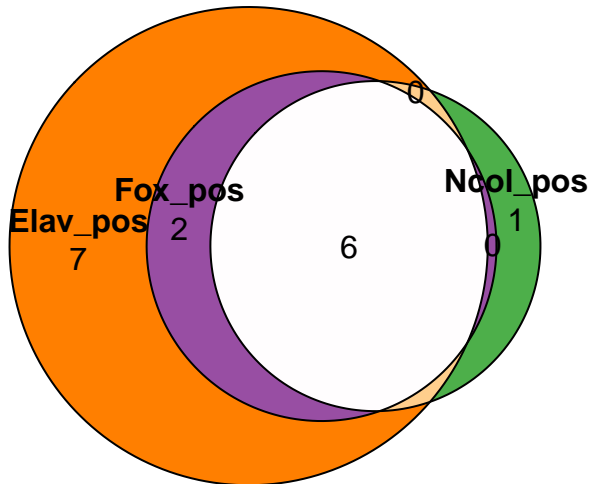


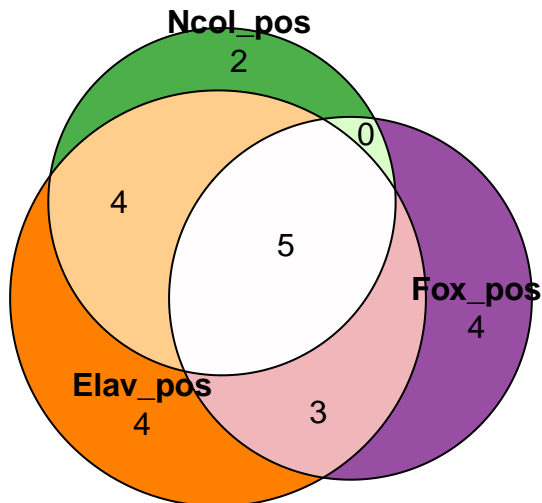
Nvec_vc1.1_XM_048732187.1

; Exo_endo_phos_2/

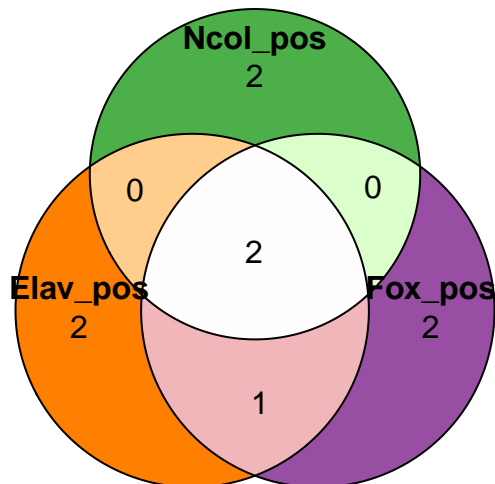


Nvec_vc1.1_XM_032373624.2

ZIP.HG1.2:BACH1/BACH2/NF; BACH1/BACH2/NFE2/NFE2L1/

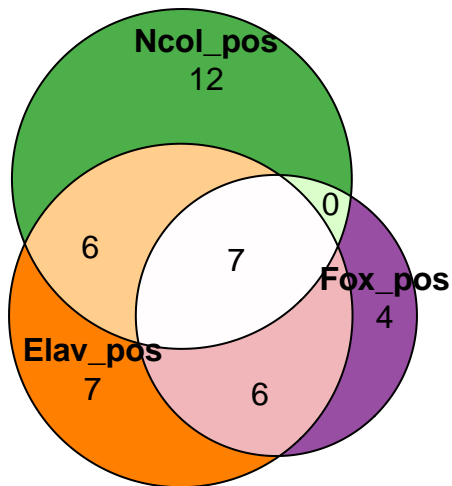


Nvec_vc1.1_XM_001638266.3
bZIP.HG1.0:CREBL2; CREBL2



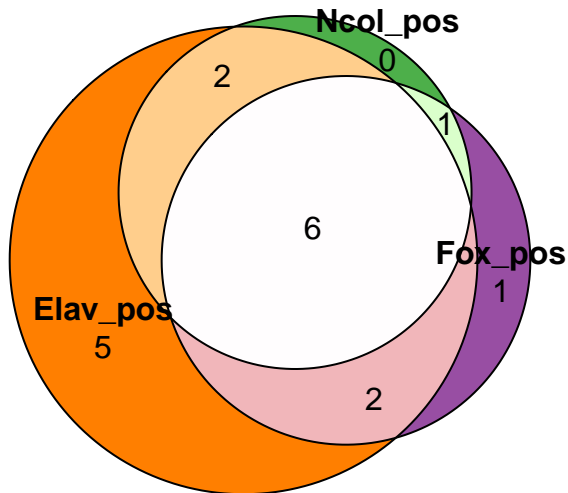
Nvec_vc1.1_XM_032364329.2

HSF.HG1.5:HSF1/HSF2/HSF4; HSF1/HSF2/HSF4



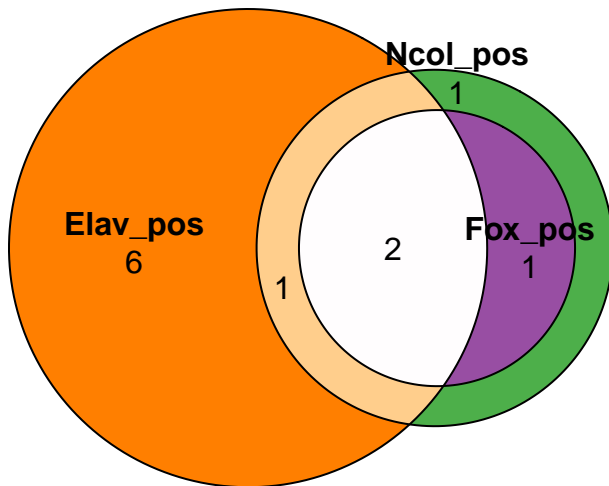
Nvec_vc1.1_XM_048733302.1

HLH.HG1.33:MITF/TFE3/TFEB; MITF/TFE3/TFEB/TFEC



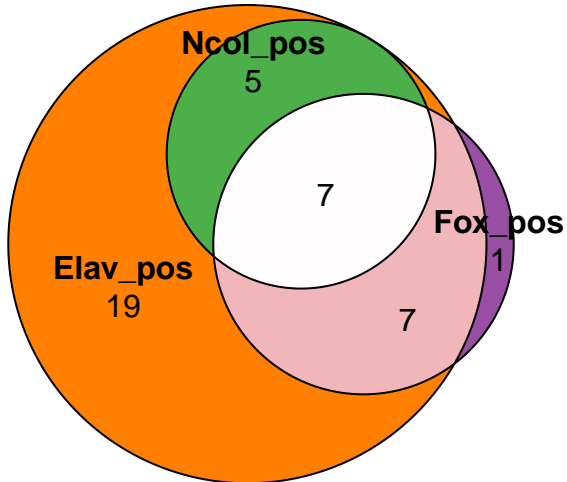
Nvec_NVE17130

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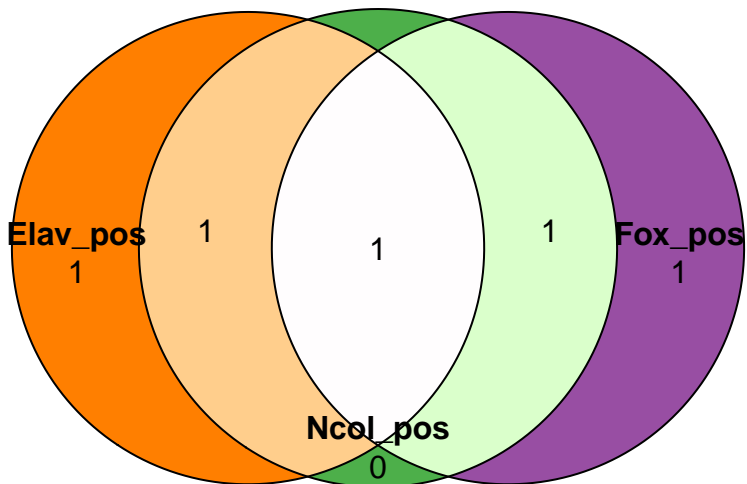
Nvec_vc1.1_XM_001634398.3

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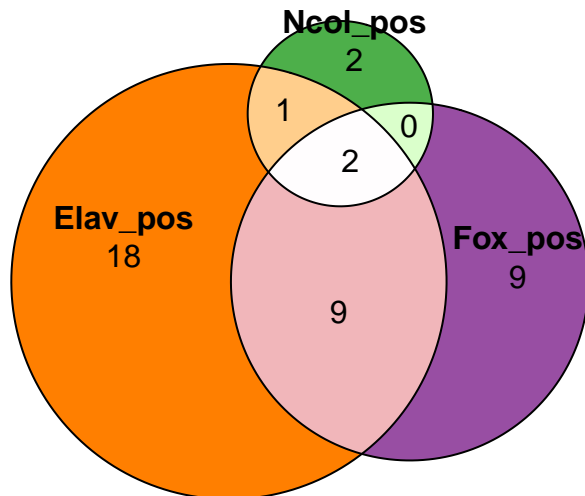
Nvec_vc1.1_XM_001629629.3

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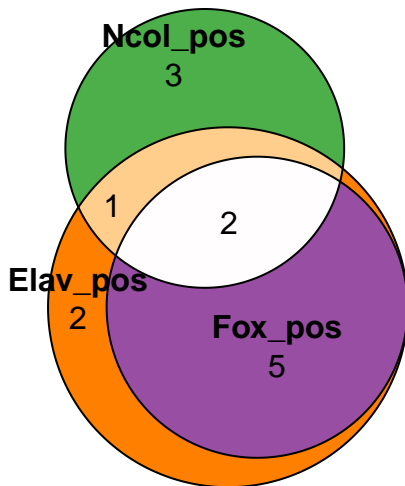
Nvec_vc1.1_XM_032364320.2

Epithelial splicing regul;



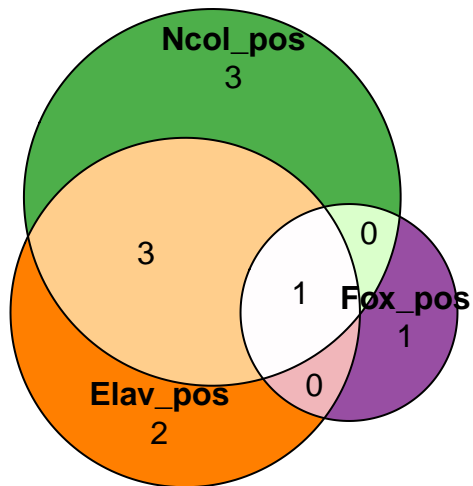
Nvec_vc1.1_XM_001630131.3

ARID_BRIGHT.HG1.2:ARID4A/; ARID4A/ARID4B



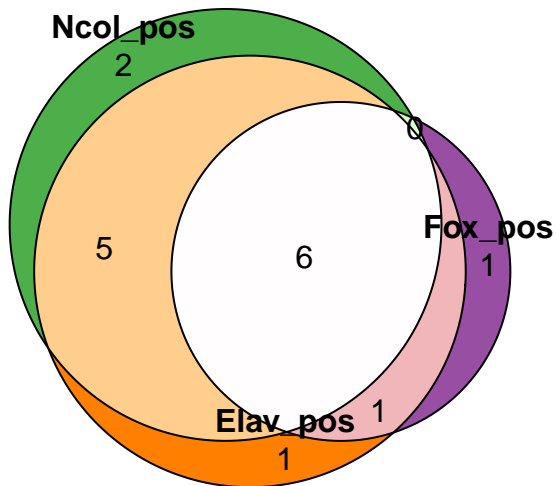
Nvec_NVE758

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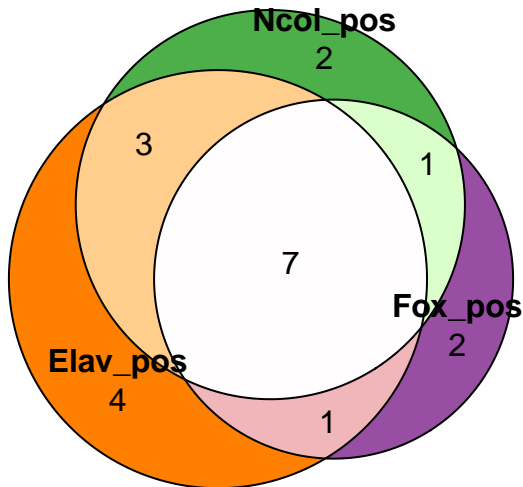
Nvec_vc1.1_XM_032379988.2

Adenosine receptor A2;



