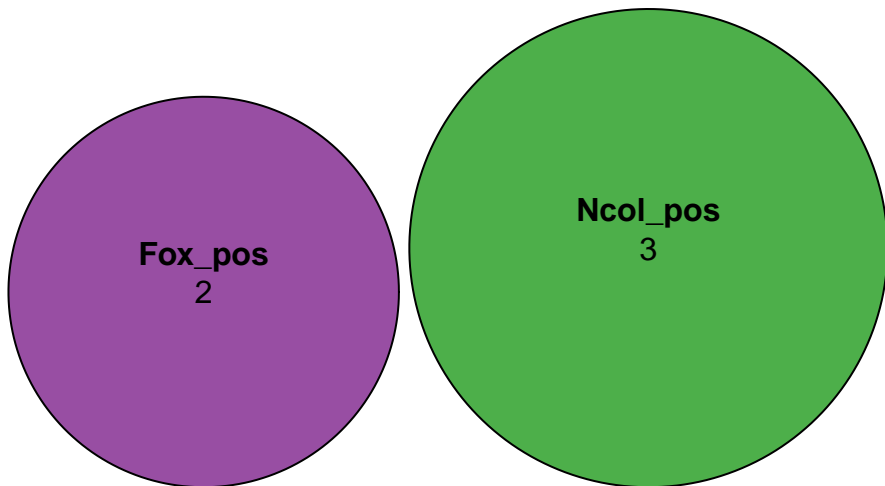
Nvec_vc1.1_XM_032372769.2

fibronectin:like:AXL/MERT; I-set/Ig_2/fn3/fn3/



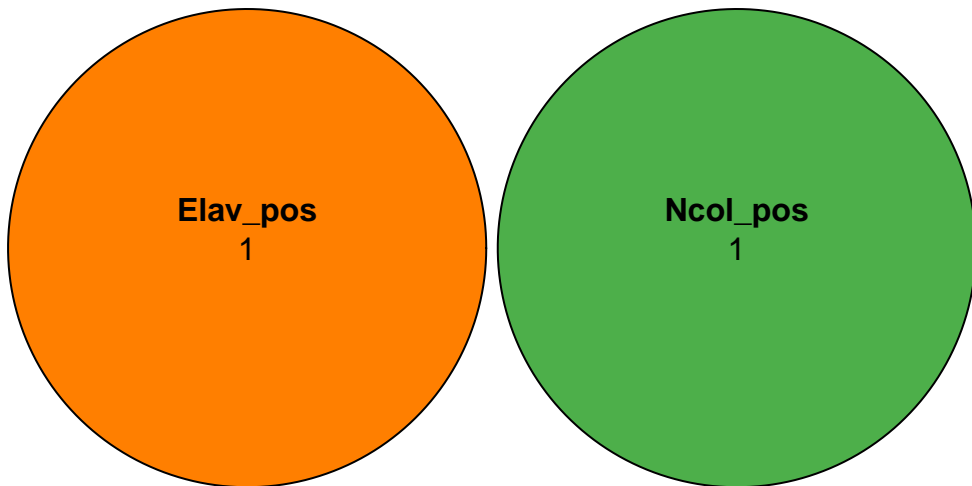
Nvec_vc1.1_XM_048734459.1

E3 ubiquitin–protein liga;



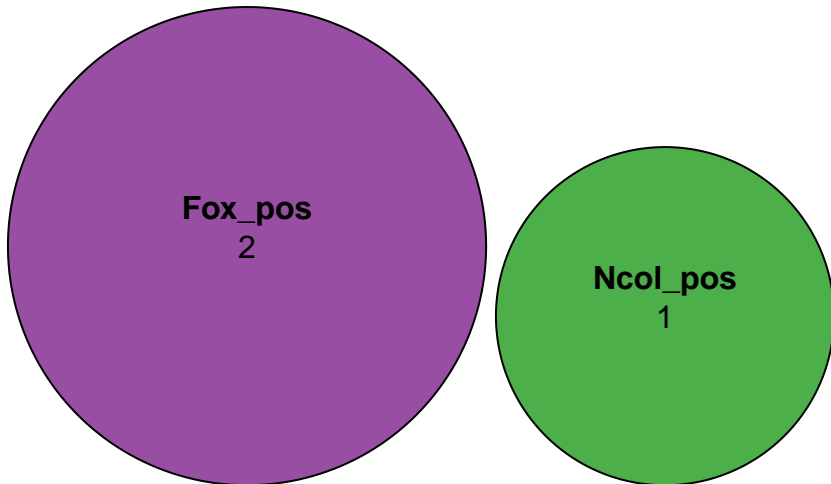
Nvec_vc1.1_XM_001636661.3

Serine/threonine-protein ; Metallophos/

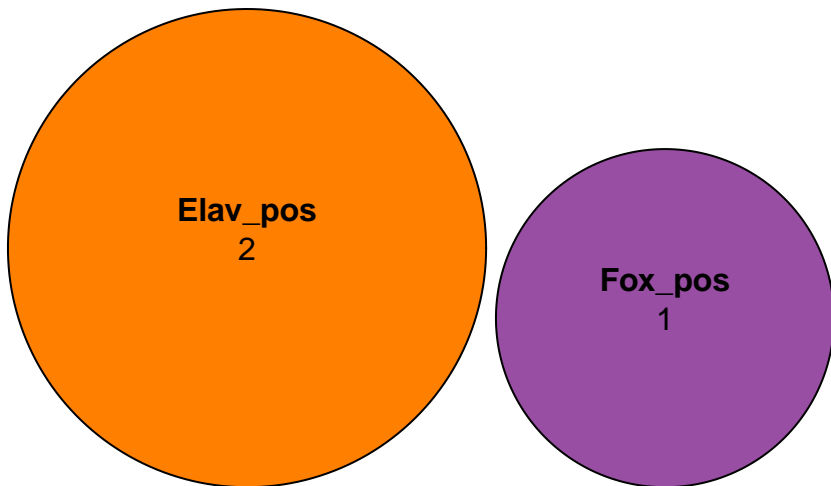


Nvec_vc1.1_XM_001639688.3

Peroxisomal acyl-coenzyme; Acyl-CoA_dh_M/Acyl-CoA_dh

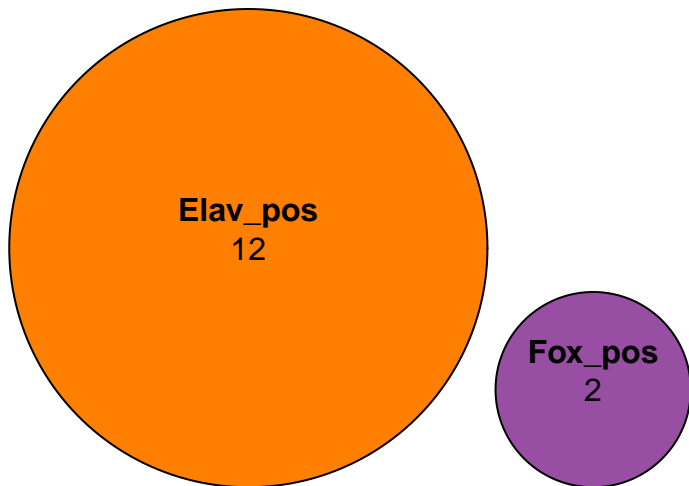


Nvec_vc1.1_XM_032374307.2
Major facilitator superfa; MFS_2/



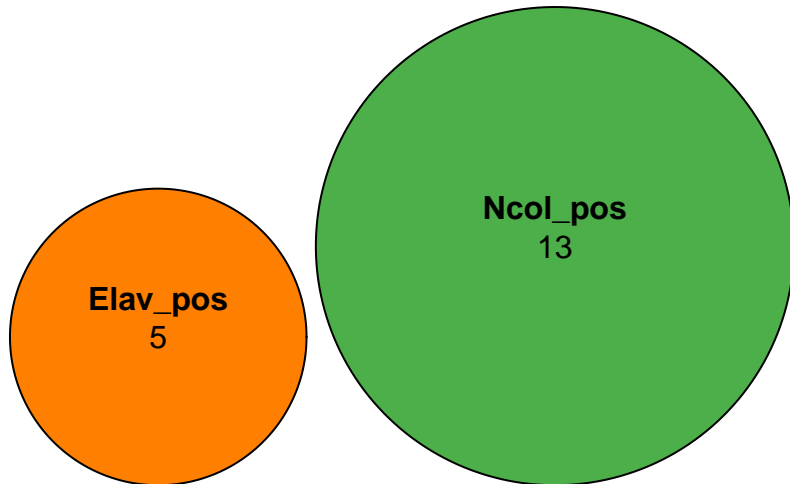
Nvec_vc1.1_XM_032365655.2

;



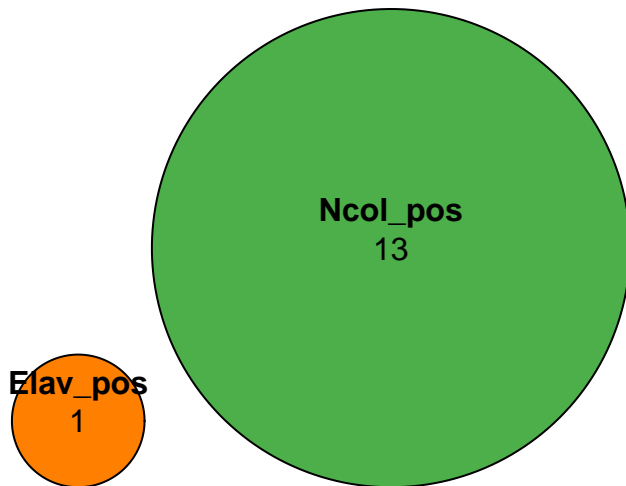
Nvec_vc1.1_XM_048720751.1

cDNA FLJ33589 fis, clone ; Asparaginase_2/

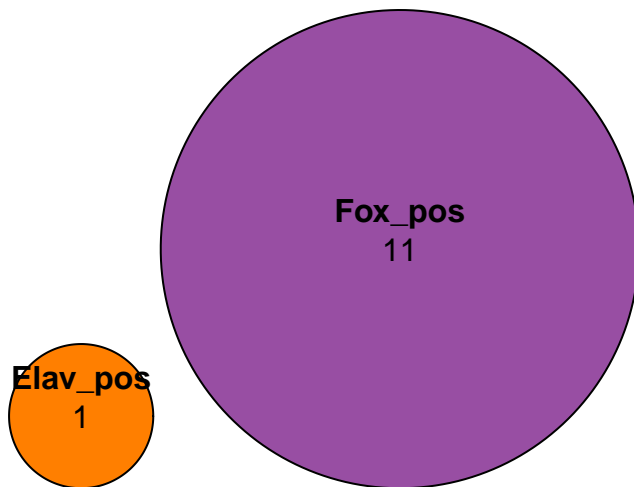


Nvec_vc1.1_XM_048720757.1

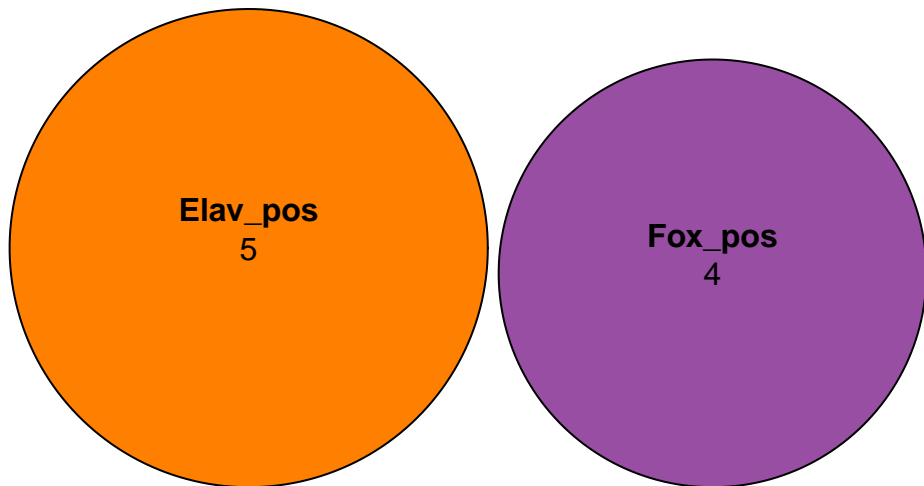
cDNA FLJ33589 fis, clone ; Asparaginase_2/Asparagina



Nvec_vc1.1_XM_032362262.2
GPCR-rhodopsin:like:CCKAR; 7tm_1/

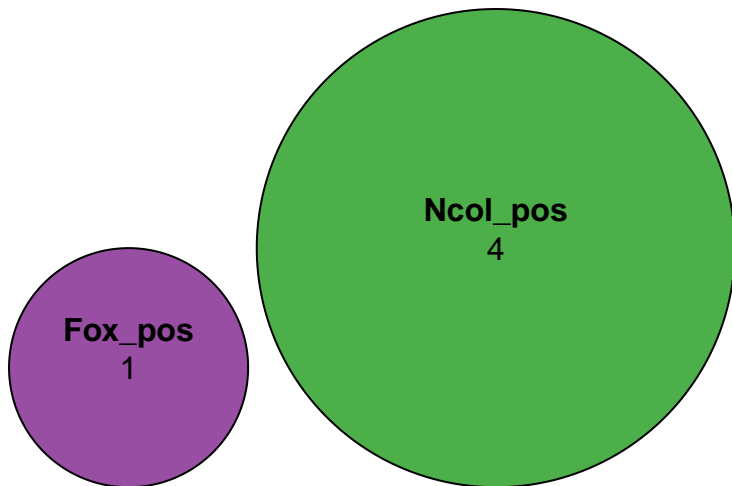


Nvec_vc1.1_XM_048734599.1
GPCR-rhodopsin:like:ADORA; 7tm_1/



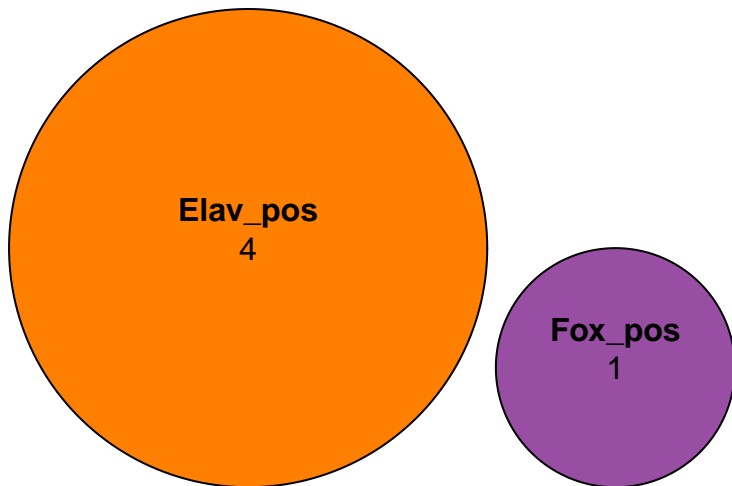
Nvec_vc1.1_XM_048723891.1

;