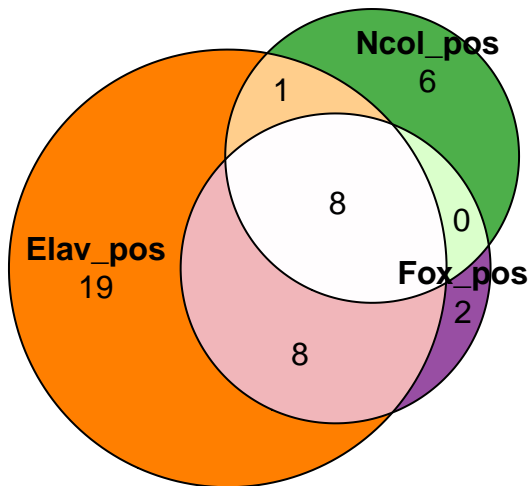
Nvec_vc1.1_XM_032378239.2

Pkinase:ULK1/ULK2; Pkinase.HG107.1:ULK1/ULK2



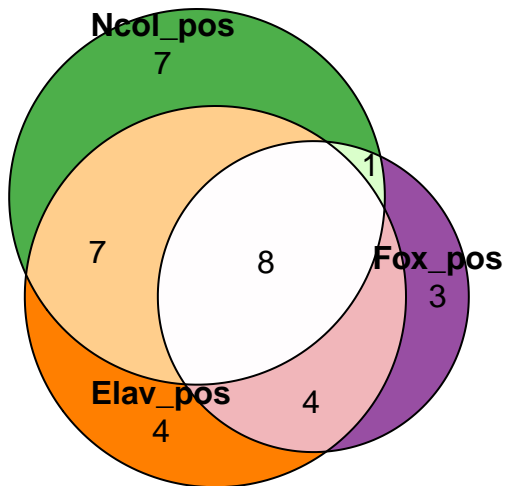
Nvec_vc1.1_XM_032375521.2

Protein phosphatase 1 reg; Ank_2/



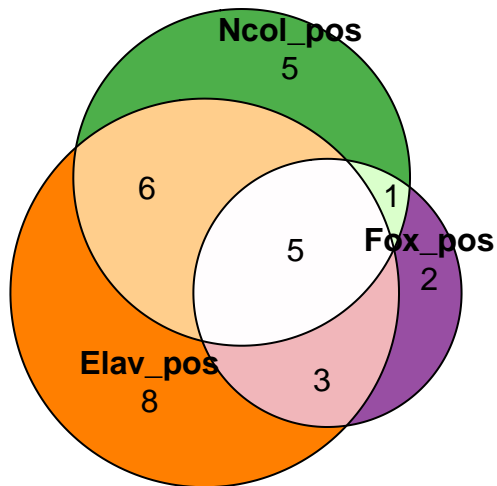
Nvec_NVE11782

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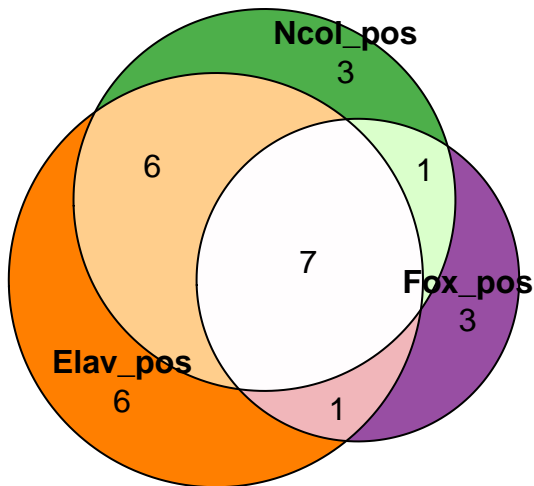
Nvec_vc1.1_XM_001637706.3