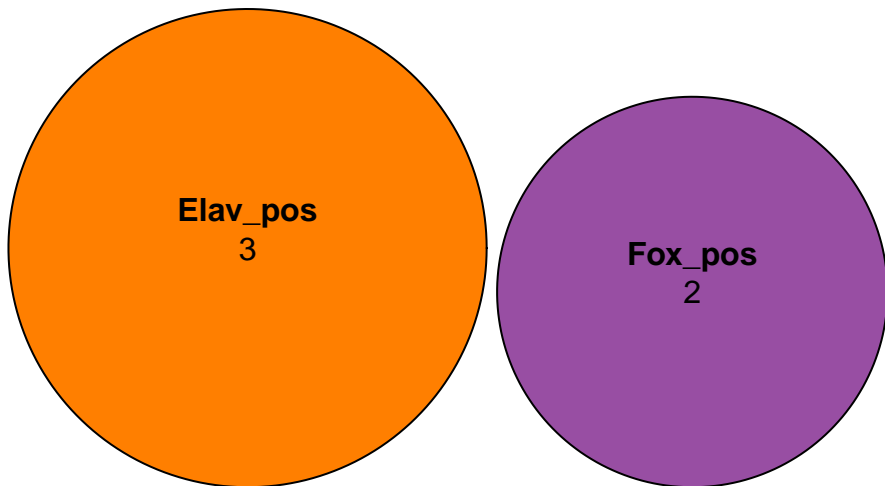
Nvec_vc1.1_XM_032386096.2

Exportin-T; Xpo1/



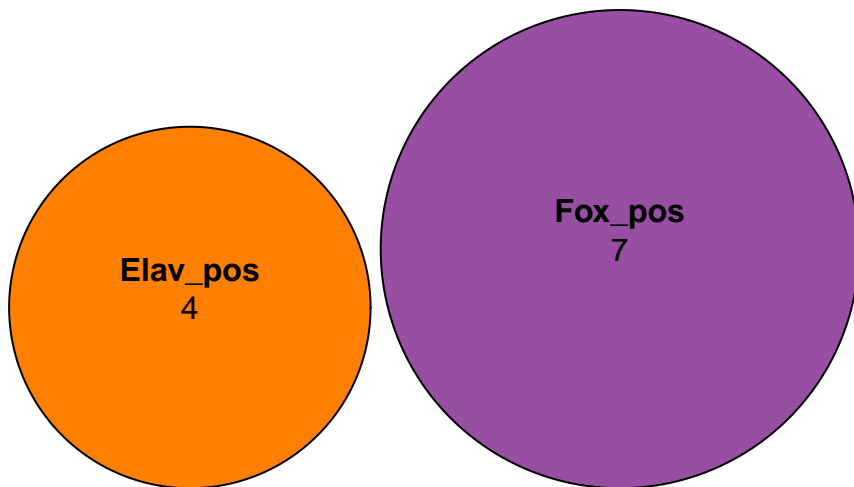
Nvec_vc1.1_XM_032363157.2

;

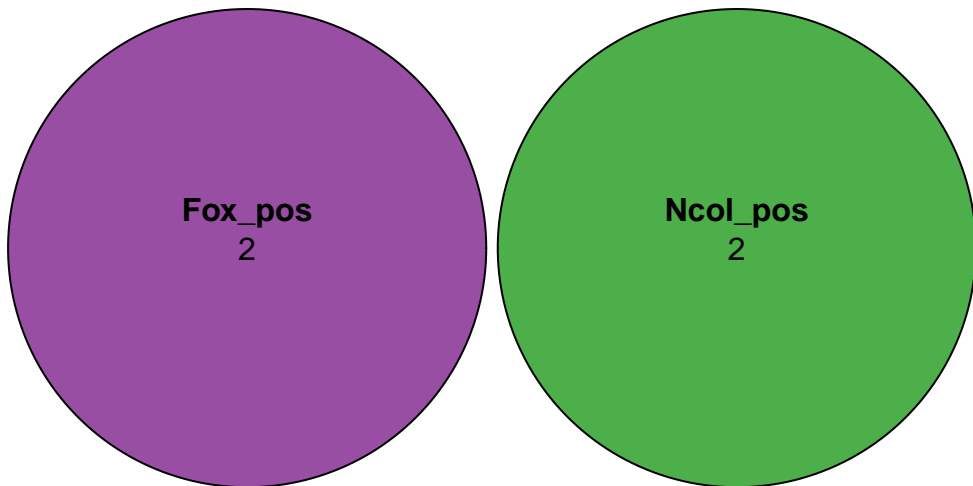


Nvec_vc1.1_XM_001633244.3

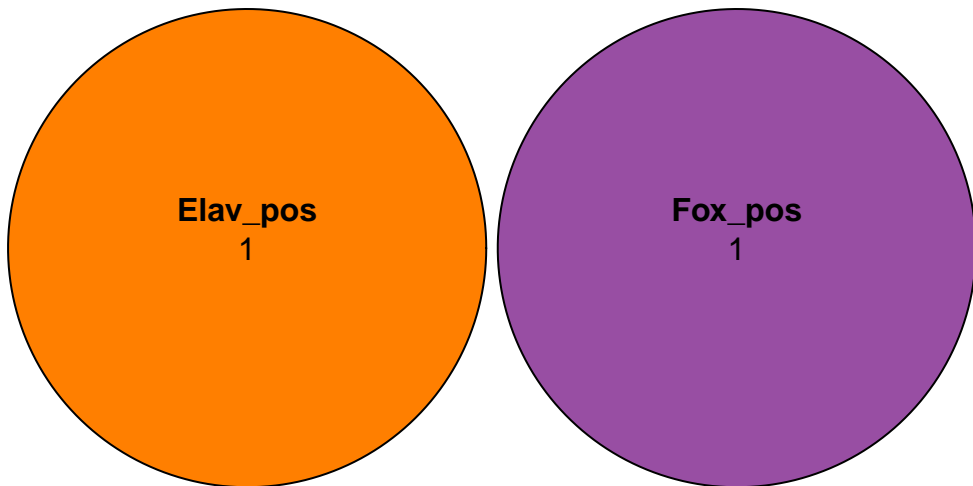
Synaptotagmin:C2CD4C/C2CD; Synaptotagmin.HG1.17:C2CD



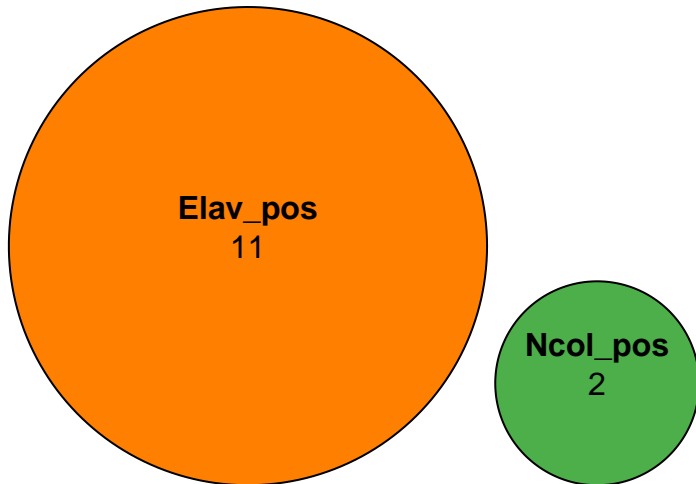
Nvec_vc1.1_XM_001639954.3
GPCR-rhodopsin:like:GPR27; 7tm_1/



Nvec_vc1.1_XM_032372557.2
Synaptotagmin:like:FO6814; C2/C2/

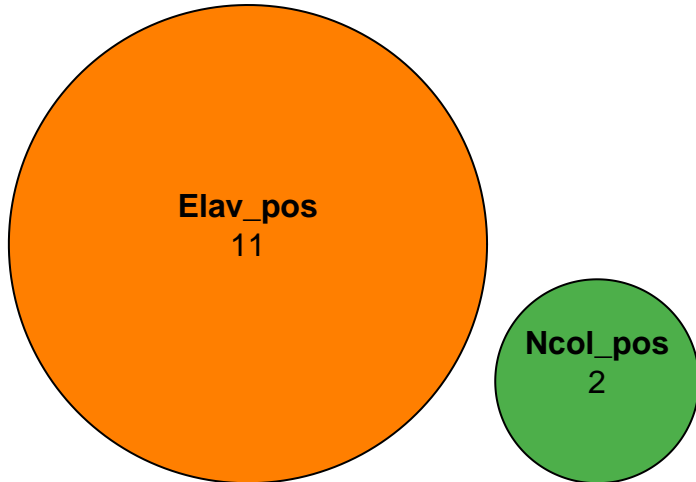


Nvec_vc1.1_XM_001636882.3
ADP-ribosylation factor-1; Arf/



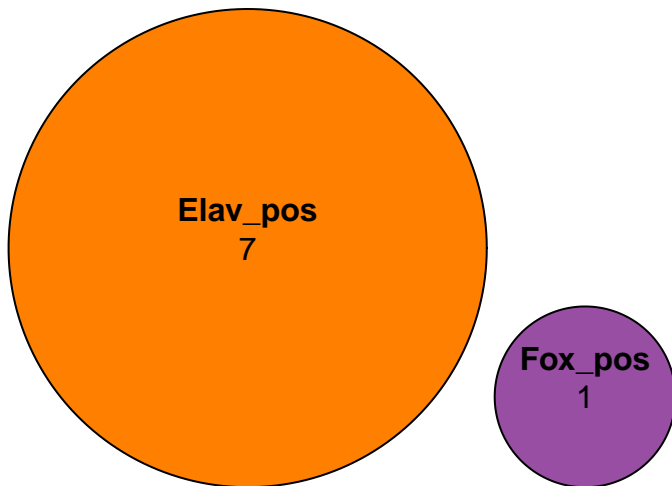
Nvec_vc1.1_XM_048719664.1

;



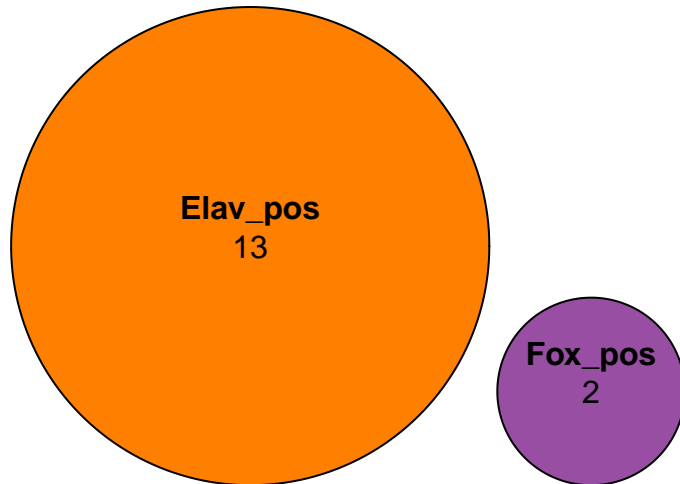
Nvec_vc1.1_XM_048725869.1

;