Pkinase:MAPK4/MAPK6; Pkinase.HG51.1:MAPK4/MAPK



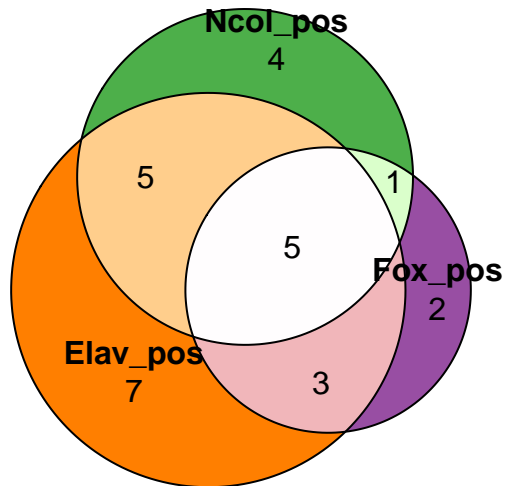
Nvec_vc1.1_XM_001634754.3

PDZ:like:PAR6A/PARD6B/PA; PDZ/



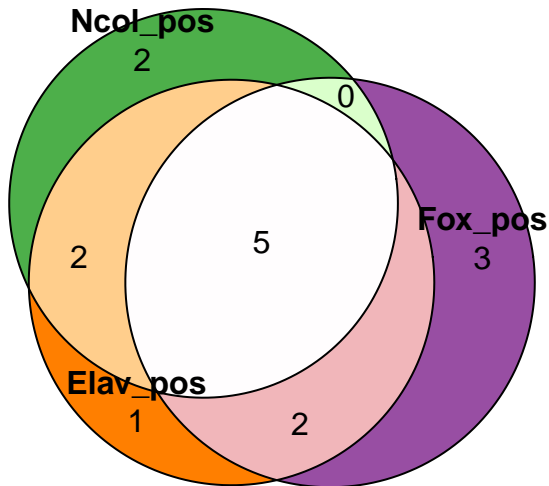
Nvec_vc1.1_XM_001624233.3

Huntingtin interacting pr; TPR_11/Fic/



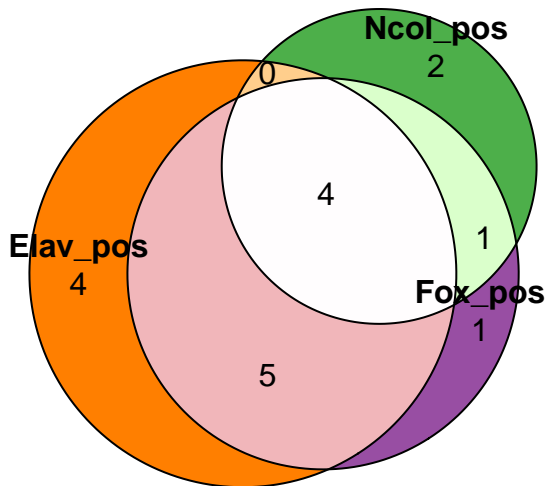
Nvec_vc1.1_XM_001624063.3

Ubiquitin-conjugating enz; UQ_con/



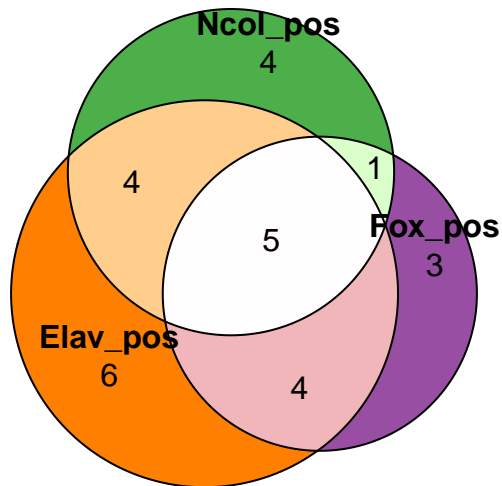
Nvec_vc1.1_XM_001623004.3

Perilipin-2; Perilipin/



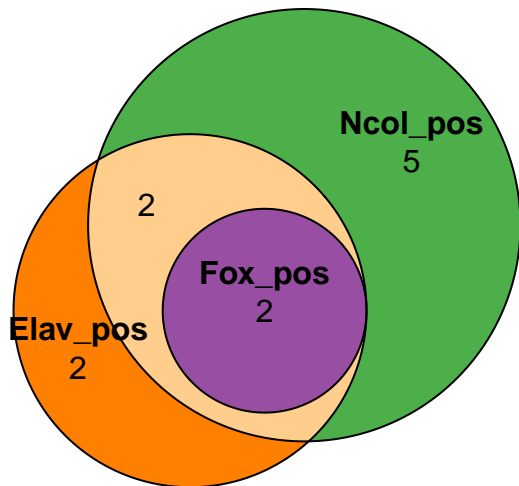
Nvec_vc1.1_XM_032380187.2

ATP-binding cassette sub-; ABC_tran/ABC2_membrane/



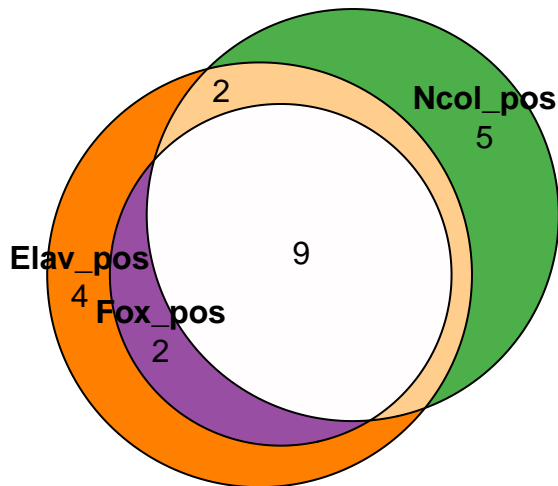
Nvec_vc1.1_XM_032377439.2

PHD finger protein 20-lik; Tudor-knot/



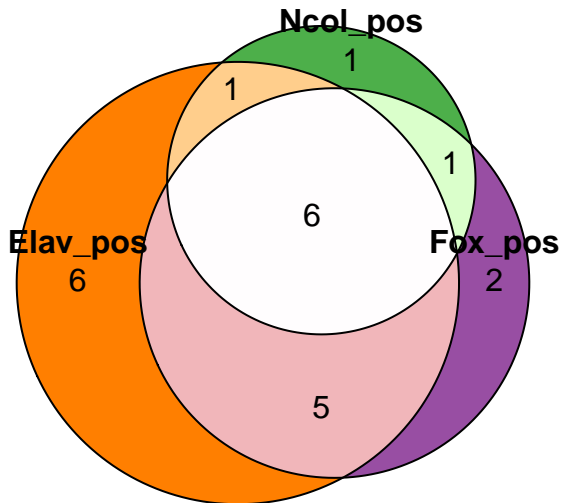
Nvec_vc1.1_XM_032370798.2

CAAX prenyl protease 1 ho; Peptidase_M48/



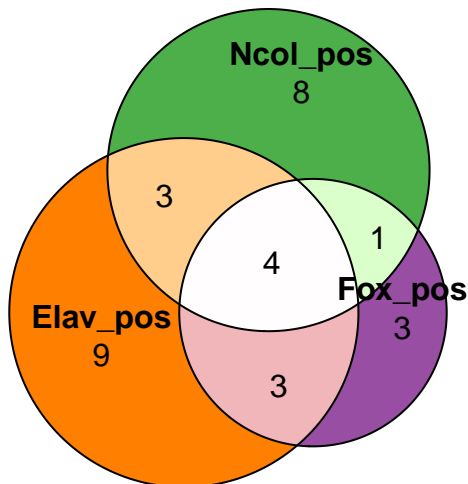
Nvec_vc1.1_XM_032366838.2

cDNA FLJ78634, highly sim; Na_Pi_cotrans/Na_Pi_cotra

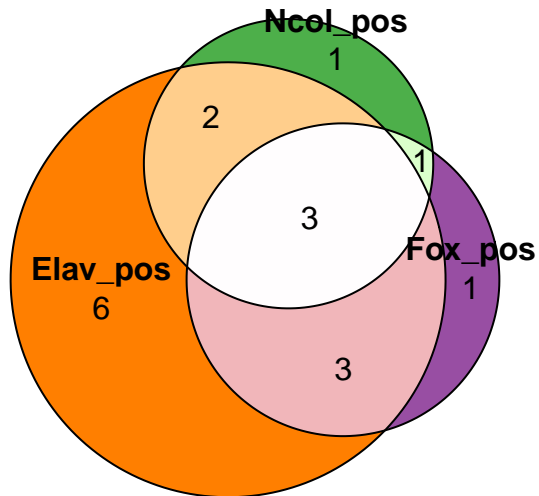


Nvec_vc1.1_XM_001639547.3

zf-C2H2.HG2.11:KLF10/KLF1; KLF10/KLF11

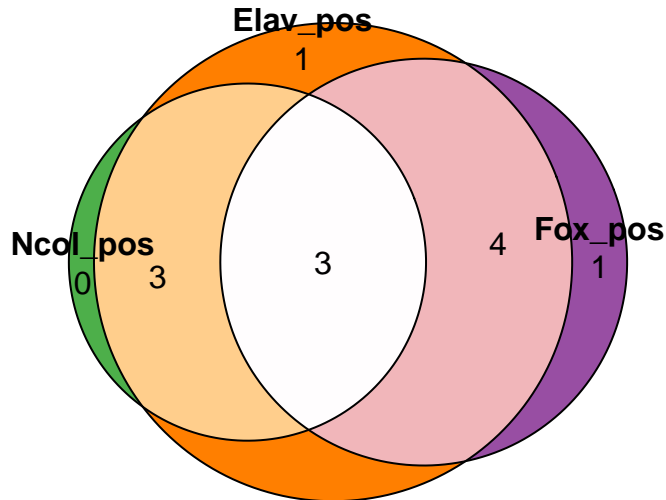


Nvec_vc1.1_XM_001632350.3
Patatin-like phospholipas; Patatin/



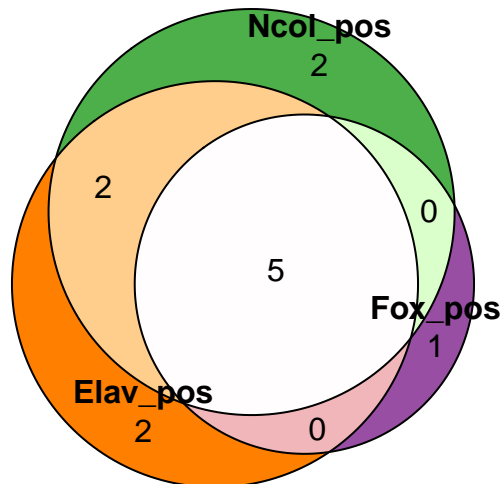
Nvec_vc1.1_XM_001627986.3

Tubulin beta-4B chain; Tubulin/Tubulin_C/



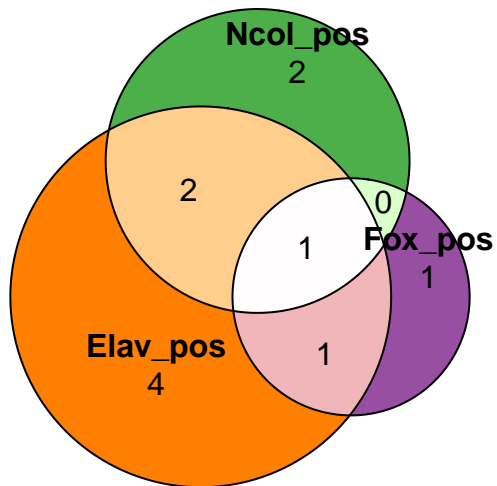
Nvec_vc1.1_XM_001633290.3

Iron/zinc purple acid pho; Metallophos/Metallophos_C



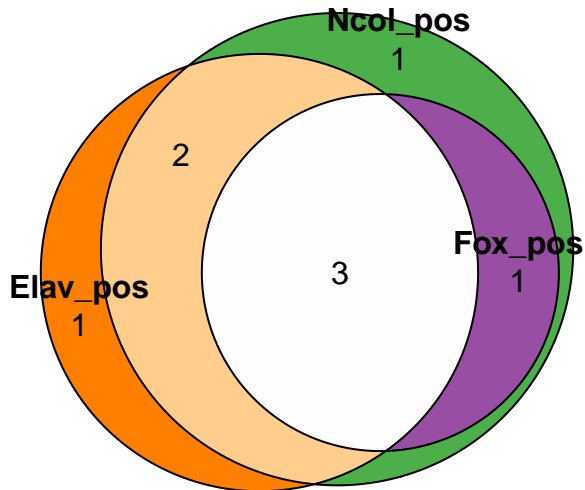
Nvec_NVE3972

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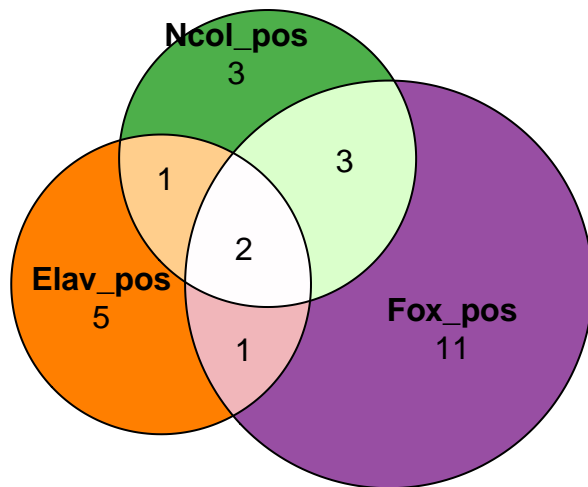


Nvec_vc1.1_XM_032381364.2

Family with sequence simi; FAM53/FAM53/

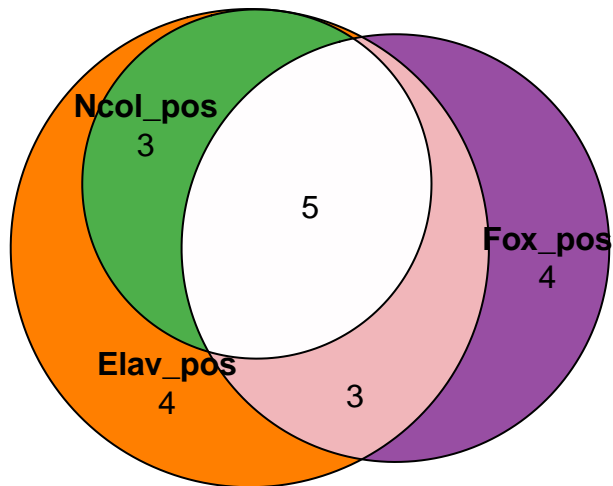


Nvec_vc1.1_XM_048724882.1
cDNA FLJ57120, highly sim; UBX/



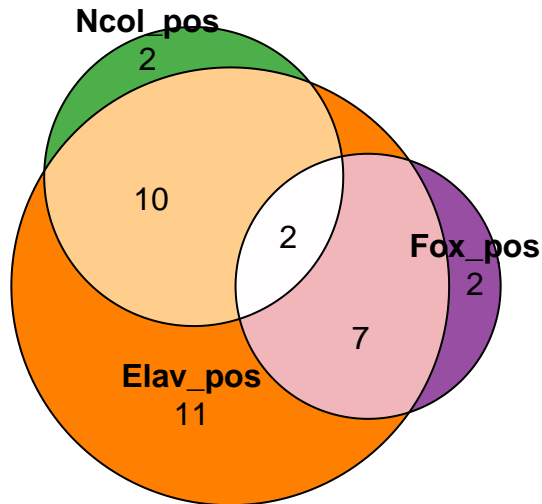
Nvec_vc1.1_XM_001640398.3

Guanine nucleotide-binding; G-gamma/



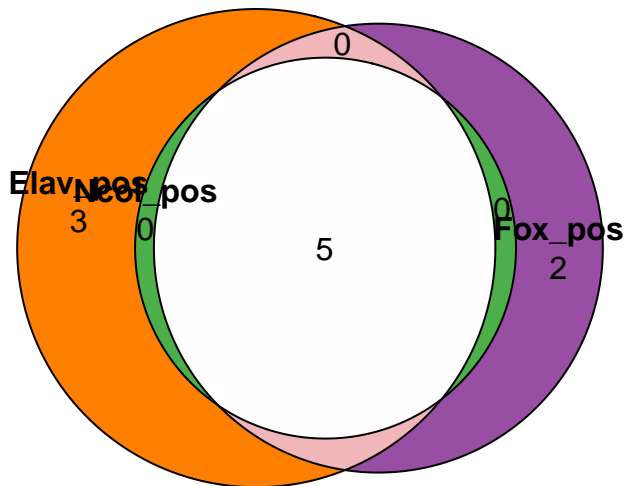
Nvec_vc1.1_XM_001634854.3

Pkinase:TRIB1/TRIB2/TRIB3; Pkinase.HG149.1:TRIB1/TRI



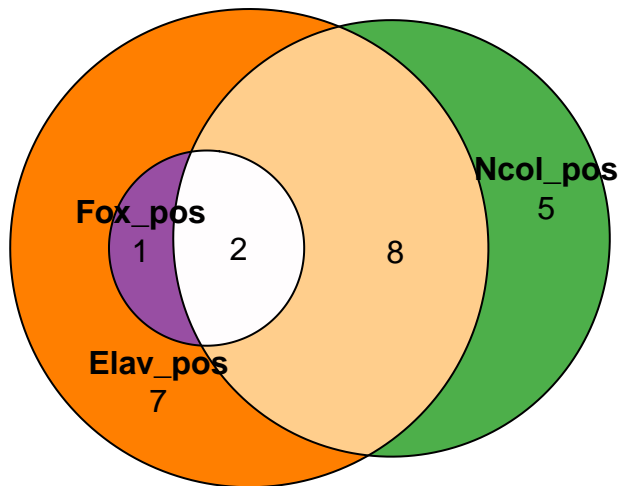
Nvec_vc1.1_XM_048727123.1

Myb.HG1.9:NCOR1/NCOR2; NCOR1/NCOR2



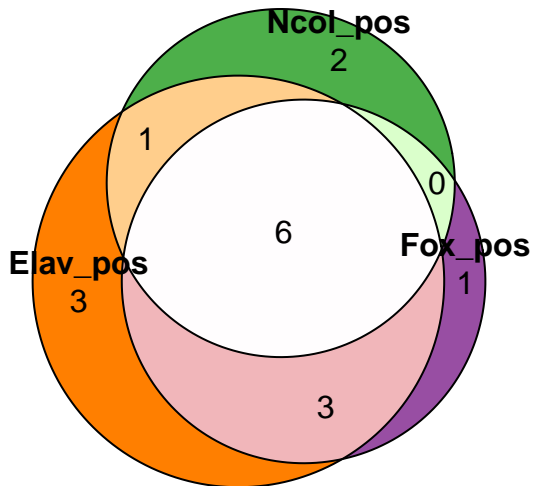
Nvec_vc1.1_XM_048731832.1

Calcium channel, voltage-; Ion_trans/



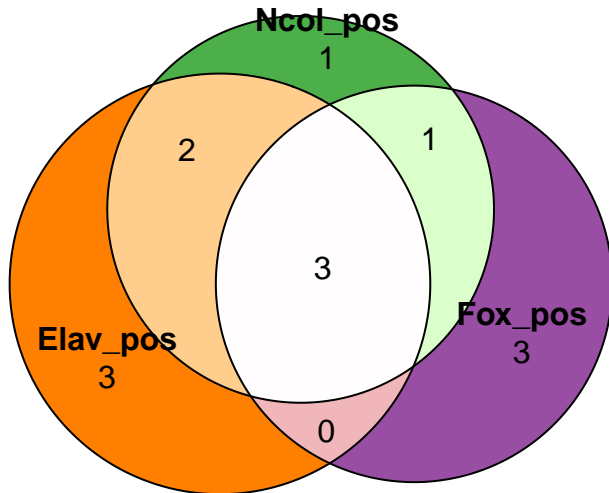
Nvec_vc1.1_XM_032377385.2

Cathepsin Z; Peptidase_C1/



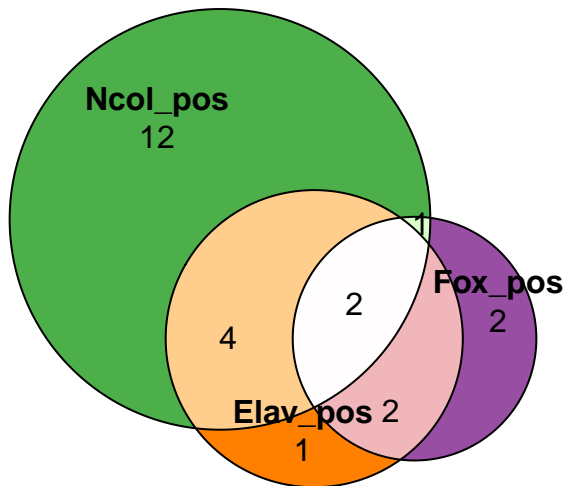
Nvec_vc1.1_XM_032366722.2

Ldl_recept_b:LRP5/LRP5L/L; Ldl_recept_b.HG1.16:LRP5/



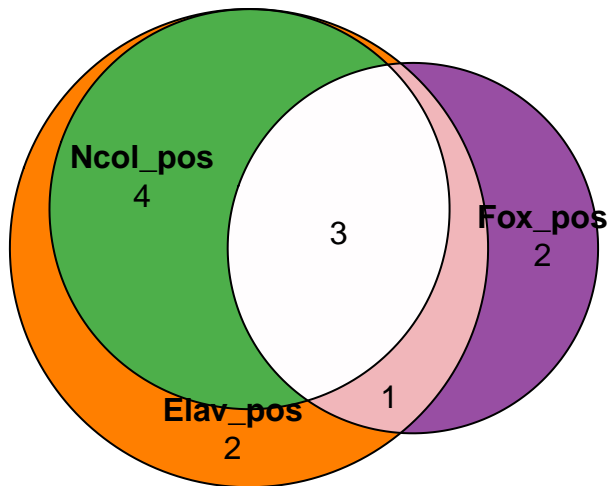
Nvec_vc1.1_XM_032382923.2

E3 ubiquitin–protein liga; Filamin/NHL/NHL/NHL/NHL/N

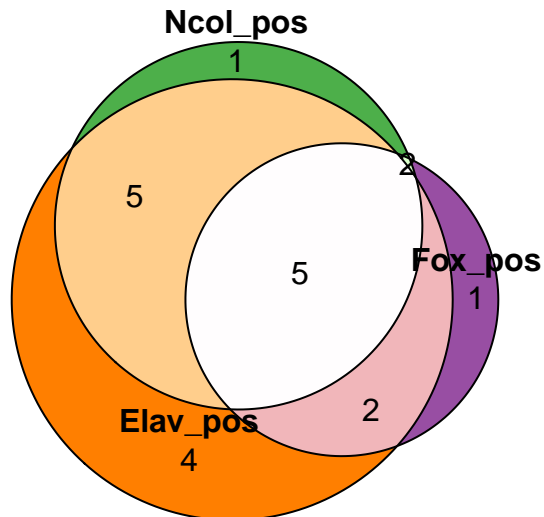


Nvec_vc1.1_XM_048723054.1

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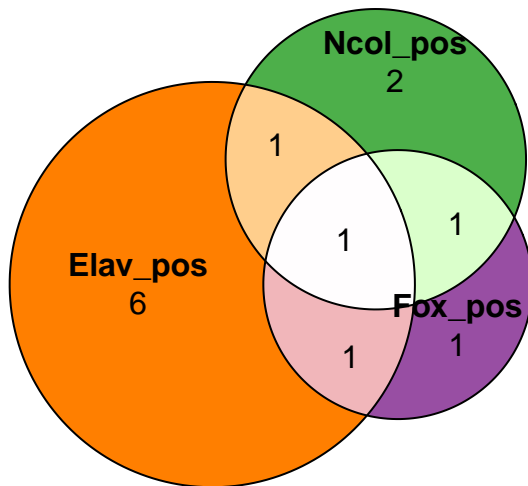


Nvec_vc1.1_XM_048733736.1
RHD_Rel.HG1.0:like:NFAT5/;

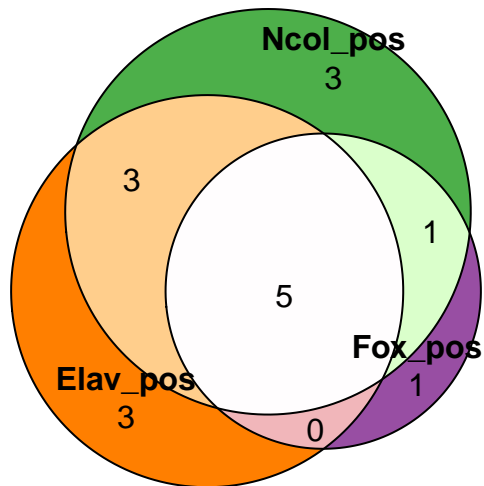


Nvec_vc1.1_XM_001638150.3

Pkinase:LIMK1/LIMK2/TESK1; Pkinase.HG45.1:LIMK1/LIMK

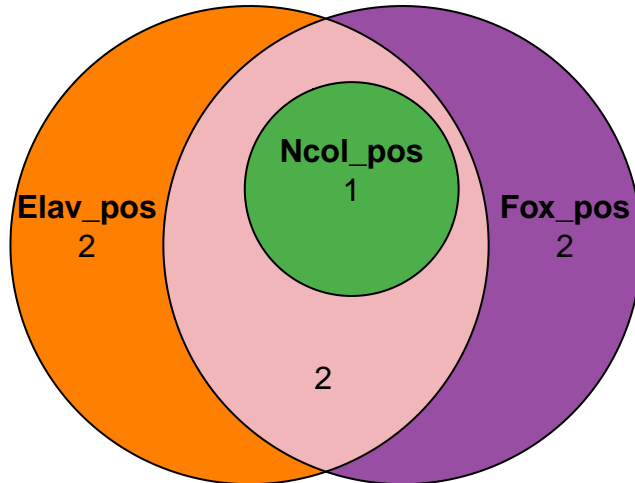


Nvec_vc1.1_XM_032372114.2
bZIP.HG1.8:ATF4/ATF5; ATF4/ATF5

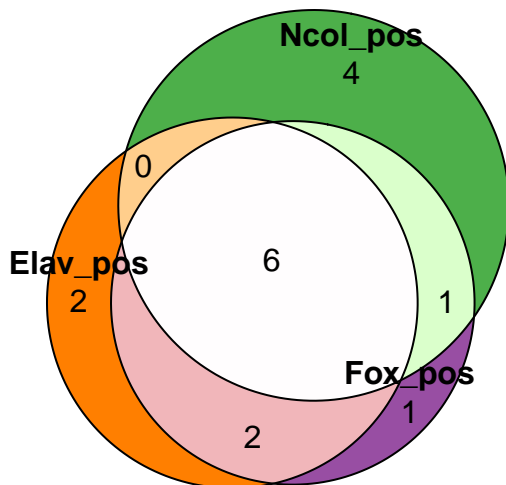


Nvec_vc1.1_XM_001626362.3

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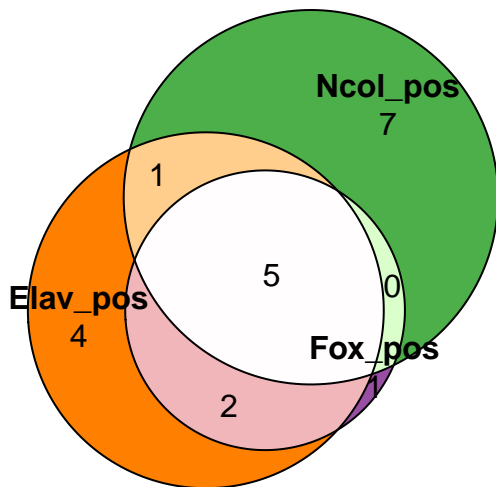


Nvec_vc1.1_XM_001633592.3
Uncharacterized protein; ubiquitin/



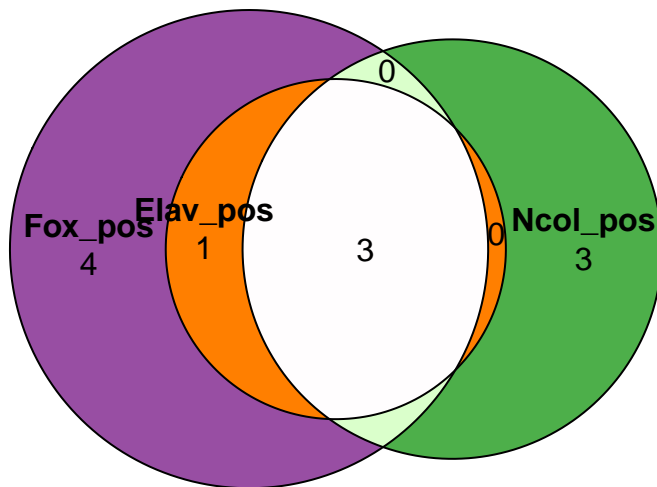
Nvec_NVE25571

Ferritin;

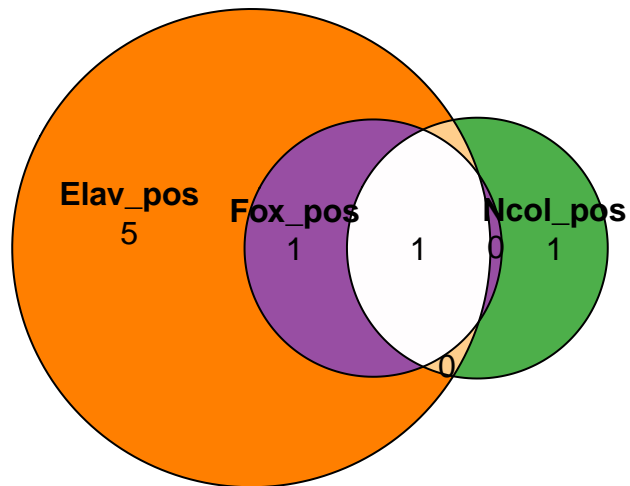


Nvec_vc1.1_XM_048726662.1

RID_BRIGHT.HG1.1:KDM5A/K; KDM5A/KDM5B/KDM5C/KDM5

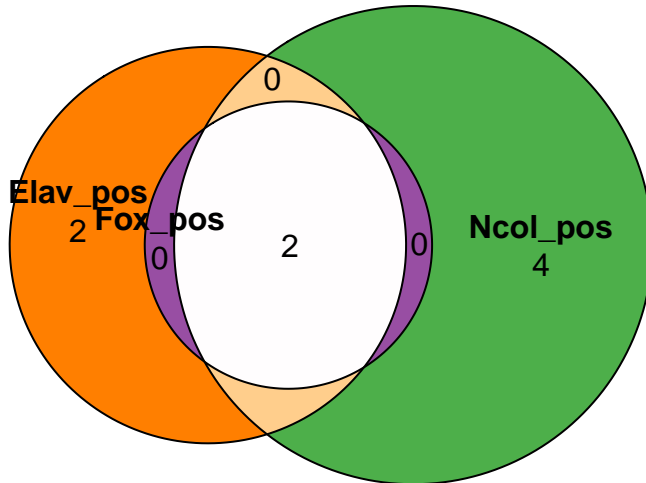


Nvec_vc1.1_XM_048723371.1
DEAD:DDX4; DEAD.HG4.6:DDX4



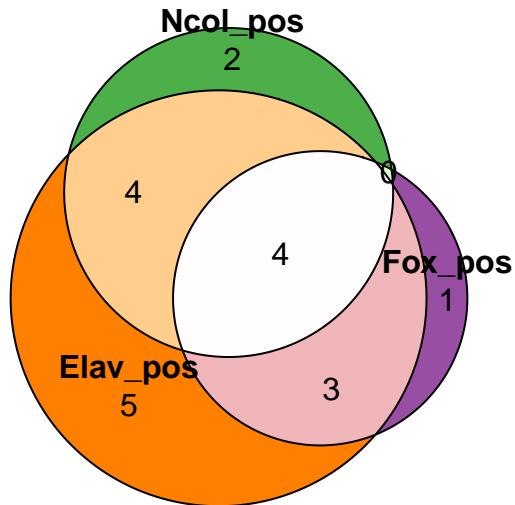
Nvec_vc1.1_XM_001641142.3

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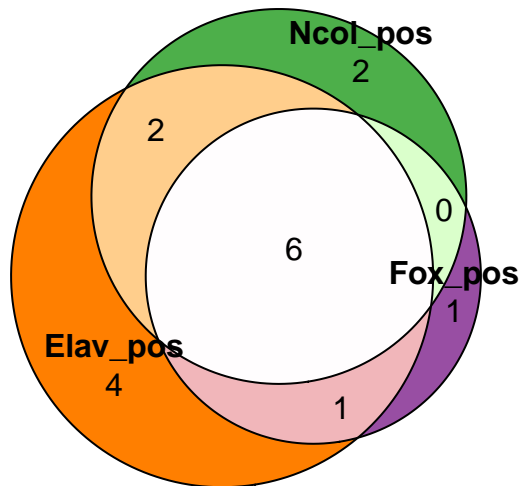
Nvec_vc1.1_XM_001623716.3

Histone H3; Histone/



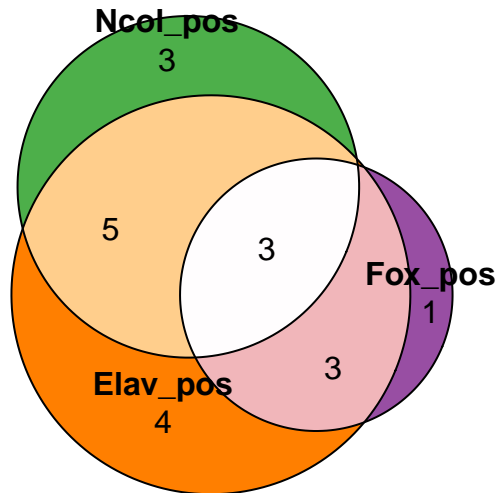
Nvec_vc1.1_XM_032380616.2

Palmitoyl-protein thioest; Palm_thioest/

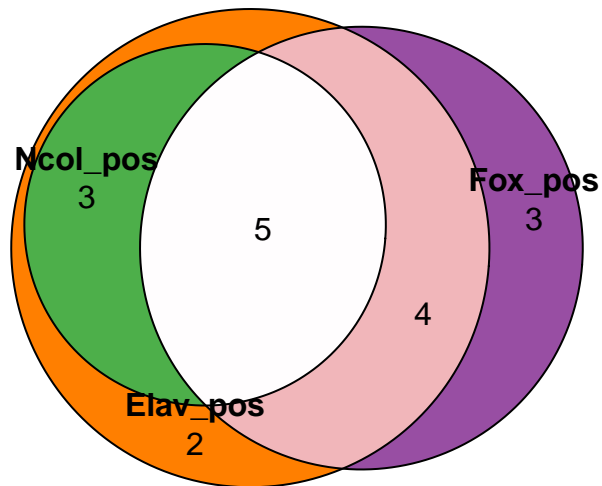


Nvec_vc1.1_XM_032374373.2

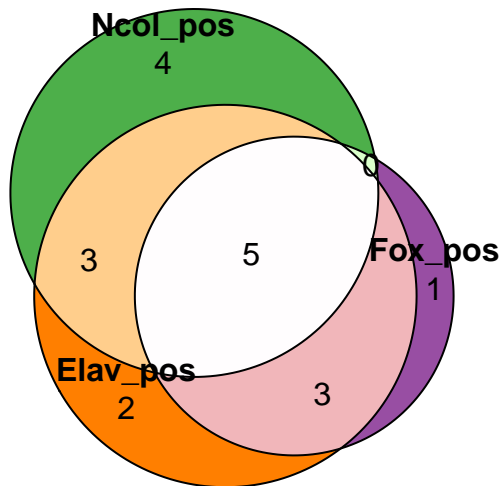
Rab geranylgeranyltransfe; Prenyltrans/



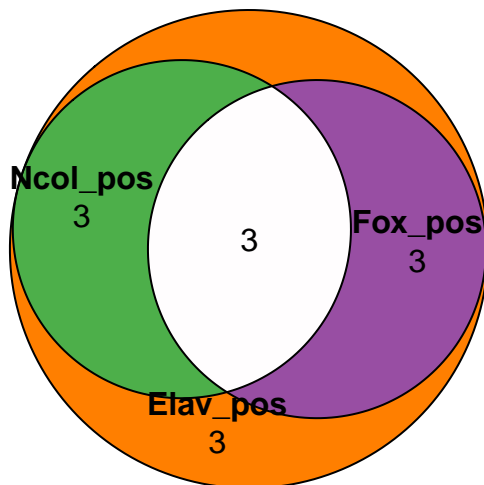
Nvec_vc1.1_XM_001626699.3
EVI5-like protein; RabGAP-TBC/