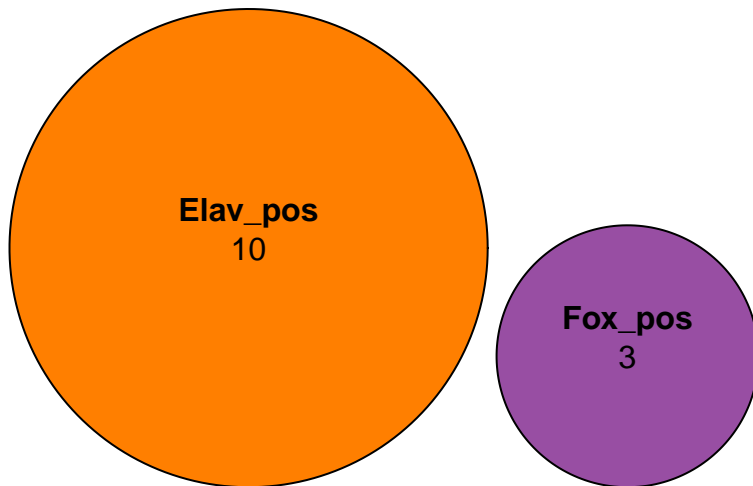
Nvec_vc1.1_XM_001632492.3

Alpha-(1,3)-fucosyltransf; Glyco_transf_10/



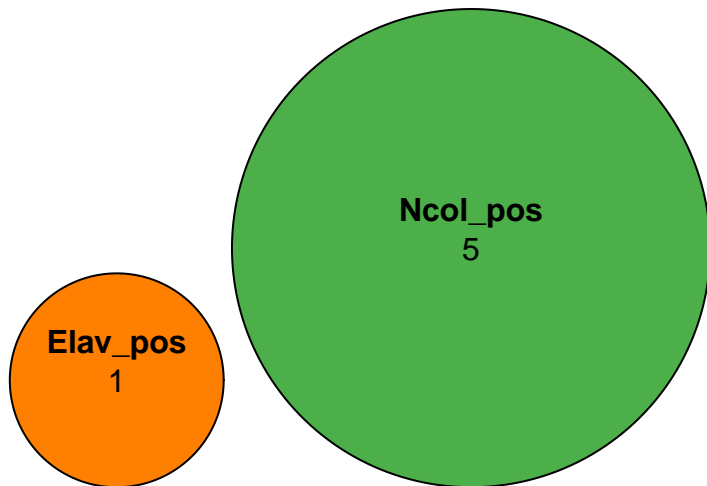
Nvec_vc1.1_XM_001639681.3

Potassium channel, subfam; lon_trans_2/lon_trans_2/



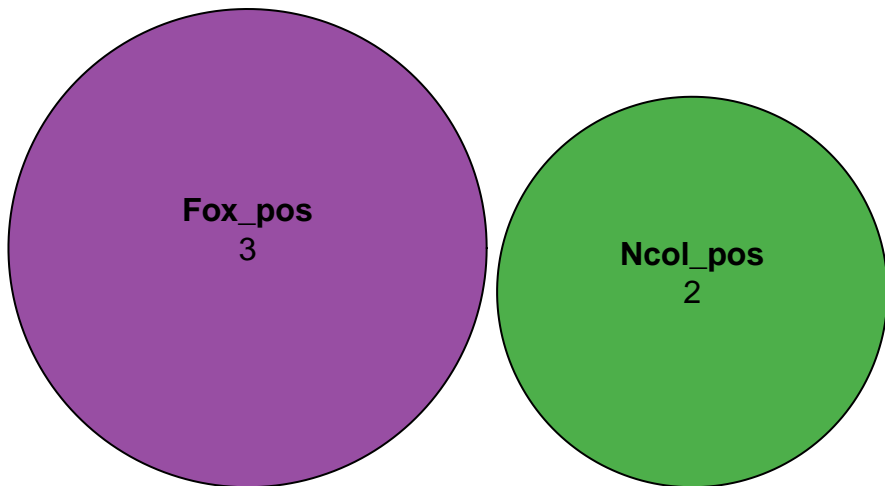
Nvec_vc1.1_XM_048729870.1

60 kDa lysophospholipase; Asparaginase/Ank_2/



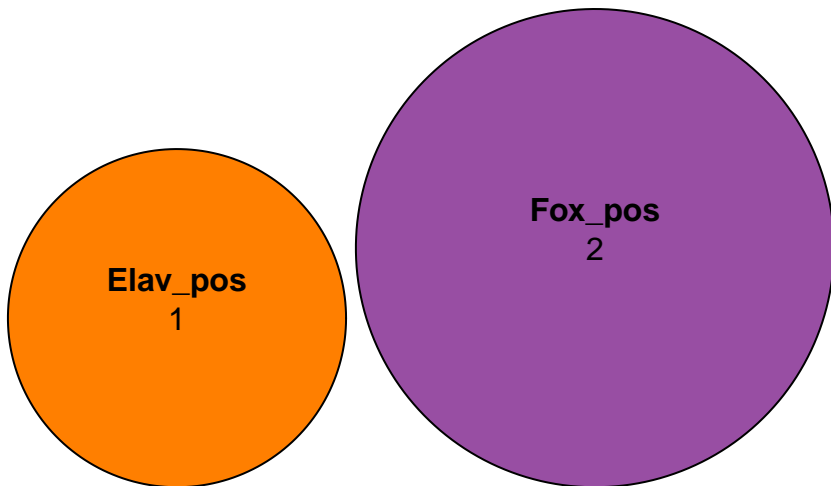
Nvec_NVE13928

;



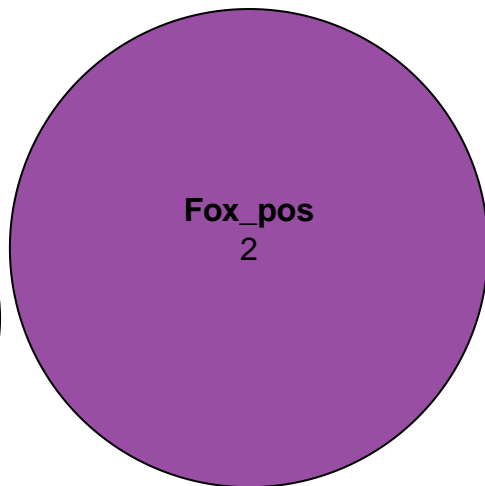
Nvec_vc1.1_XM_032364349.2

Aspartate aminotransferas; Aminotran_1_2/



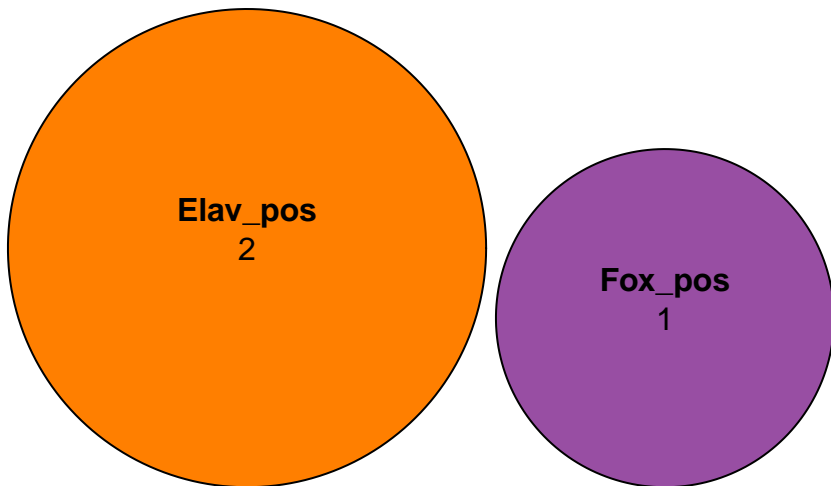
Nvec_vc1.1_XM_032376979.2

;

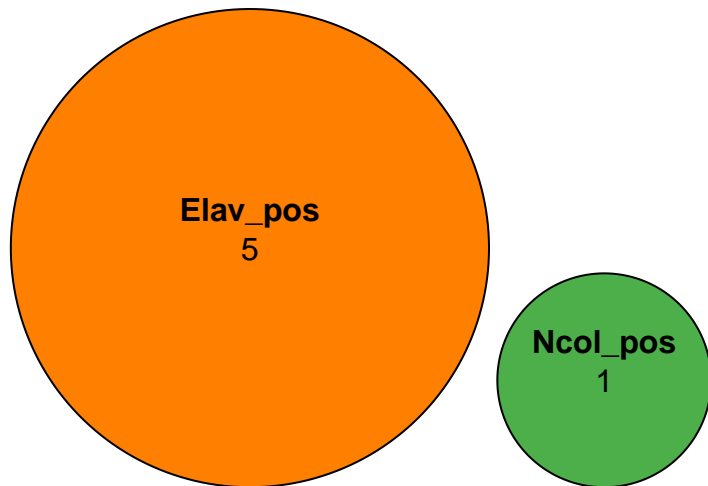


Nvec_vc1.1_XM_048730108.1

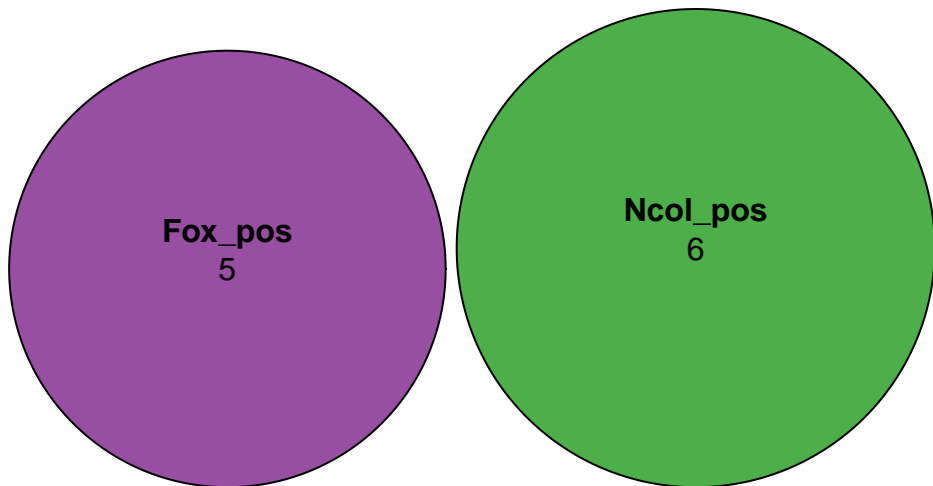
GPCR-rhodopsin:NA; 7tm_1/



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

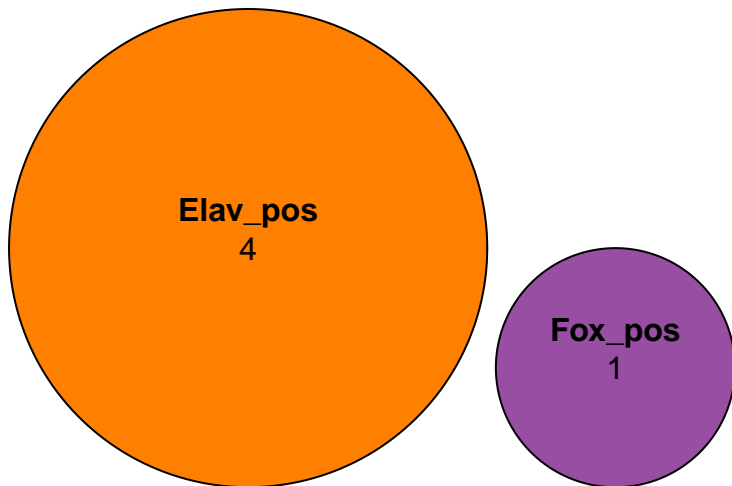


Nvec_vc1.1_XM_032382117.2
GPCR-rhodopsin:like:GPR17; 7tm_1/



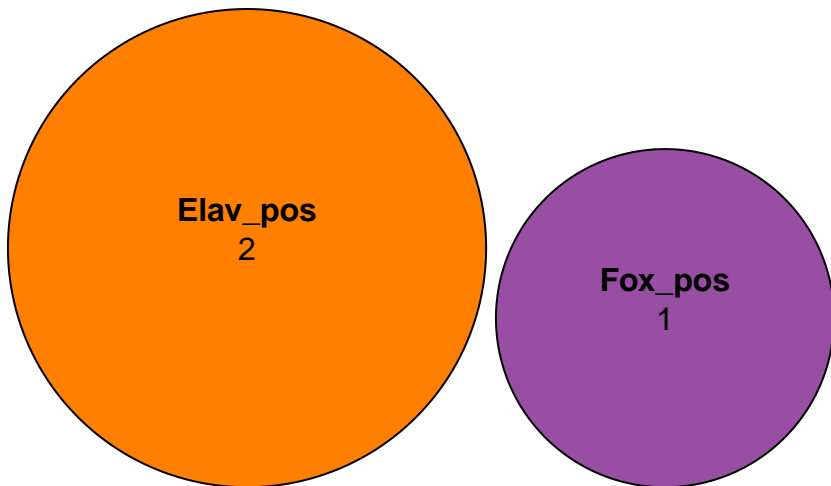
Nvec_vc1.1_XM_032386298.2

GPCR-rhodopsin:NA; 7tm_1/

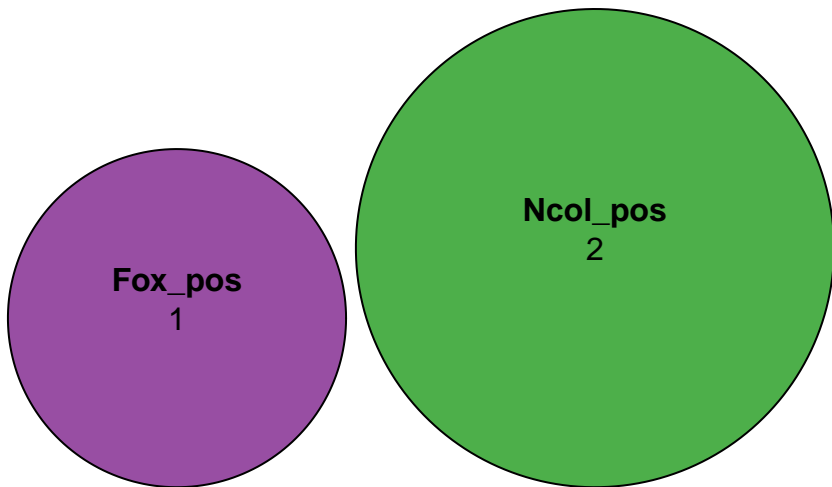


Nvec_vc1.1_XM_032362299.2

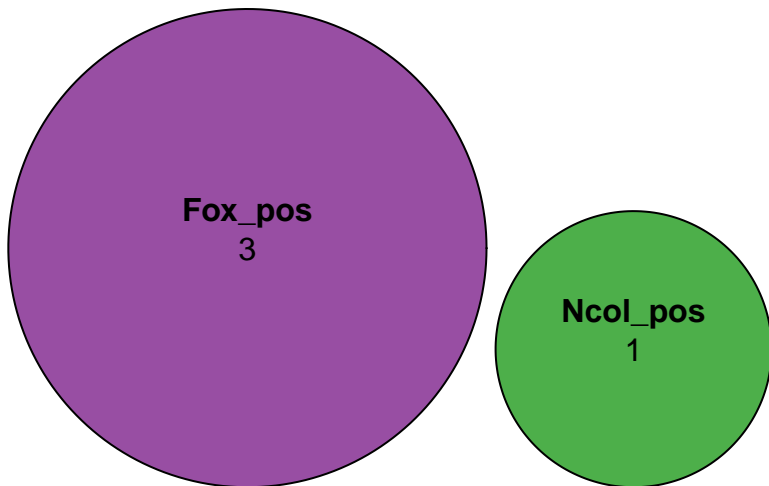
;