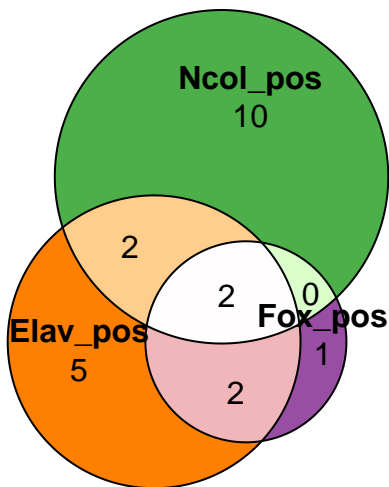
Nvec_vc1.1_XM_048721608.1
cDNA FLJ52165, highly sim; UQ_con/



Nvec_vc1.1_XM_032385038.2
Death-associated protein ; Daxx/

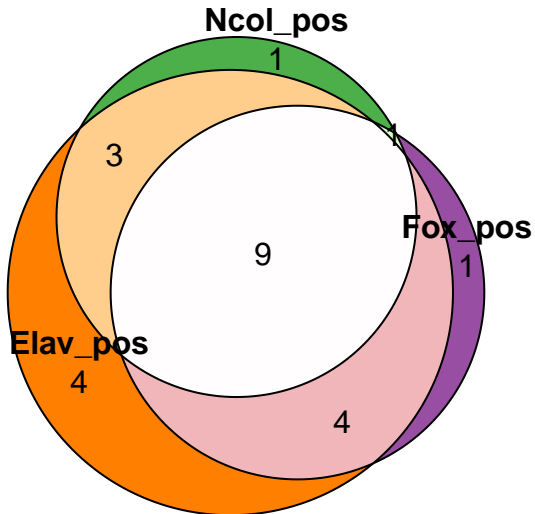


Nvec_vc1.1_XM_001629728.3
Protein phosphatase 1 reg; CBM_21/



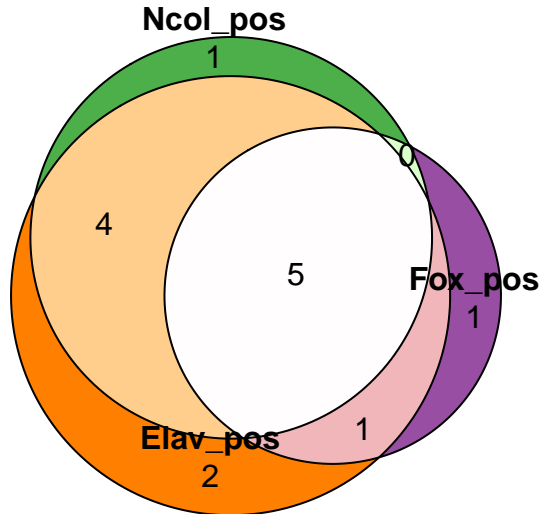
Nvec_vc1.1_XM_001629618.3

; Usp/



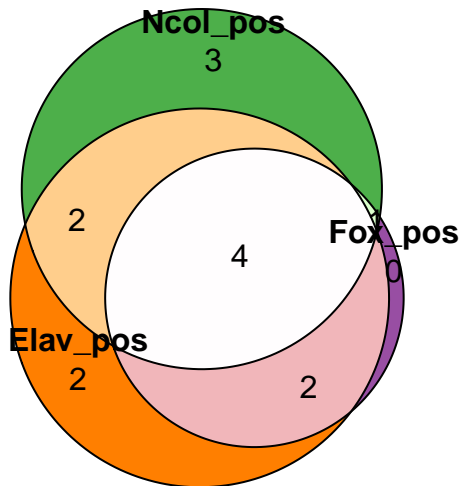
Nvec_vc1.1_XM_032373806.2

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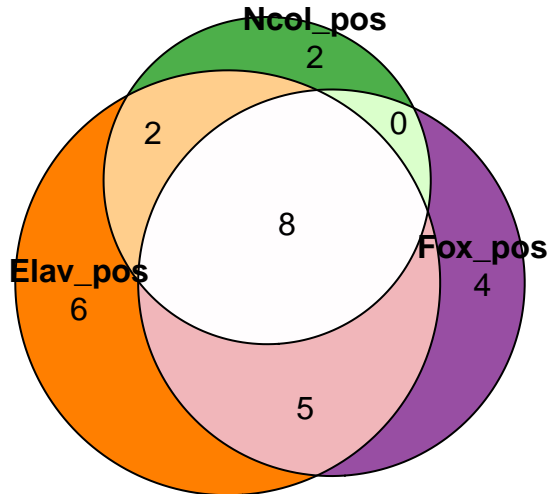


Nvec_vc1.1_XM_001639991.3

Beta tropomyosin isoform; Tropomyosin/

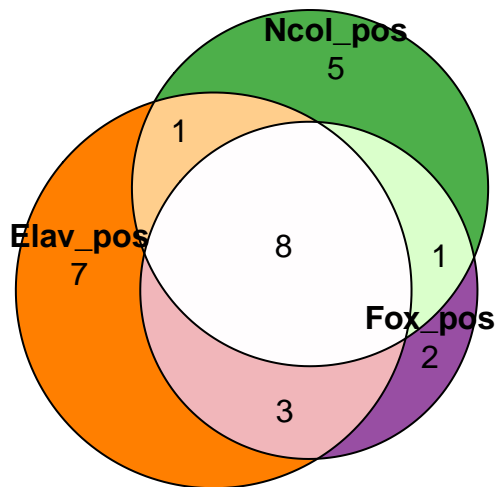


Nvec_vc1.1_XM_048723161.1
cDNA FLJ50585, highly sim;



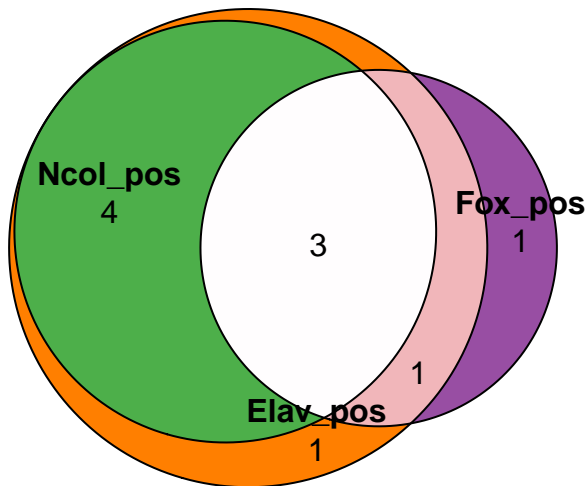
Nvec_vc1.1_XM_032364382.2

chromo:CHD6/CHD7/CHD8/CHD; Chromo.HG4.0:CHD6/CHD7/CHD8/CHD9

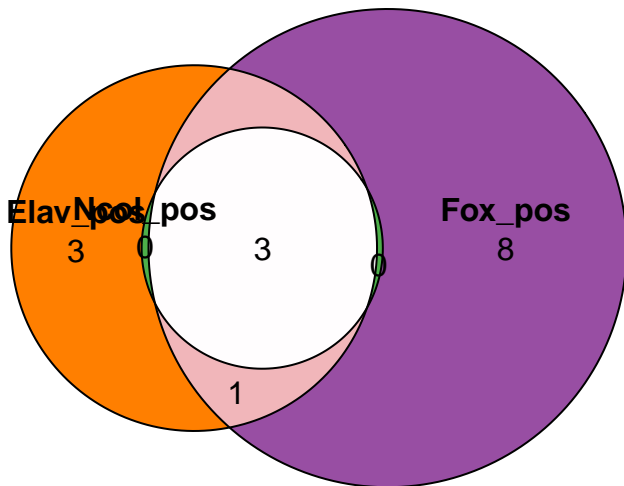


Nvec_vc1.1_XM_032380170.2

F-box only protein 22; F-box-like/FIST/

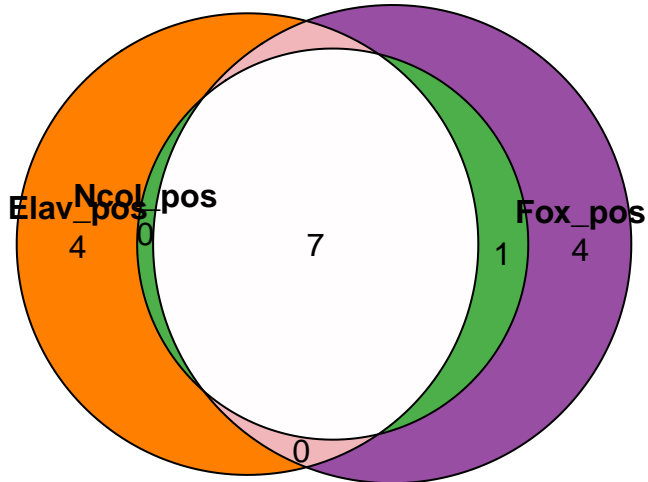


Nvec_vc1.1_XM_001627423.3
HMGbox_Sox.HG1.10:BBX; BBX



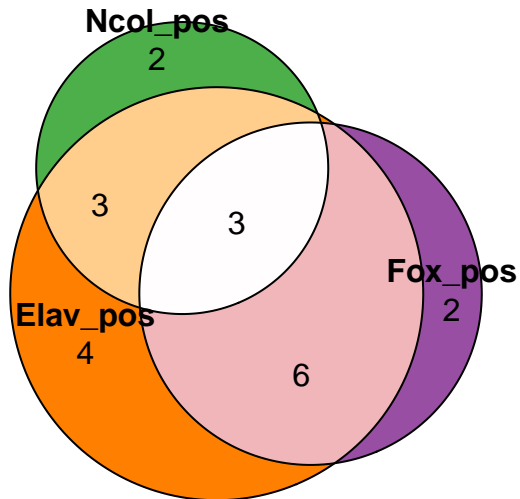
Nvec_vc1.1_XM_032377982.2

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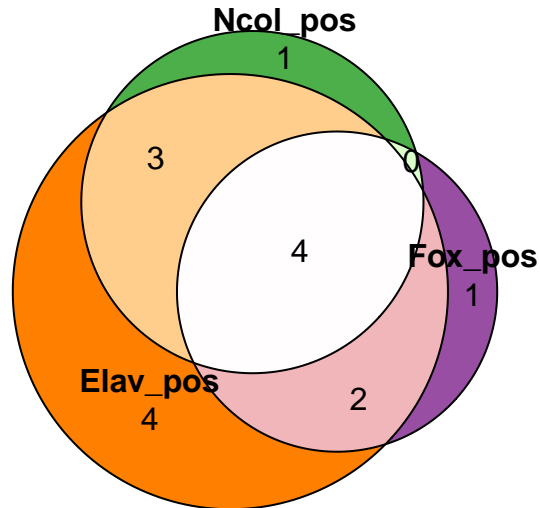
Nvec_vc1.1_XM_032372722.2

RRM:PABPC1/PABPC1L/PABPC1; RRM.HG4.13:PABPC1/PABPC1



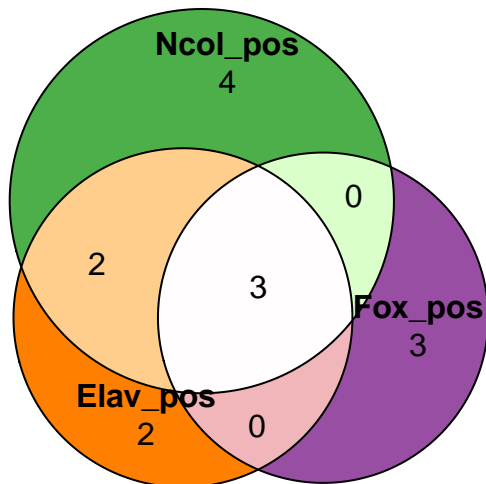
Nvec_vc1.1_XM_032384654.2

Transmembrane anterior po; DUF747/



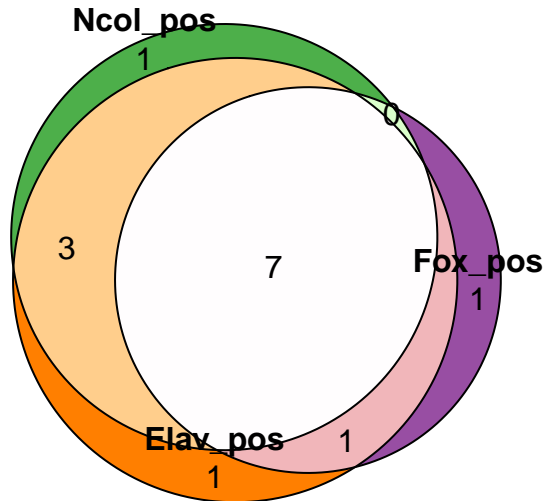
Nvec_vc1.1_XM_001641577.3

RelA-associated inhibitor; Ank_2/SH3_1/



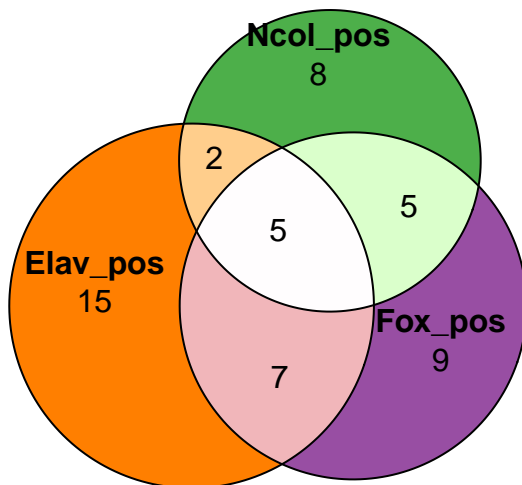
Nvec_vc1.1_XM_032375203.2

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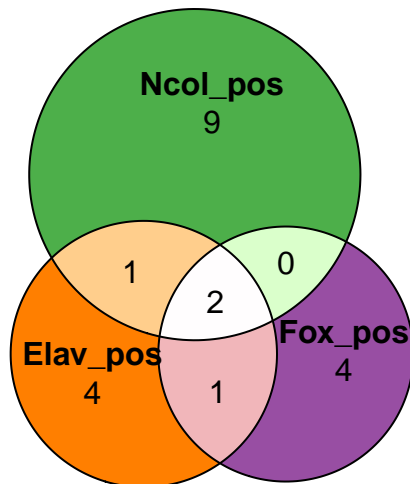
Nvec_vc1.1_XM_001633854.3