Nvec_vc1.1_XM_001633615.3
HCG2044780; Kua-UEV1_localn/

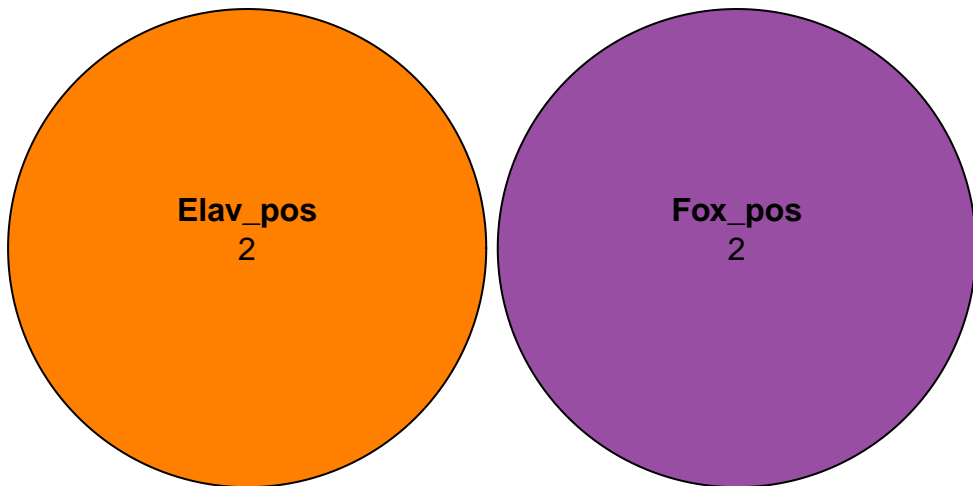


Nvec_vc1.1_XM_001636545.3
; RNase_P_Rpp14/

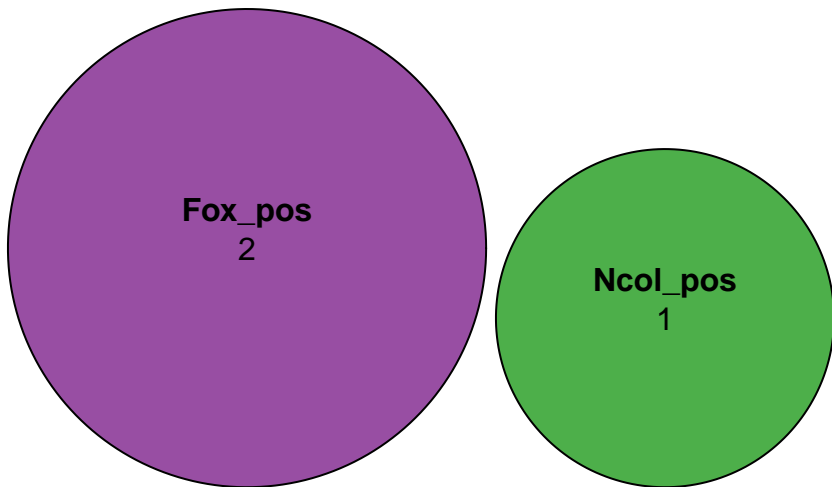


Nvec_vc1.1_XM_048720443.1

C-Maf-inducing protein;

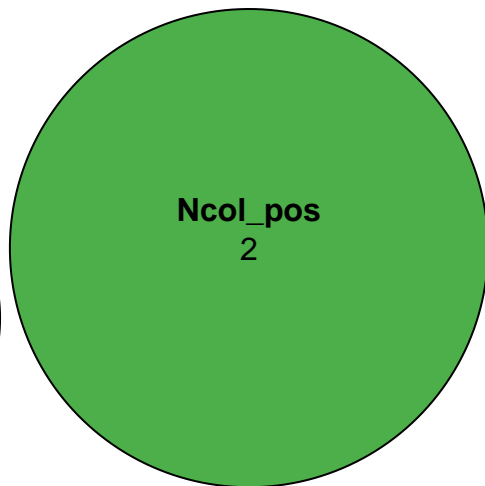


Nvec_vc1.1_XM_032384695.2
Monocarboxylate transport; MFS_1/



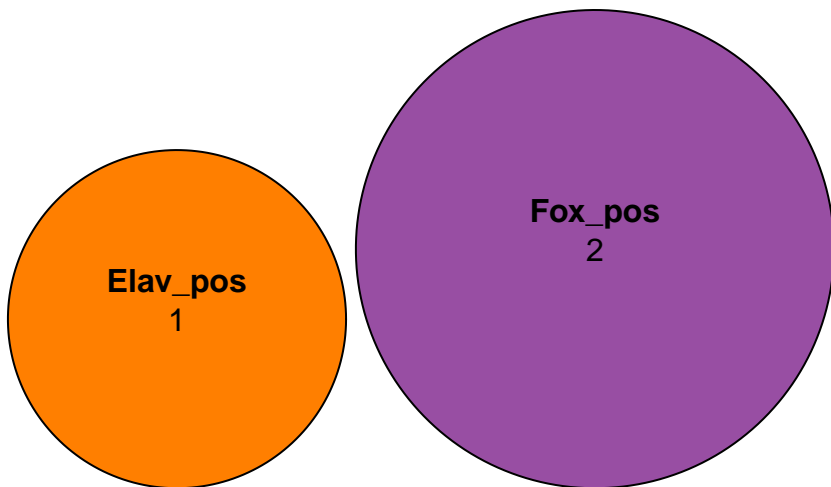
Nvec_vc1.1_XM_048731044.1

;

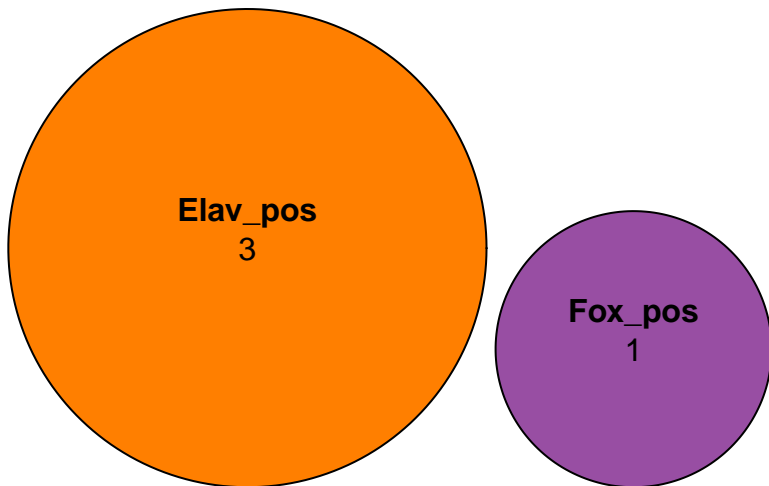


Nvec_vc1.1_XM_001629129.3

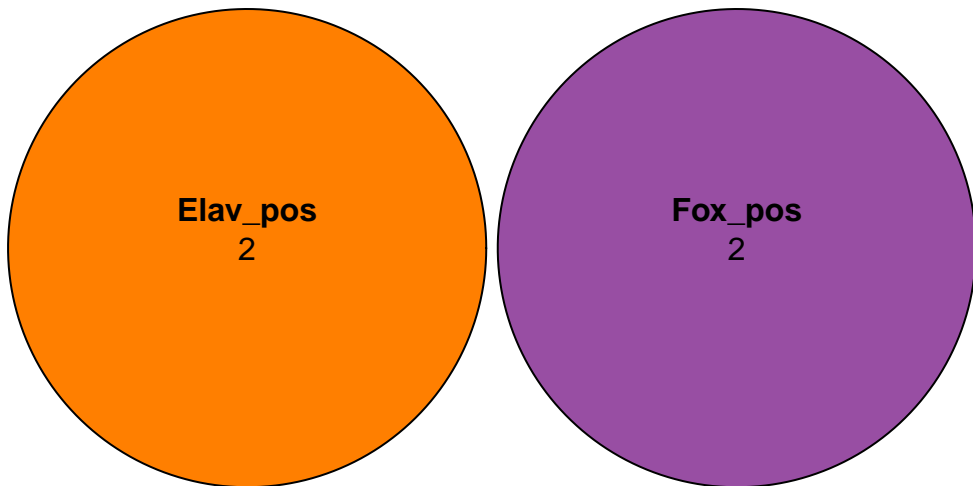
Probable aminopeptidase N; Peptidase_M17/



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

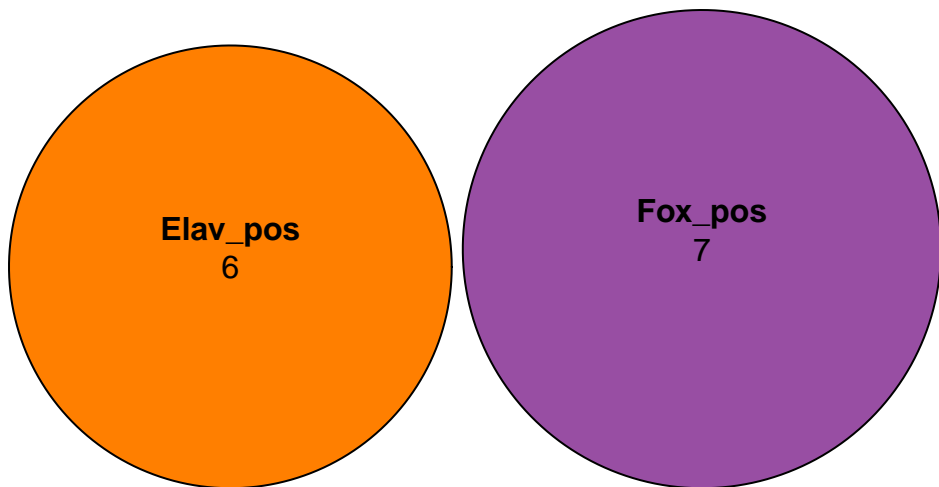


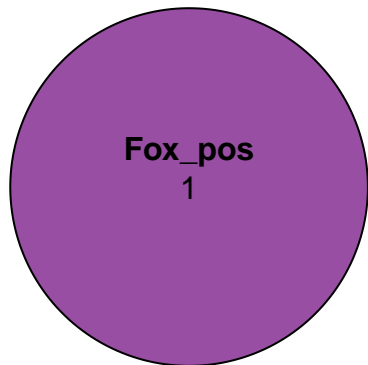
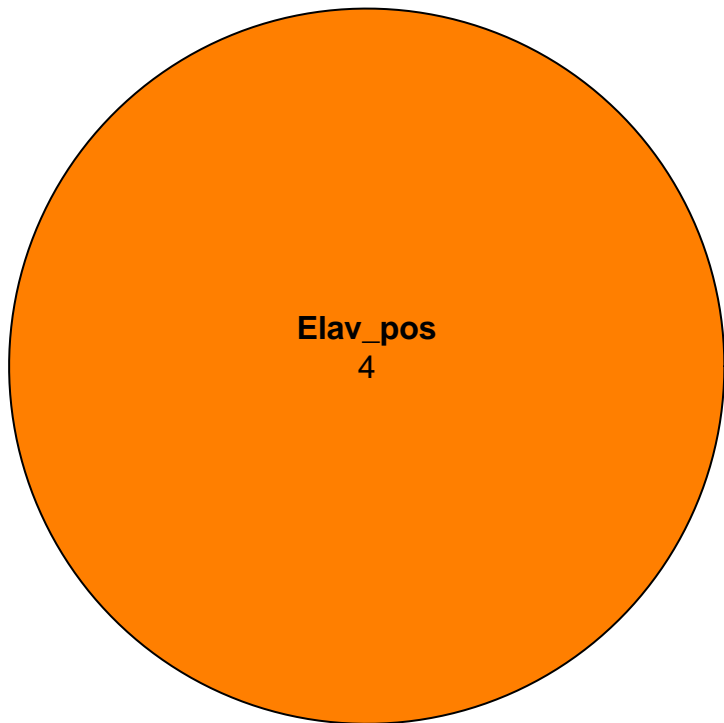
Nvec_vc1.1_XM_032373148.2
S1:SUPT6H; S1.HG5.0:SUPT6H



Nvec_vc1.1_XM_032381433.2

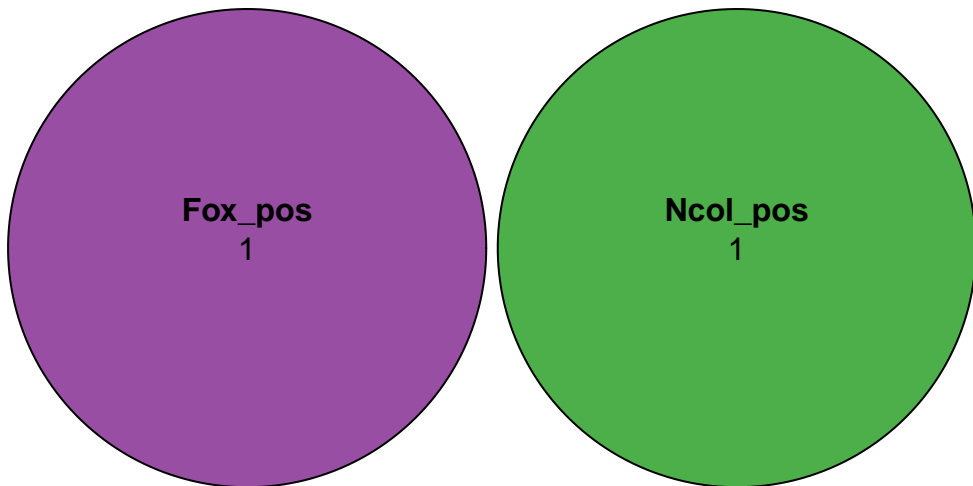
Guanylate cyclase soluble; HNOB/HNOBA/Guanylate_cyc/