GPCR-glutamate:like:GABBR; 7tm_3/



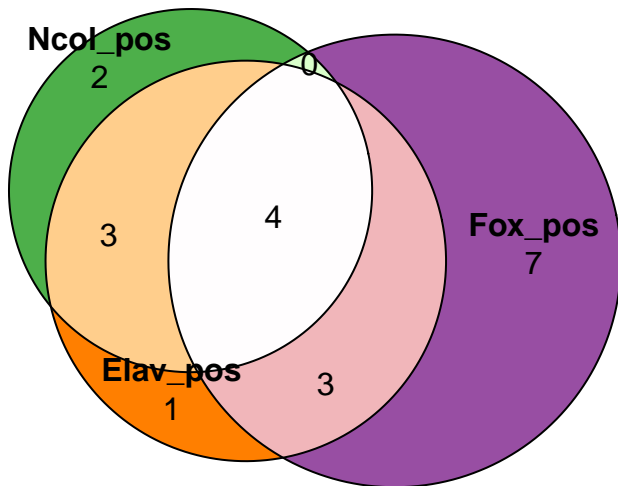
Nvec_vc1.1_XM_048729934.1

BTB:like:CCIN/GAN/KLHL11;; BTB/BACK/Kelch_1/Kelch_1/



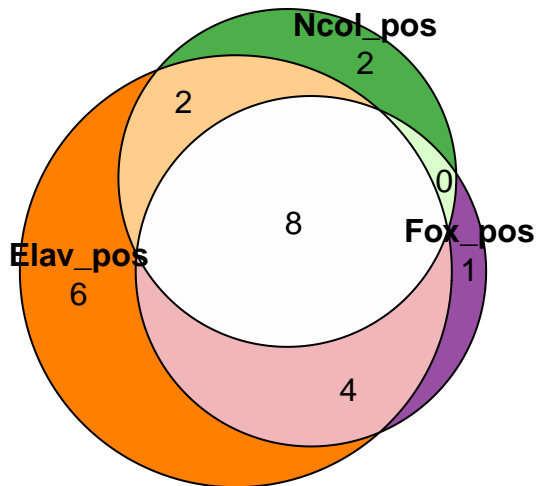
Nvec_vc1.1_XM_001622410.3

;



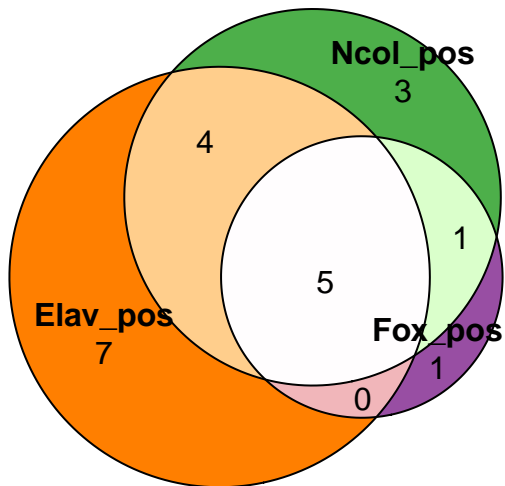
Nvec_vc1.1_XM_001634922.3

Kelch domain-containing p; Kelch_6/Kelch_4/Kelch_1/K



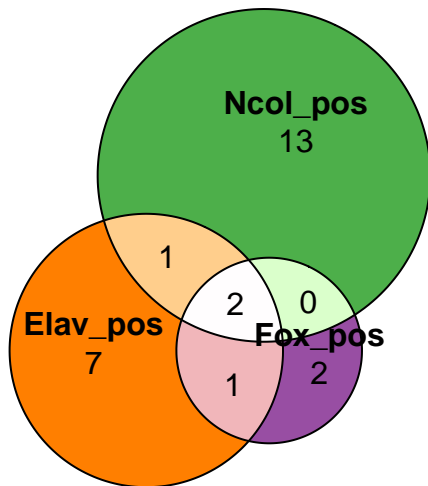
Nvec_vc1.1_XM_001621919.3

Cysteine/serine-rich nucl;



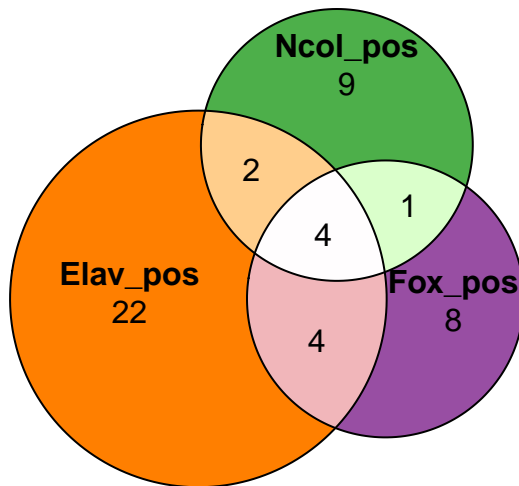
Nvec_vc1.1_XM_032378865.2

;



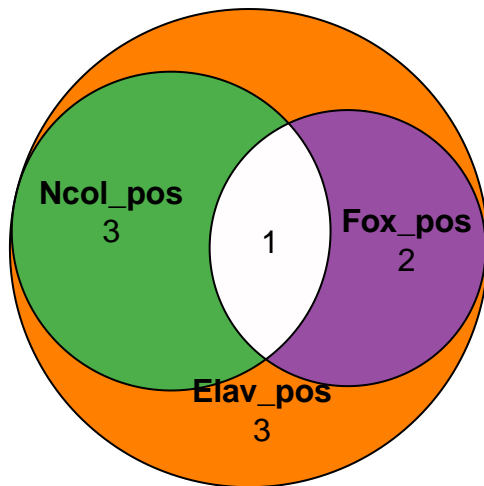
Nvec_vc1.1_XM_032377587.2

cDNA FLJ41053 fis, clone ; F-box/LRR_6/LRR_6/

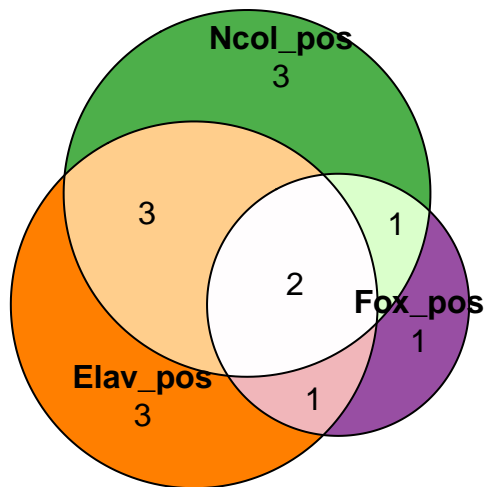


Nvec_vc1.1_XM_048724894.1

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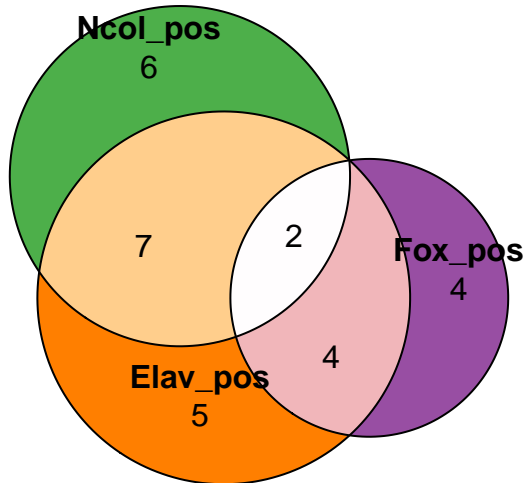


Nvec_vc1.1_XM_032386689.2
Aryl hydrocarbon receptor; PAS/

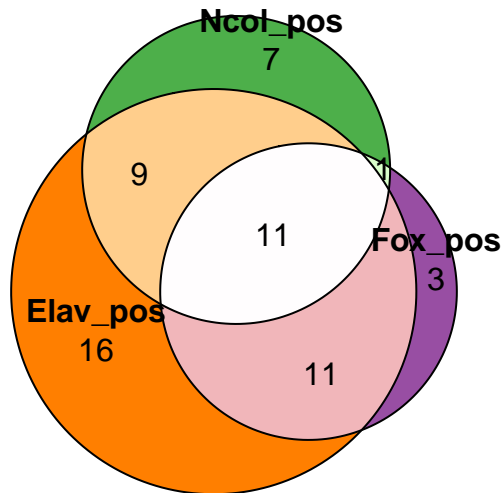


Nvec_vc1.1_XM_032379825.2

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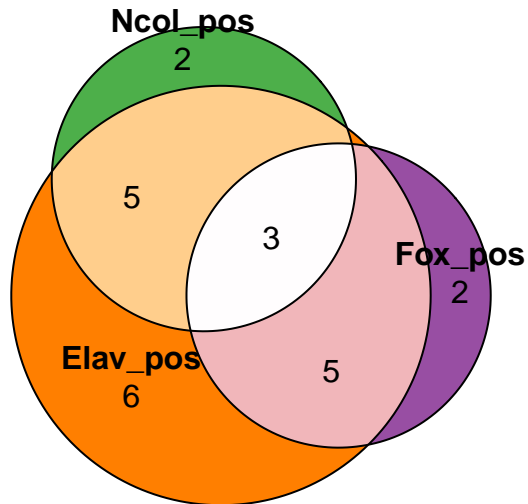


Nvec_vc1.1_XM_001638761.3
zf-C2H2.HG16.2:like:TRERF;



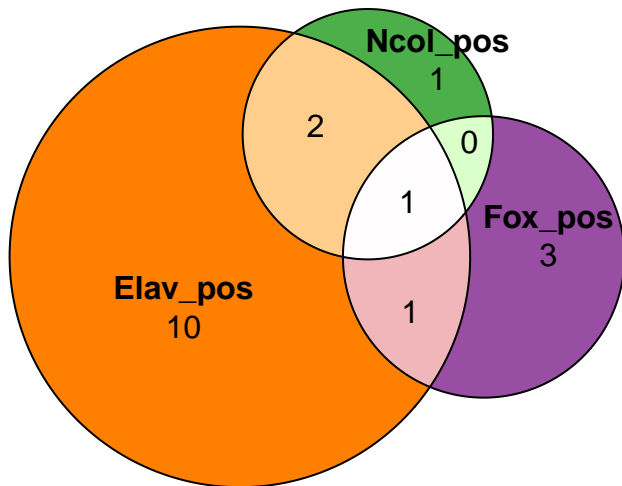
Nvec_vc1.1_XM_001639531.3

Protein yippee-like 1; Yippee-Mis18/



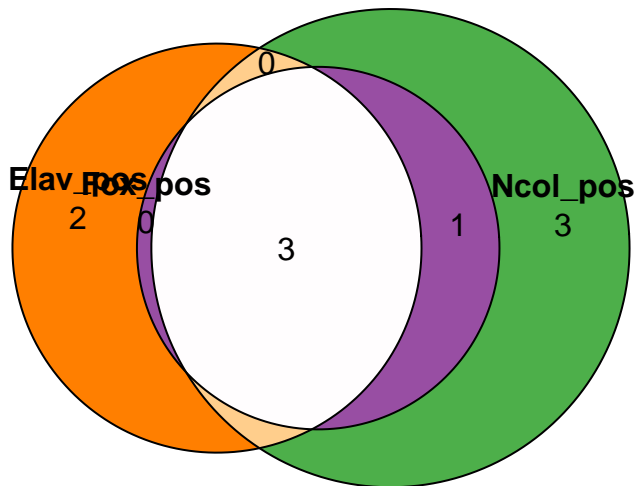
Nvec_v1g241678

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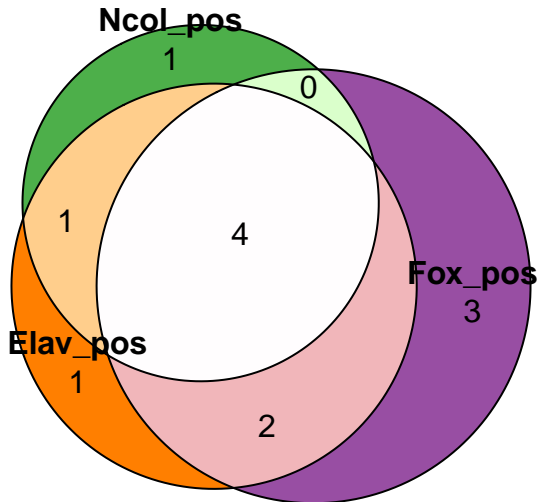