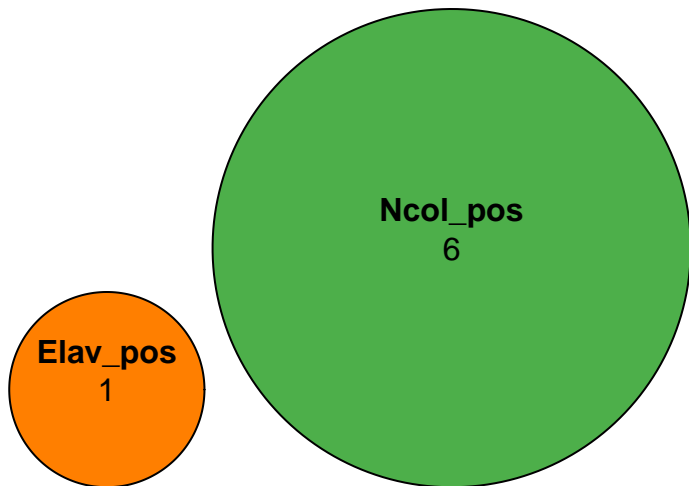
Nvec_NVE7893

;



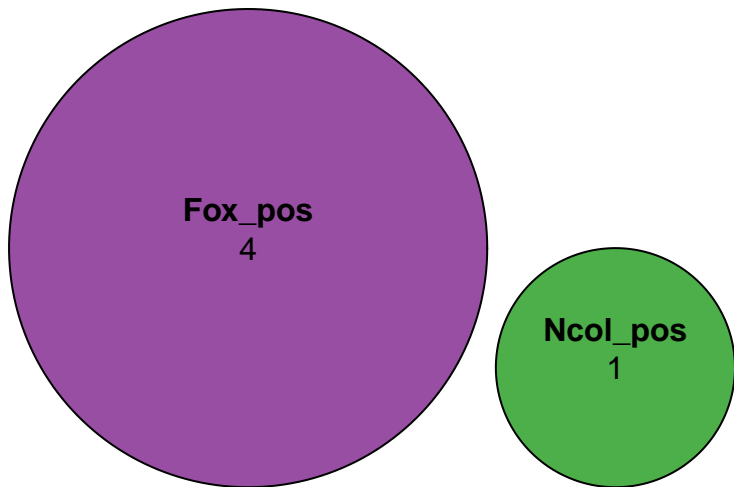
Nvec_vc1.1_XM_032372214.2

;



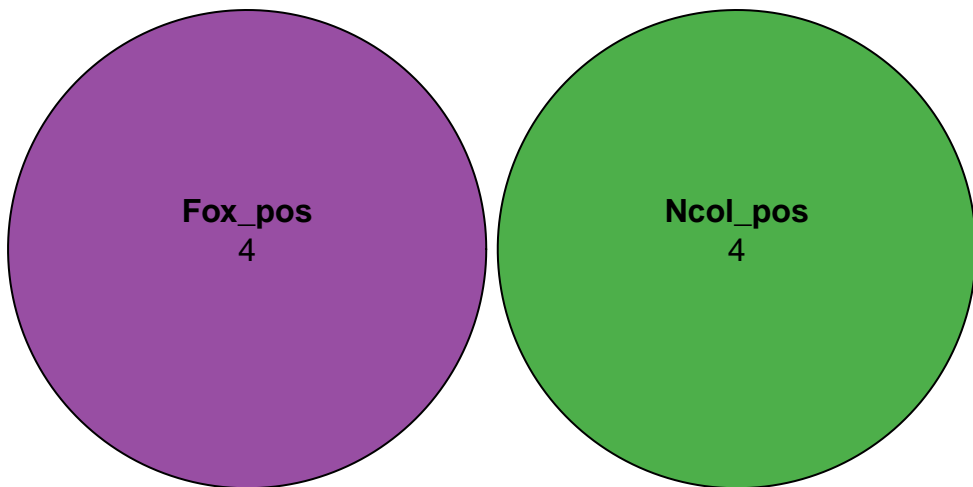
Nvec_vc1.1_XM_048730962.1

;



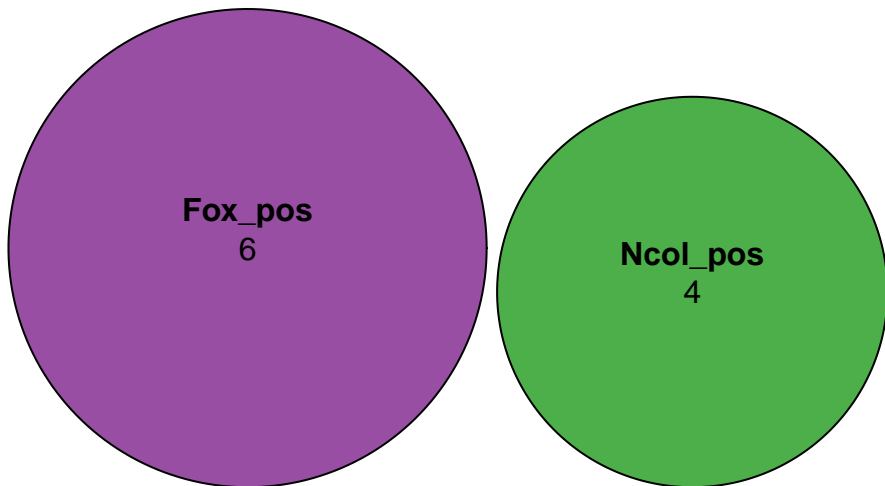
Nvec_vc1.1_XM_001637616.3

Fukutin related protein; LicD/



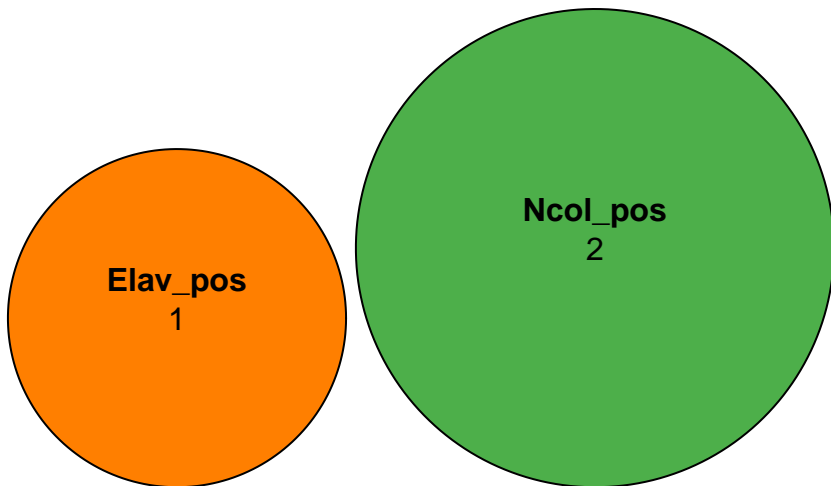
Nvec_vc1.1_XM_001628059.3

Fatty acid amide hydrolas; Amidase/

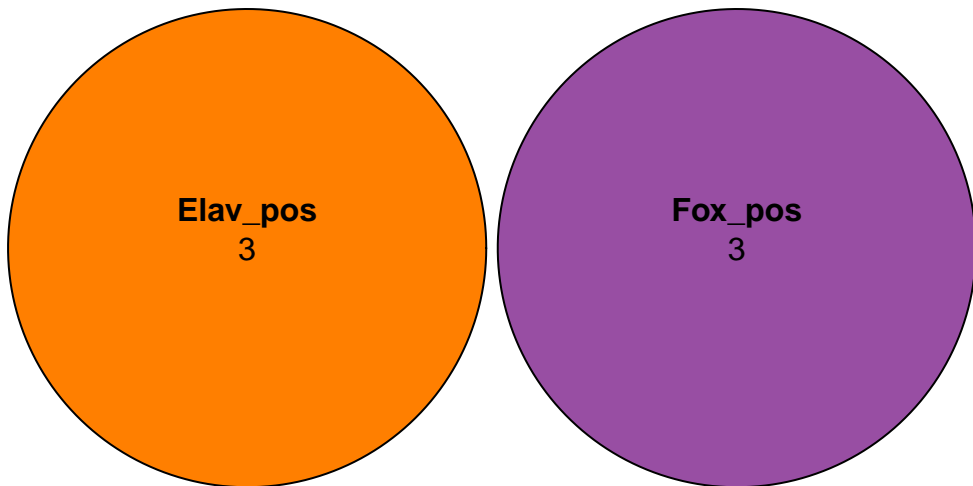


Nvec_vc1.1_XM_048730184.1

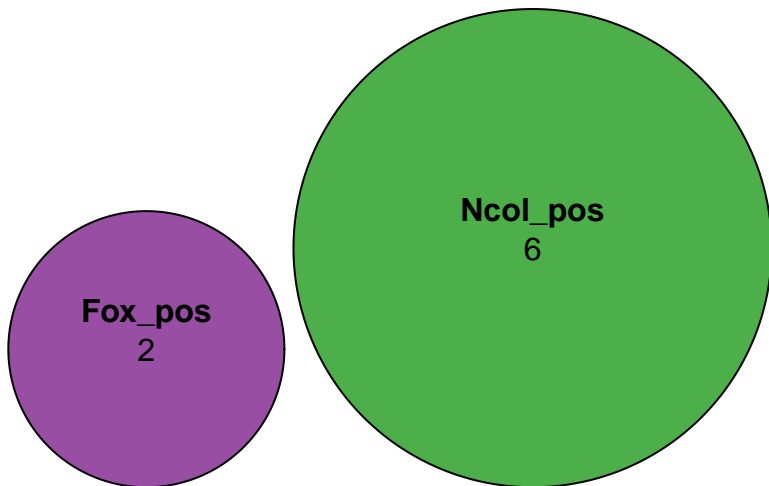
Thyroid receptor-interact;



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

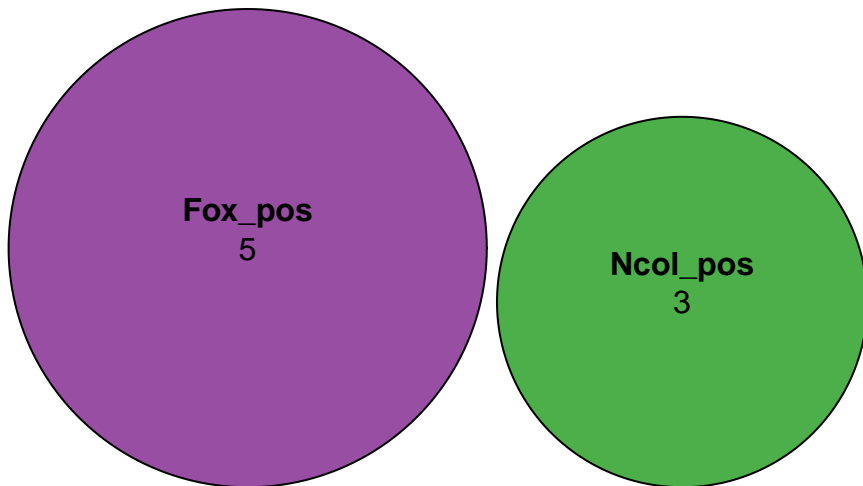


Nvec_vc1.1_XM_001641783.3
mTERF.HG1.0:MTERF4; MTERF4



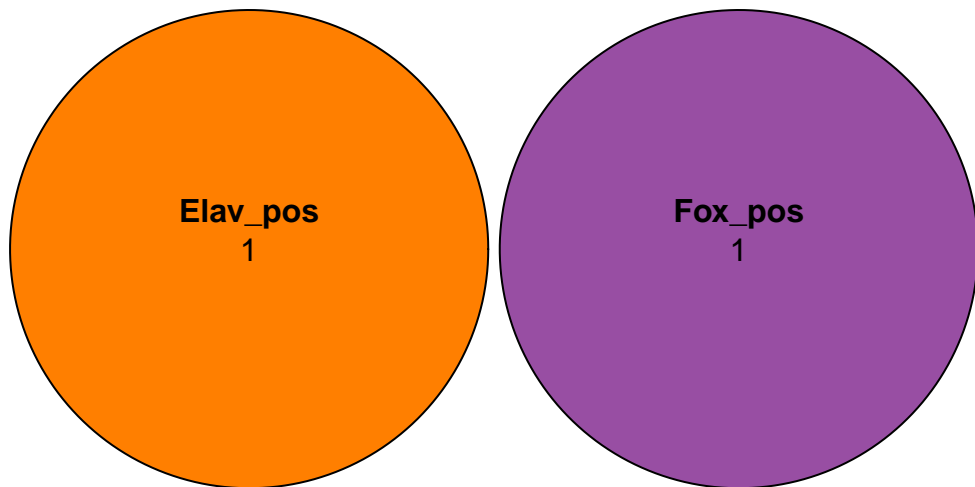
Nvec_vc1.1_XM_032374380.2

;



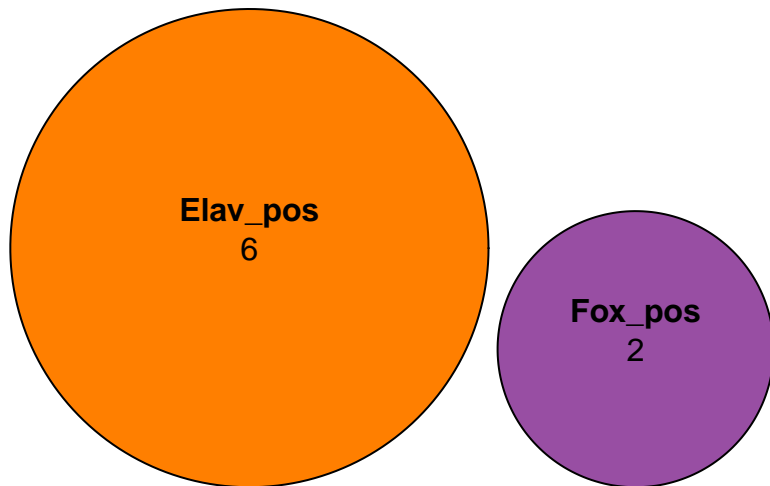
Nvec_vc1.1_XM_032384990.2

Protein odr-4 homolog; ODR4-like/ODR4-like/



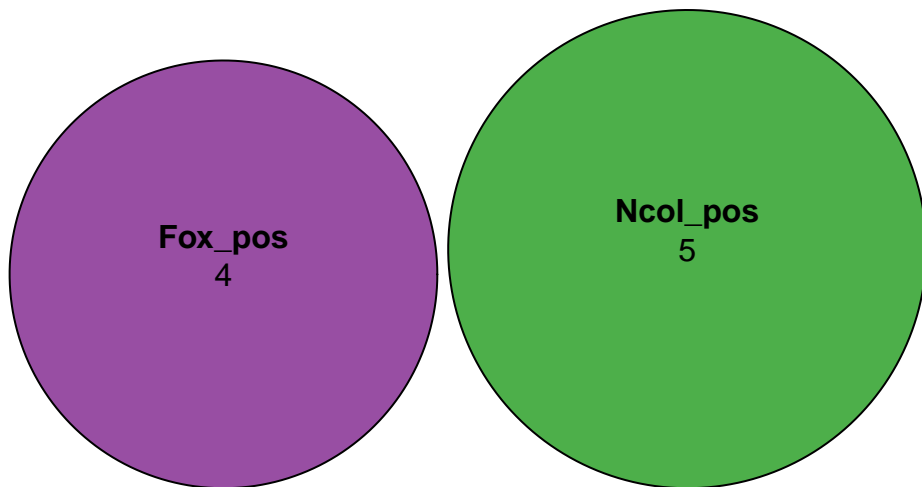
Nvec_vc1.1_XM_048729479.1

TAF5-like RNA polymerase ; TFIID_90kDa/WD40/WD40/WD40



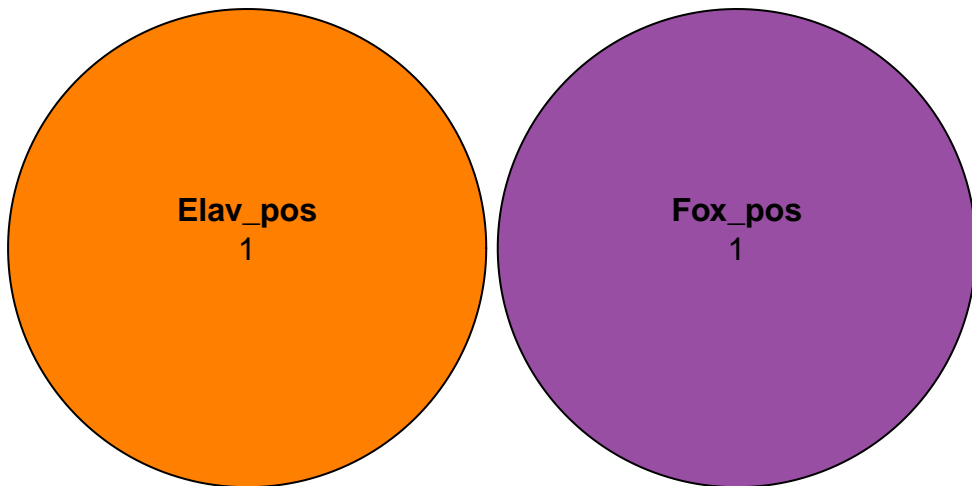
Nvec_vc1.1_XM_001639817.3

Ephrin:AL691442/EFNA1/EFN; Ephrin.HG1.0:AL691442/EFN



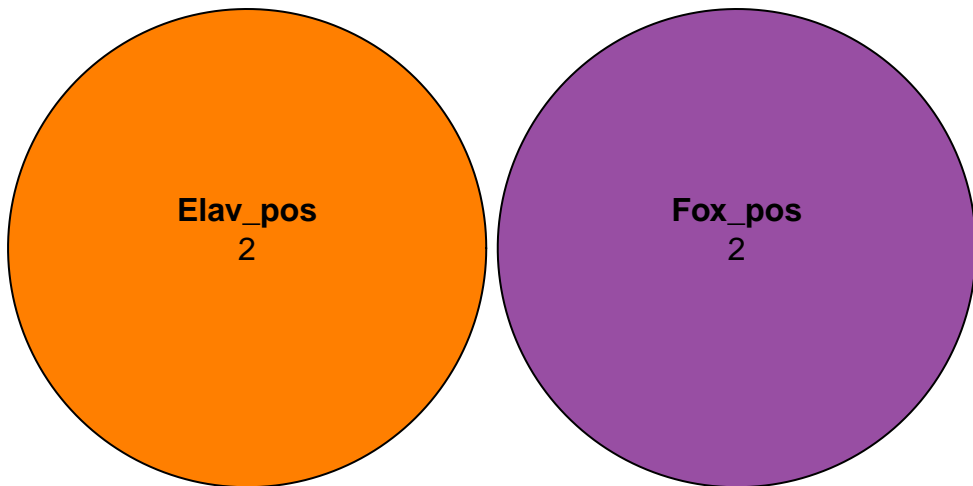
Nvec_vc1.1_XM_032366492.2

N-acetylated-alpha-linked; TFR_dimer/



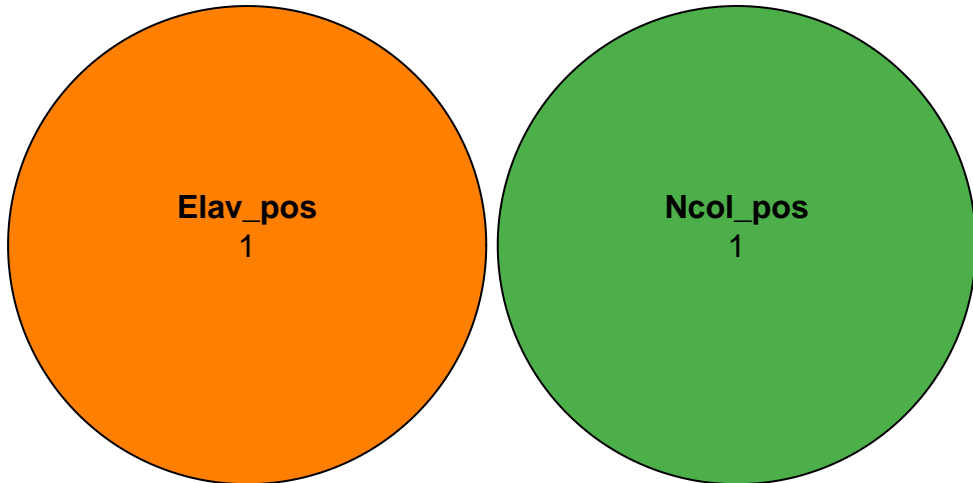
Nvec_vc1.1_XM_048729868.1

E3 ubiquitin–protein liga;

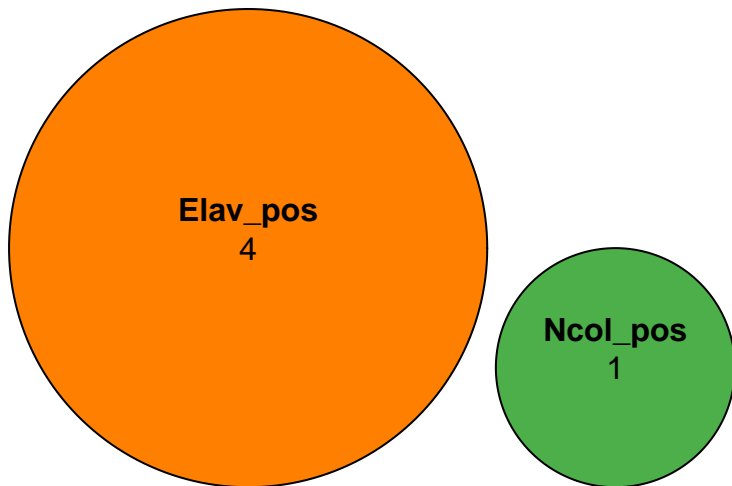


Nvec_vc1.1_XM_048734712.1

Very long-chain-specific ;

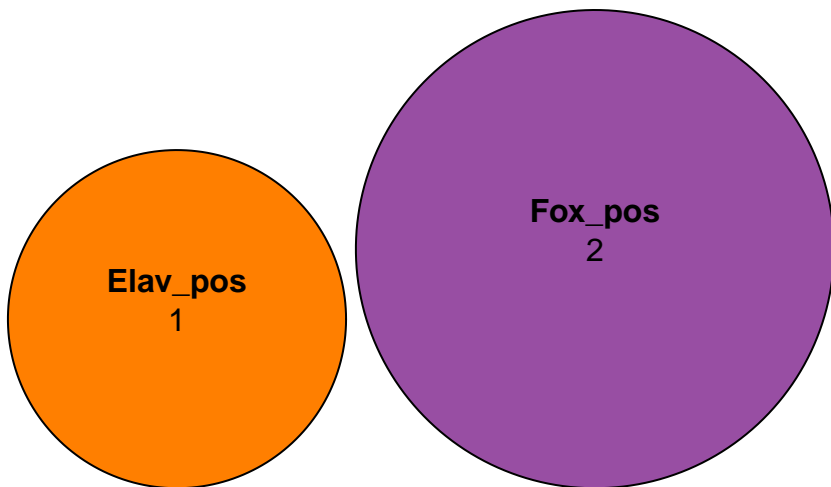


Nvec_vc1.1_XM_001628363.3
cDNA, FLJ94698, highly si; Sybindin/

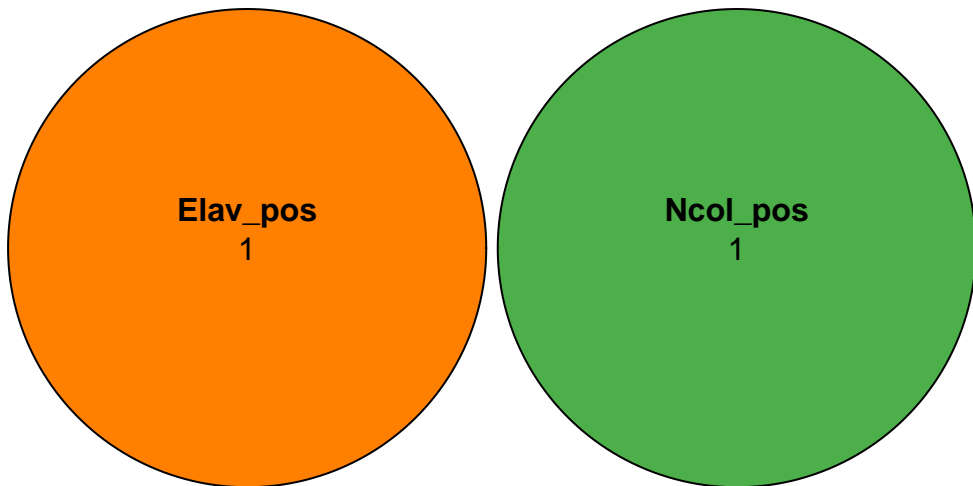


Nvec_vc1.1_XM_032387176.2

LMBR1 domain-containing p; LMBR1/

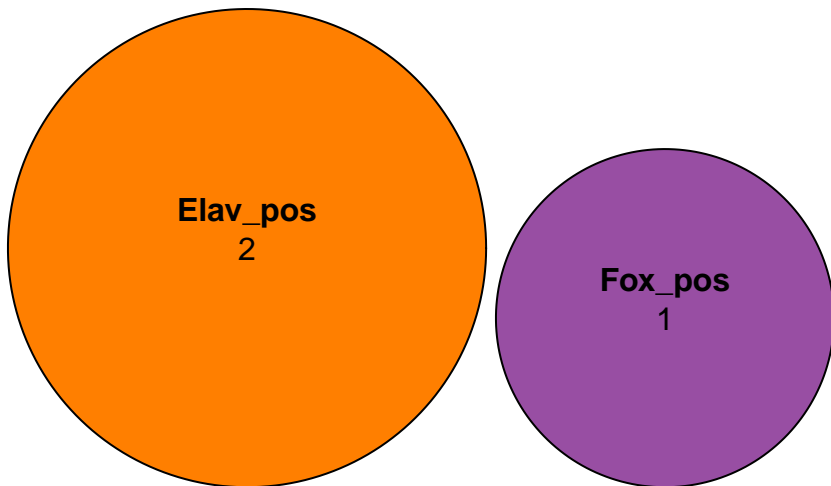


Nvec_vc1.1_XM_001631292.3
ADP-ribosylation factor r;