Nvec_vc1.1_XM_001641695.3

SAM:STIM1/STIM2; SAM.HG11.0:STIM1/STIM2

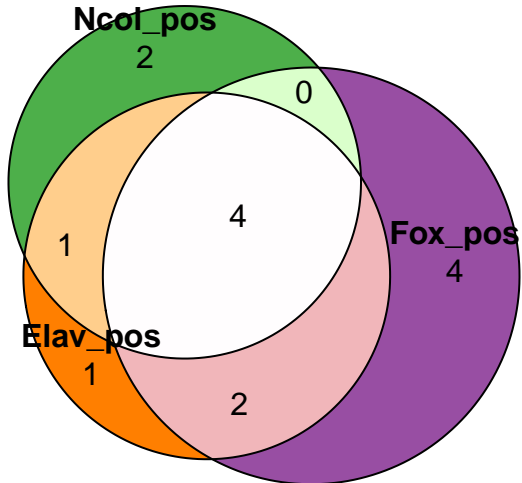


Nvec_vc1.1_XM_048727141.1
TBCC domain-containing pr; TBCC/

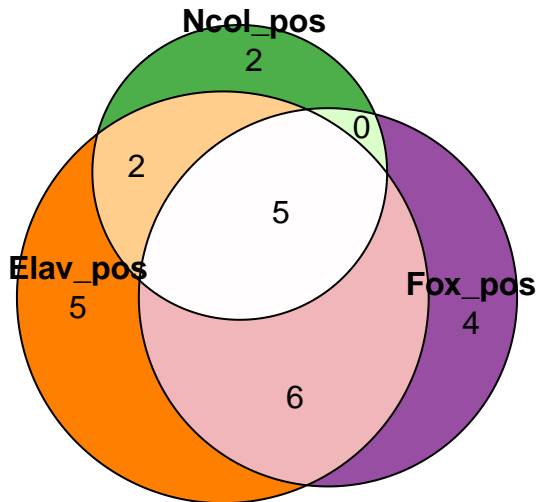


Nvec_vc1.1_XM_048722504.1

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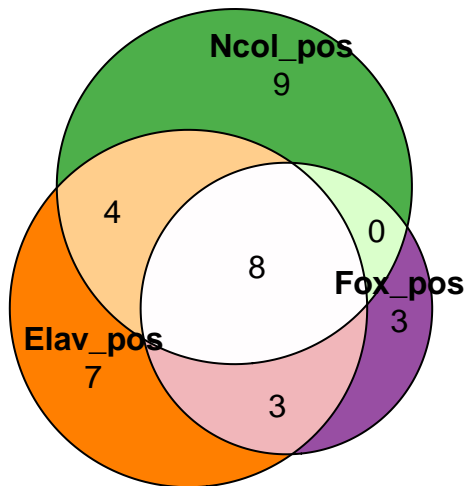


Nvec_vc1.1_XM_001626603.3
Golgi-associated plant pa; CAP/



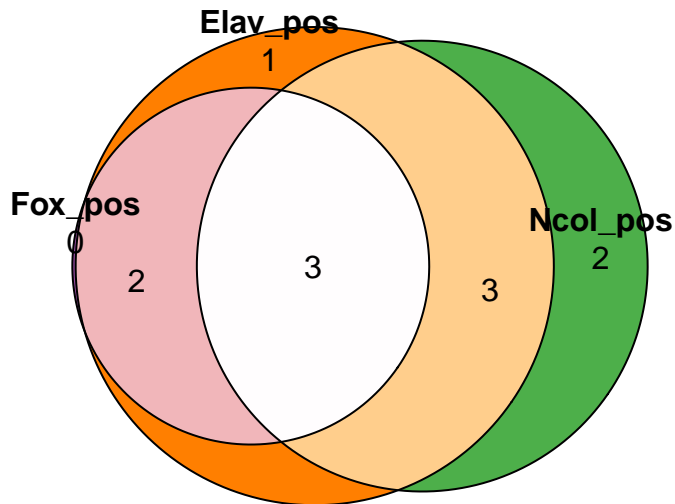
Nvec_vc1.1_XM_001639831.3

Kinesin-like protein; Kinesin/



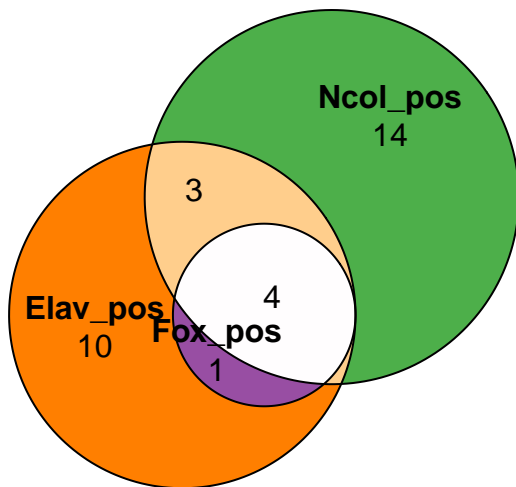
Nvec_vc1.1_XM_048722272.1

Malic enzyme; malic/Malic_M/



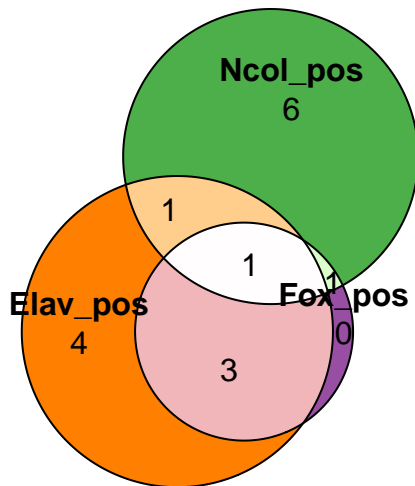
Nvec_vc1.1_XM_032362411.2

n-NaKCa:KCNH3/KCNH4/KCN; ion-NaKCa.HG10.2:KCNH3/K



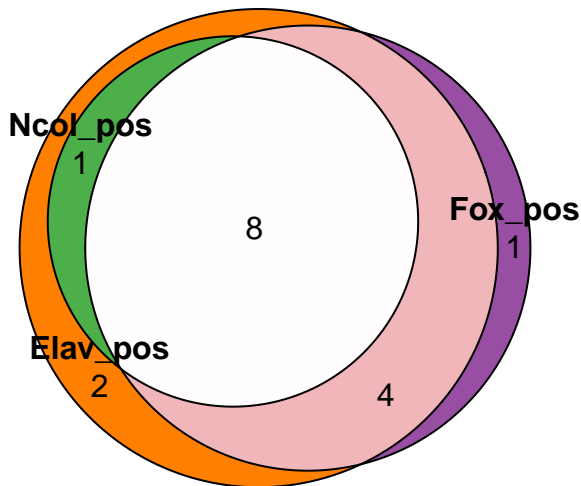
Nvec_vc1.1_XM_001626441.3

ERM_M:EPB41L4A/EPB41L4B/; FERM_M.HG1.9:EPB41L4A/EPB41L4B/

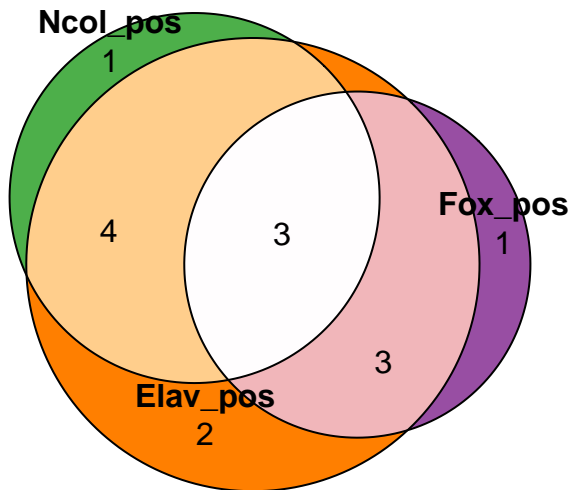


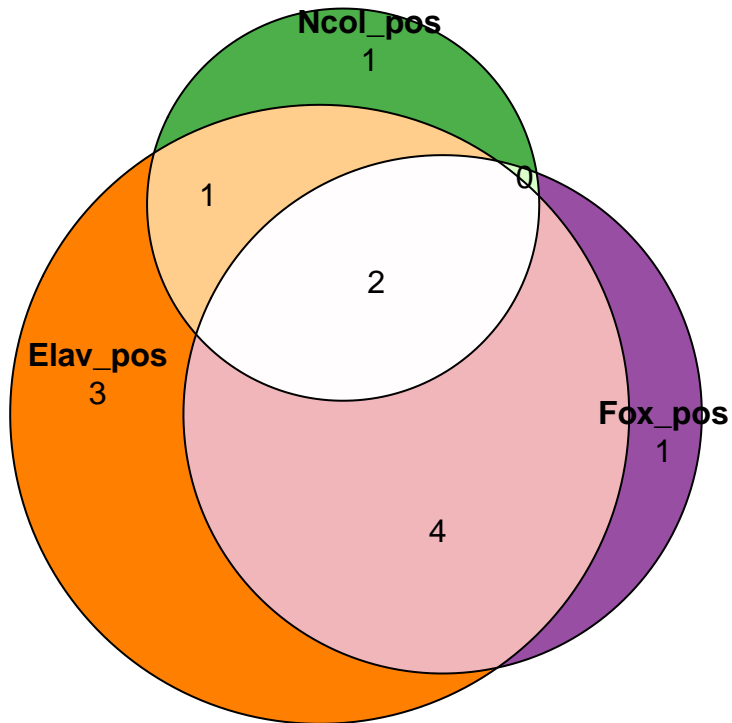
Nvec_vc1.1_XM_001631883.3

Beta-galactosidase; Glyco_hydro_35/



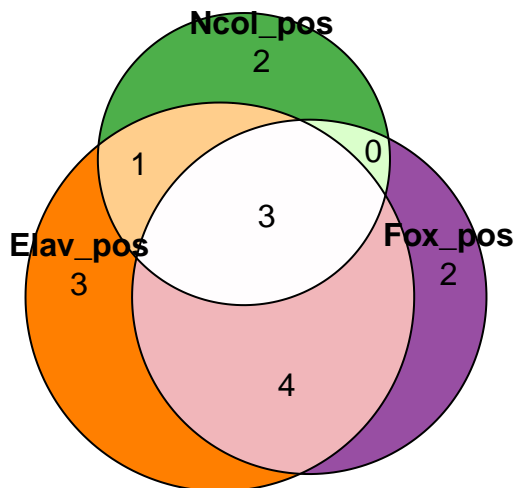
Nvec_vc1.1_XM_032382705.2
cDNA FLJ90319 fis, clone ;