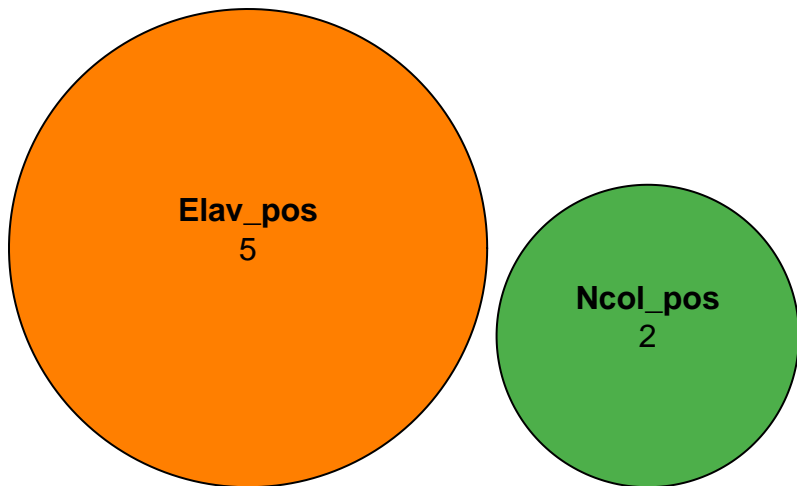


Nvec_vc1.1_XM_001631286.2

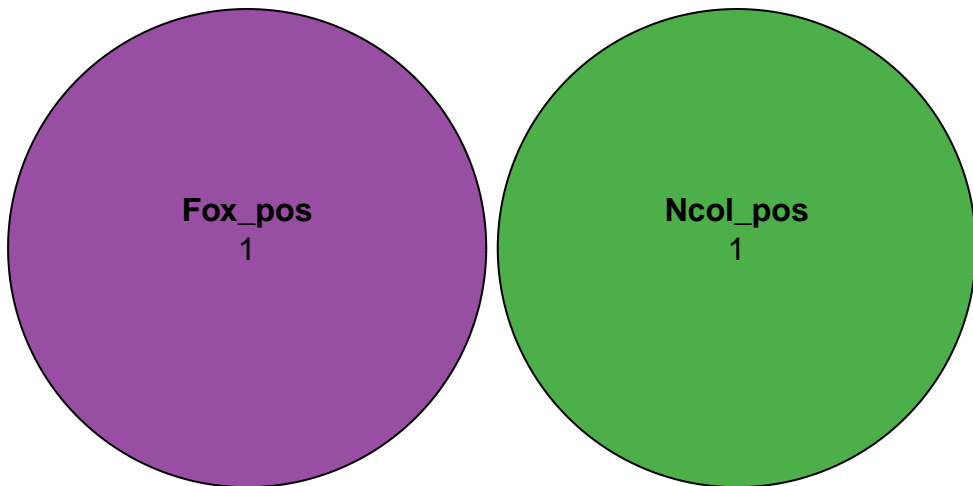
NNMT protein; NNMT_PNMT_TEMT/



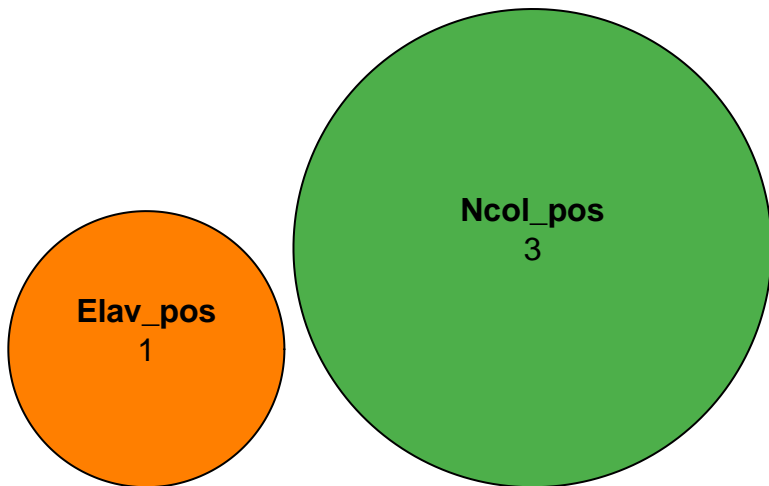
Nvec_vc1.1_XM_001624997.3
; F-box-like/



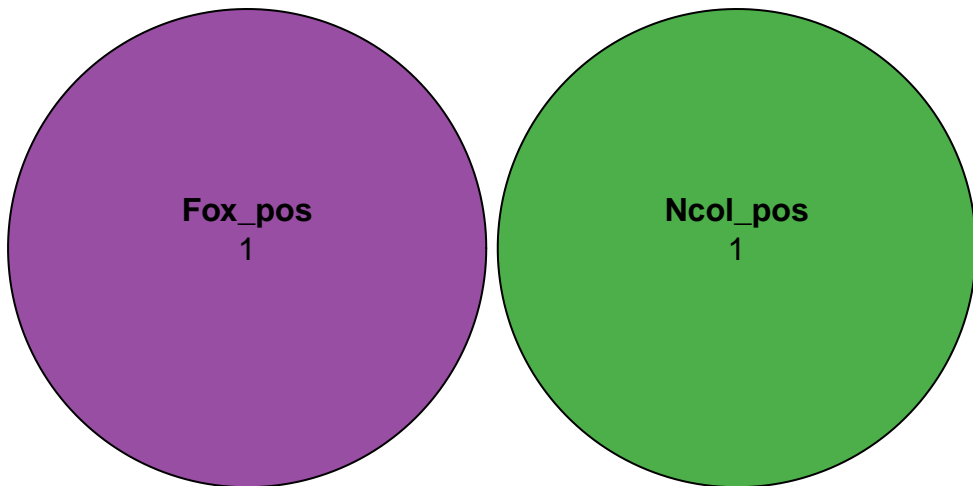
Nvec_vc1.1_XM_001641286.3
cDNA FLJ51917, highly sim; TIMP/



Nvec_vc1.1_XM_032368559.2
Protein prenyltransferase; PPTA/

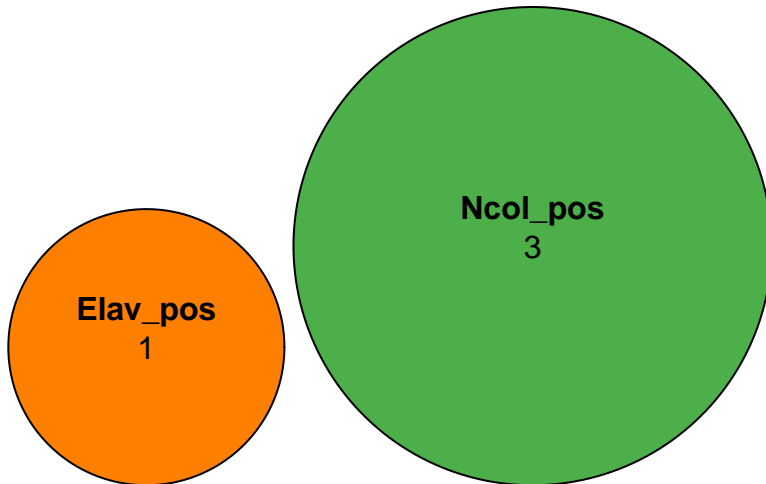


Nvec_vc1.1_XM_048727242.1
Zinc transporter 7; Cation_efflux/



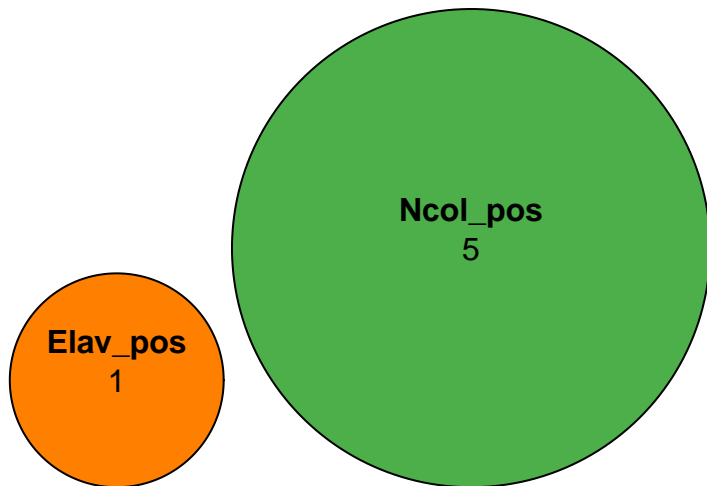
Nvec_vc1.1_XM_032372191.2

Abhydrolase domain contai; Abhydrolase_6/



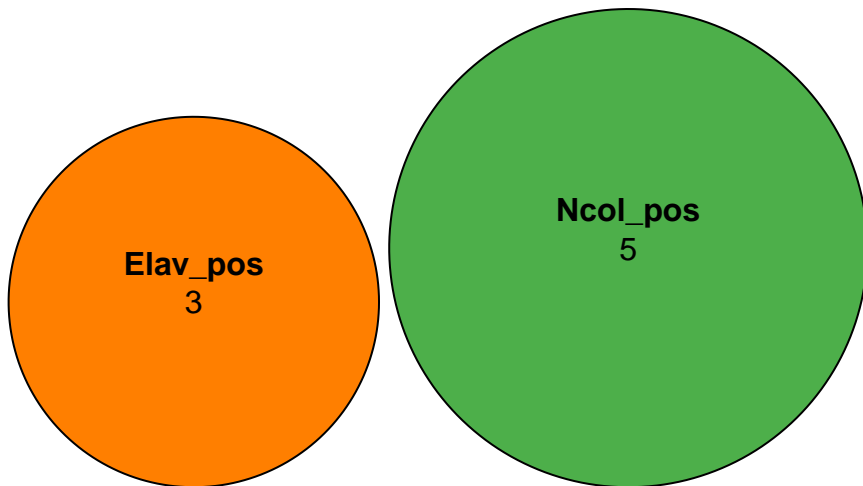
Nvec_vc1.1_XM_048730264.1

Component of oligomeric g; COG2/DUF3510/



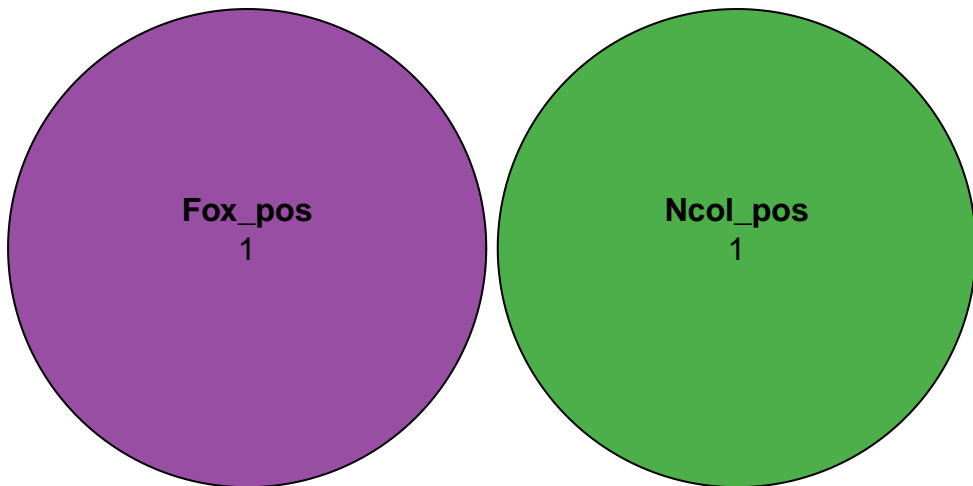
Nvec_vc1.1_XM_032375751.2

Transmembrane protein 115; DUF1751/

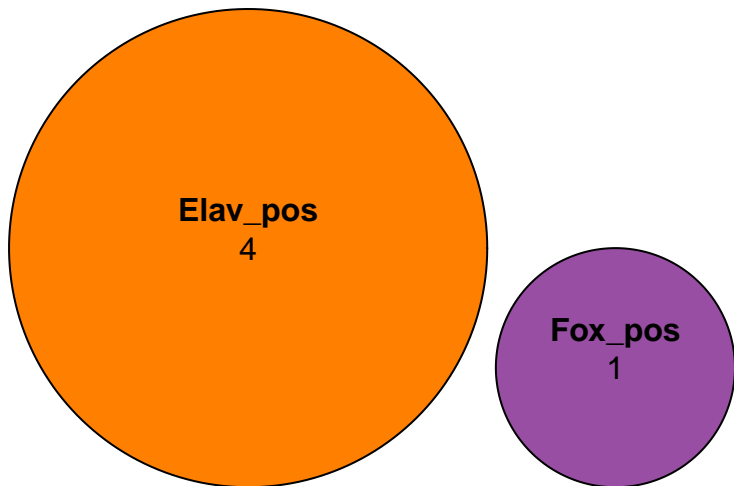


Nvec_vc1.1_XM_048725016.1

;

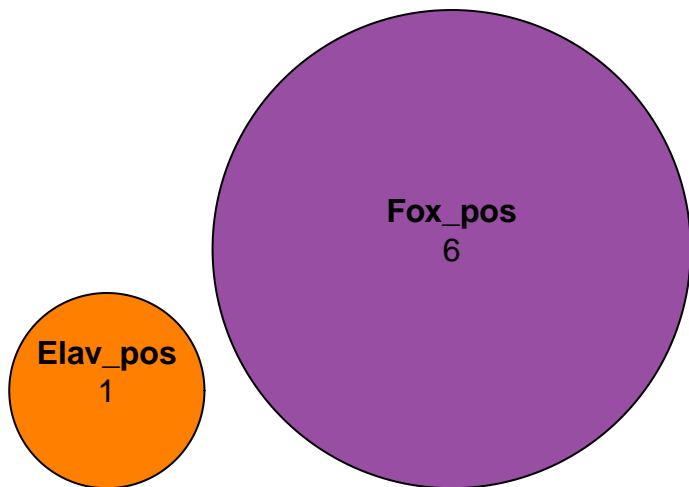


Nvec_vc1.1_XM_001638793.3
Zinc finger CCHC domain-c; PSP/



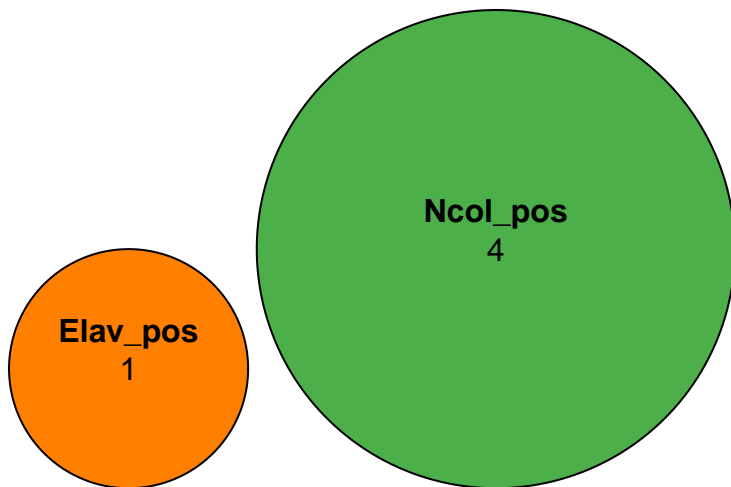
Nvec_vc1.1_XM_032363581.2

GPCR-rhodopsin:NA; 7tm_1/

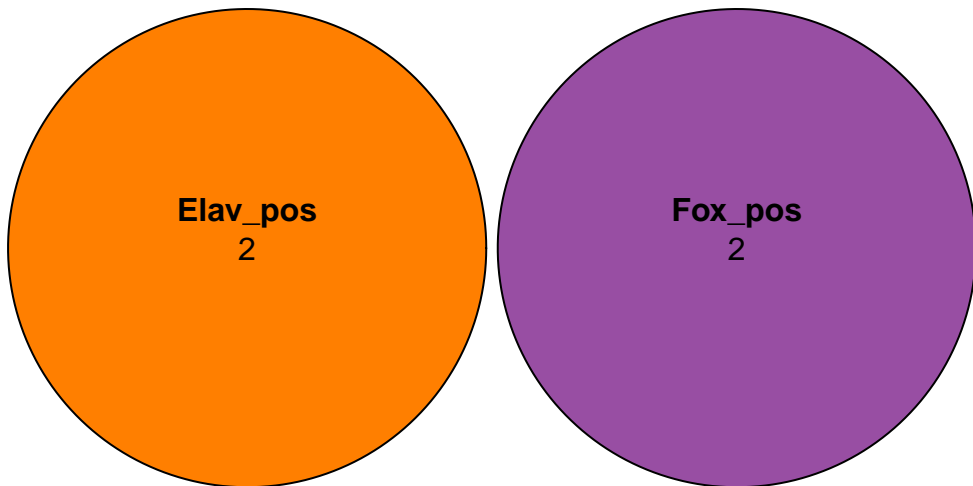


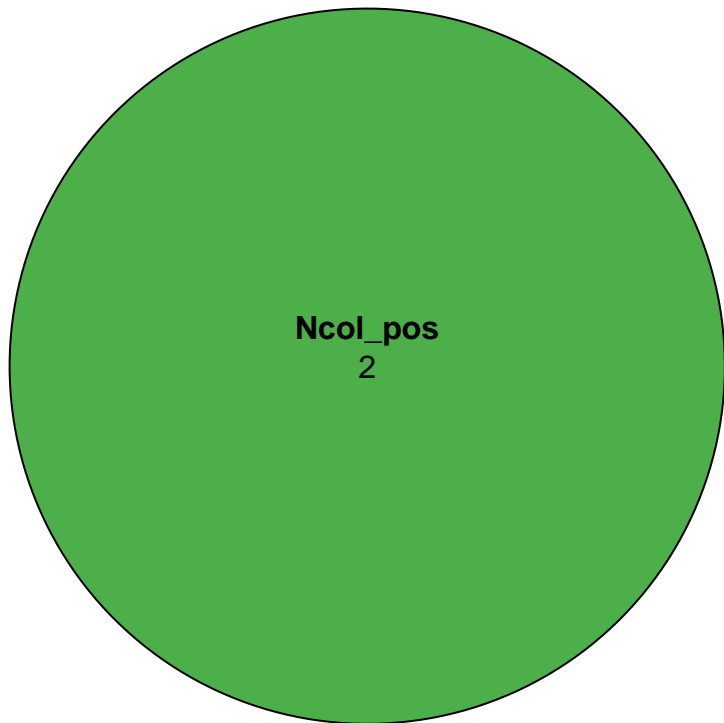
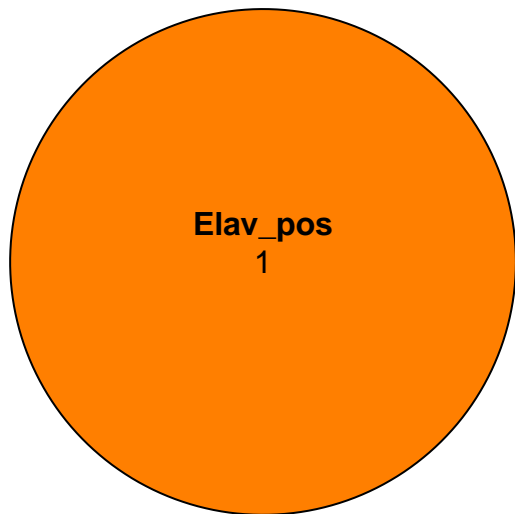
Nvec_vc1.1_XM_032364770.2

GPCR-rhodopsin:NA; 7tm_1/



Nvec_vc1.1_XM_032367592.2
cDNA, FLJ79298, highly si;





Nvec_v1g243589

;

Fox_pos

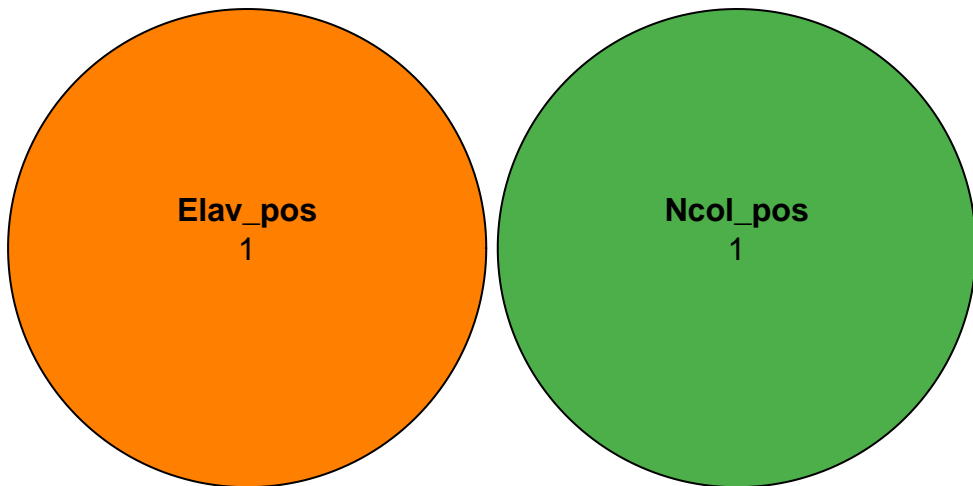
1

Ncol_pos

1

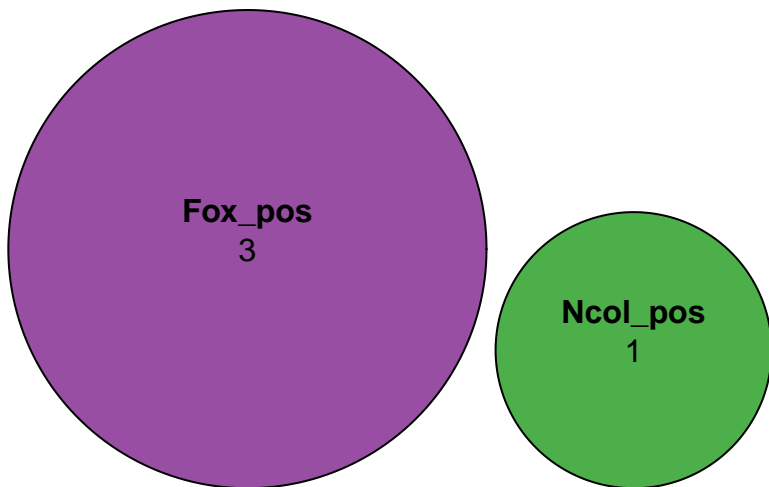
Nvec_vc1.1_XM_032385089.2

Ets.HG1.7:GABPA; GABPA



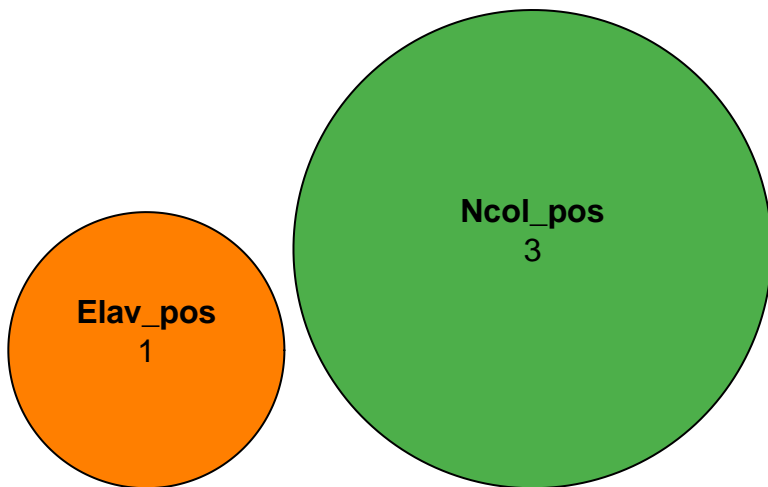
Nvec_vc1.1_XM_048720705.1

Ubiquitin carboxyl-termin;



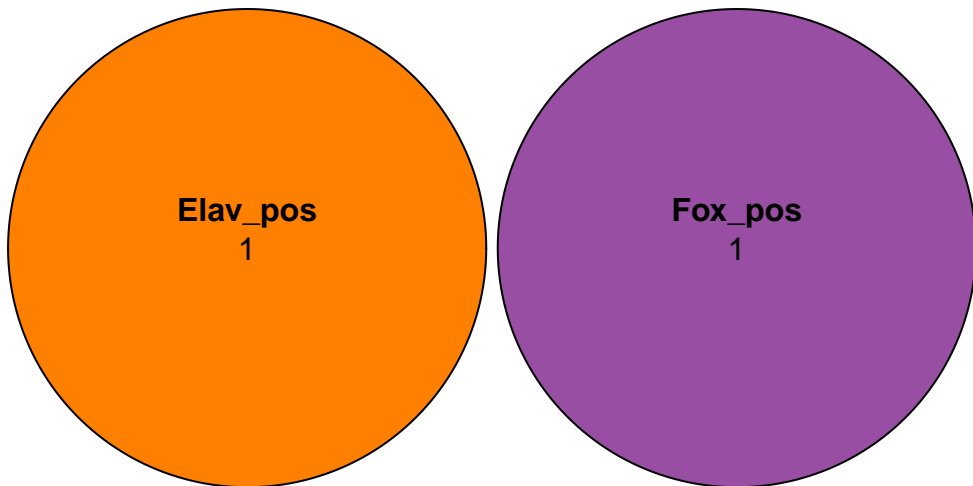
Nvec_vc1.1_XM_032362554.2

Synembryn-A; Ric8/

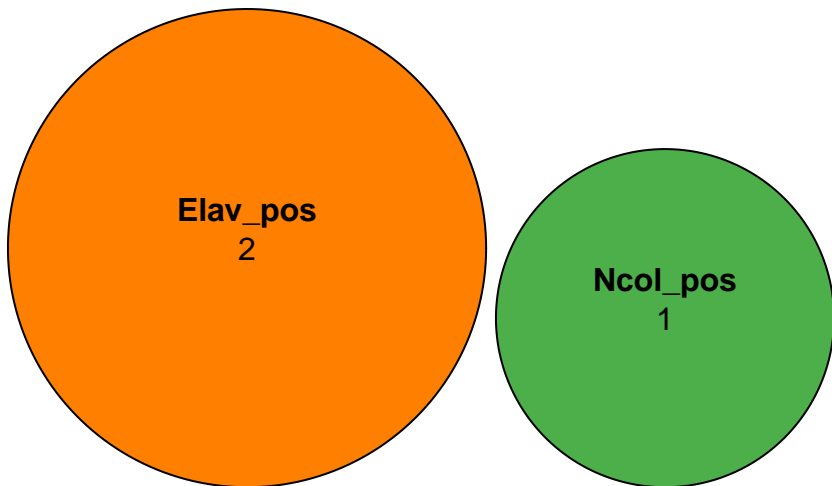


Nvec_vc1.1_XM_048726963.1

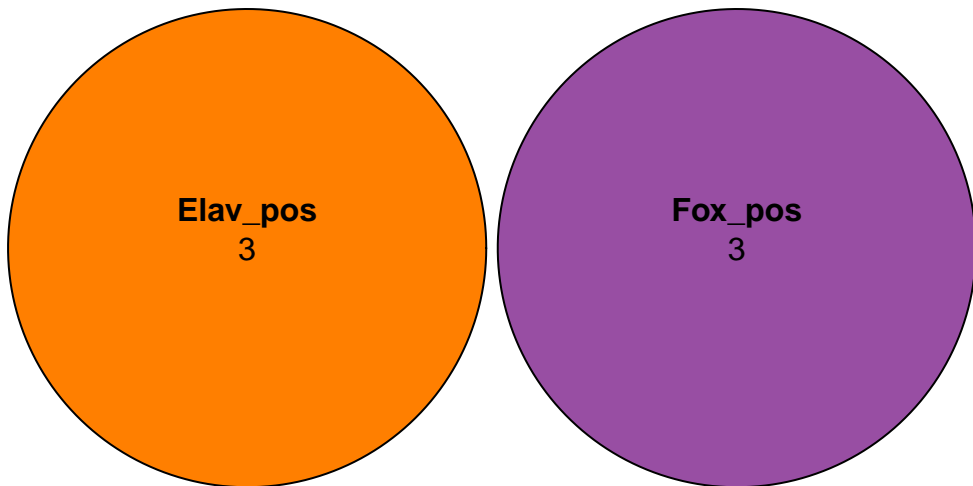
;



Nvec_vc1.1_XM_032382564.2
Uncharacterized protein; TMEM219/



Nvec_vc1.1_XM_001630938.3
TM2 domain-containing pro; TM2/



Nvec_vc1.1_XM_048730070.1

FAM179B protein;