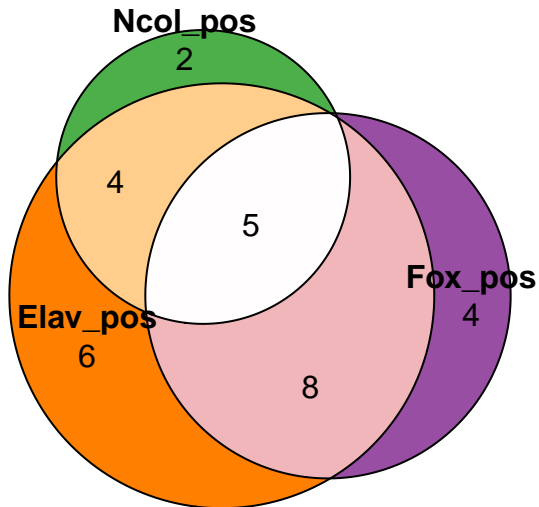


Nvec_vc1.1_XM_032380497.2

PURA protein; PurA/

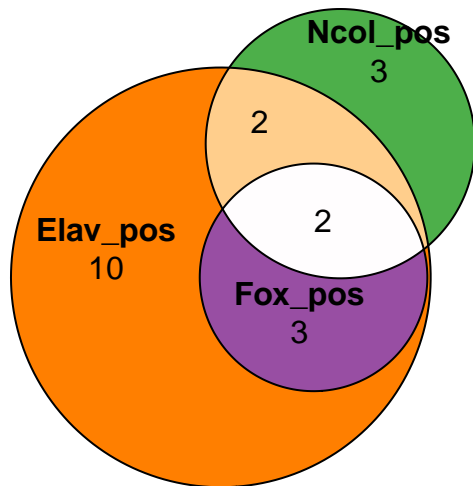


Nvec_vc1.1_XM_032381269.2
; DUF4419/



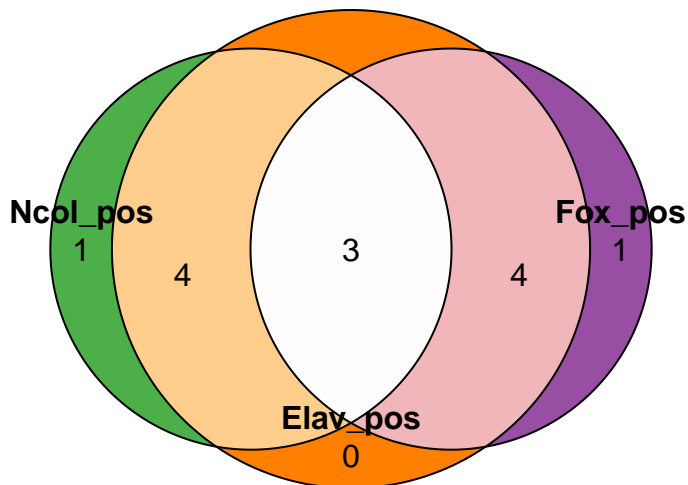
Nvec_vc1.1_XM_001640639.3

Epididymis luminal secret; HATPase_c_3/HSP90/



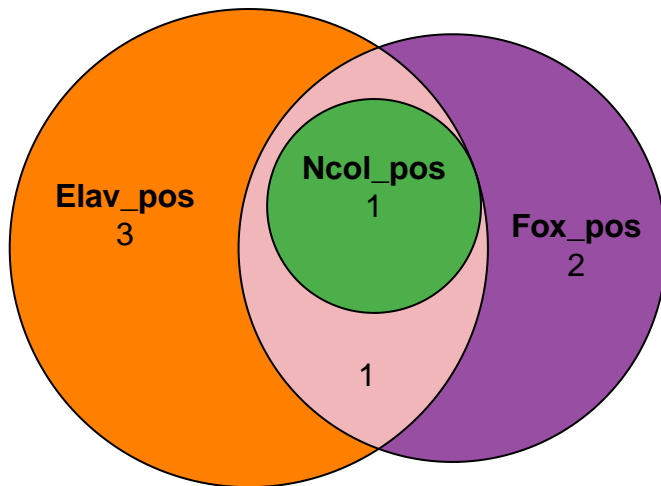
Nvec_vc1.1_XM_032363989.2

on-NaKCa:PKD2/PKD2L1/PKD; ion-NaKCa.HG2.2:PKD2/PKD

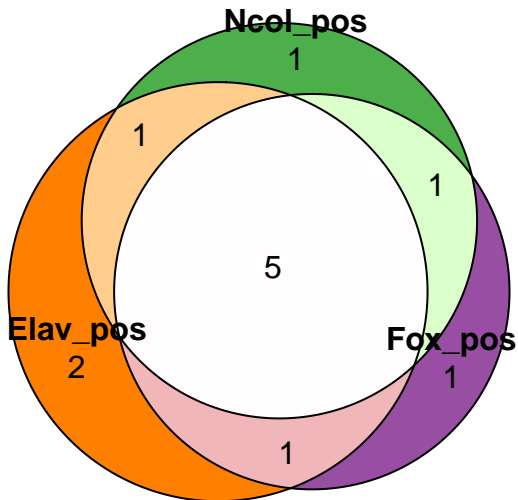


Nvec_vc1.1_XM_001623108.3

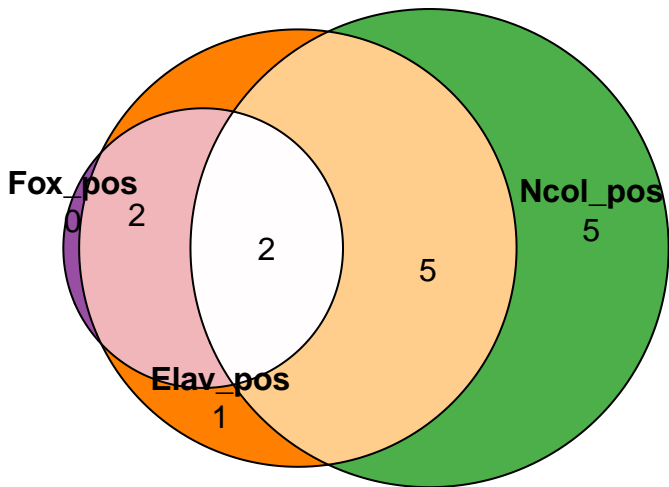
Graves disease carrier pr; Mito_carr/Mito_carr/Mito_



Nvec_vc1.1_XM_032374436.2
Sphingosine-1-phosphate p; PAP2/

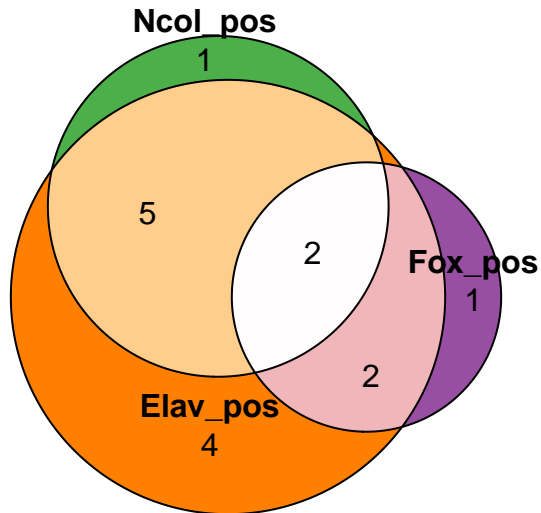


Nvec_vc1.1_XM_001637926.3
Gamma-aminobutyric acid r; Atg8/



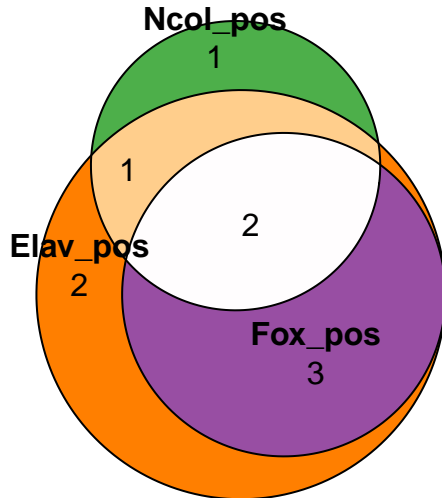
Nvec_vc1.1_XM_001626637.3

Sorting nexin-11; PX/



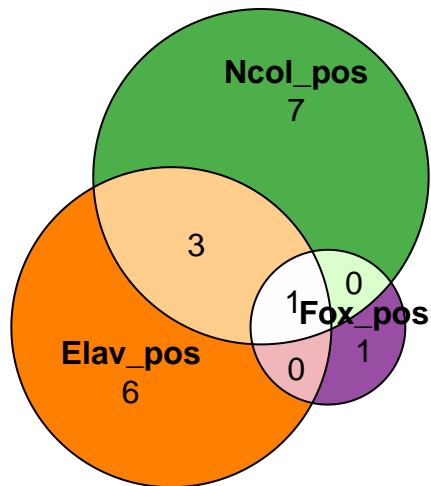
Nvec_vc1.1_XM_001629636.3

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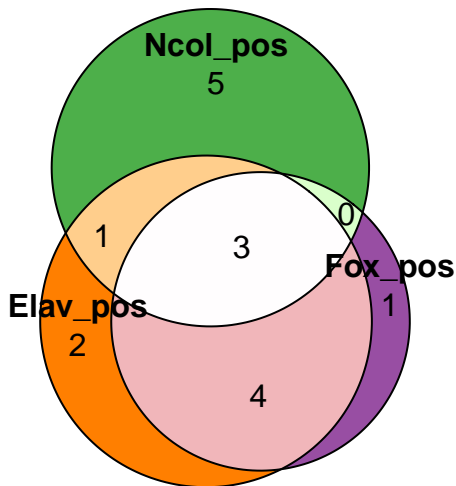


Nvec_vc1.1_XM_032385582.2

PDZ domain containing rin; zf-C3HC4/

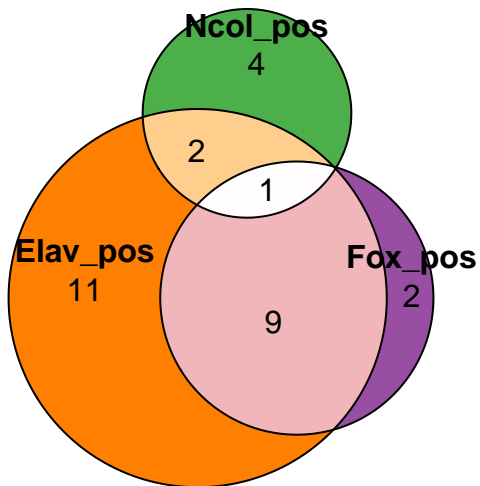


Nvec_vc1.1_XM_001639183.3
BTB:KLHL20; BTB.HG1.36:KLHL20



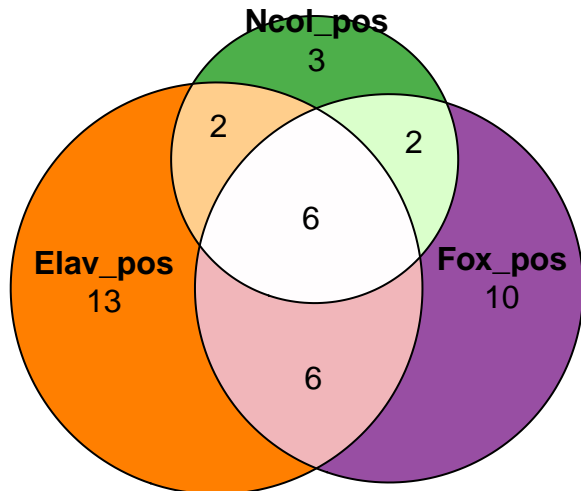
Nvec_v1g248496

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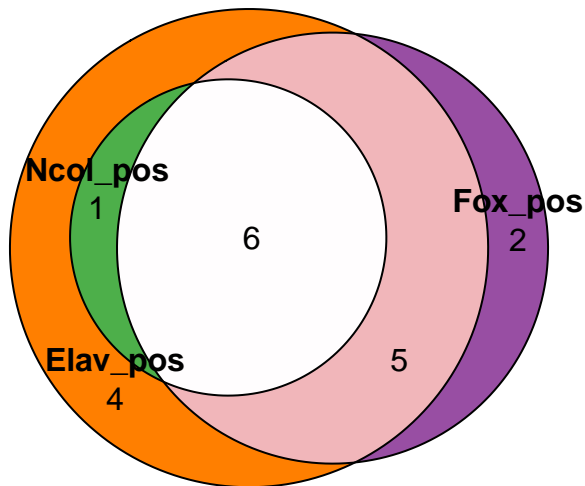
Nvec_vc1.1_XM_048727546.1

Pkinase:like:KSR1/KSR2:li; Pkinase_Tyr/



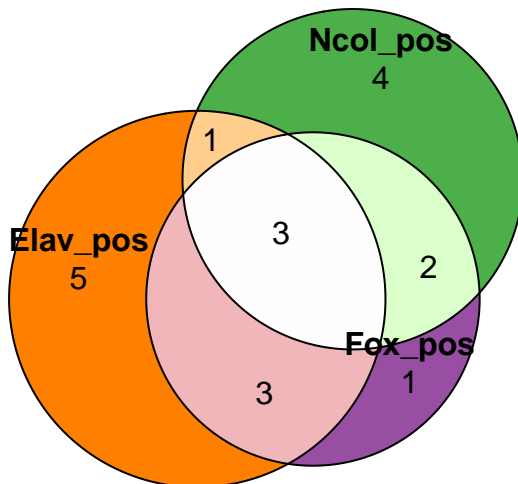
Nvec_vc1.1_XM_001636508.3

FA_hydroxylase:FAXDC2; FA_hydroxylase.HG1.12:FAX



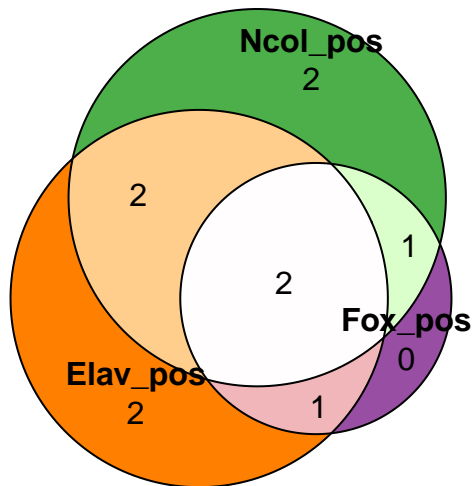
Nvec_vc1.1_XM_001630376.3

Dinucleotide oxidase disu; EMP70/



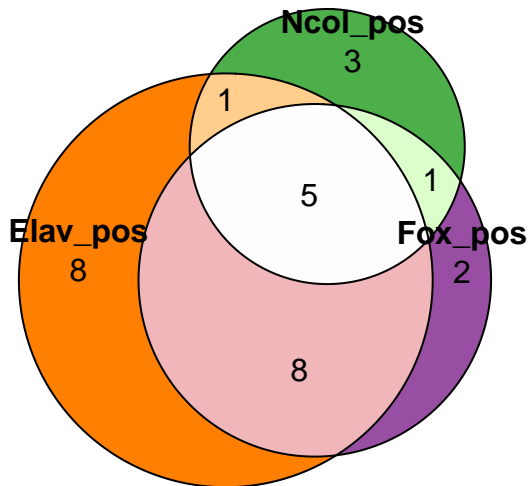
Nvec_vc1.1_XM_048730914.1

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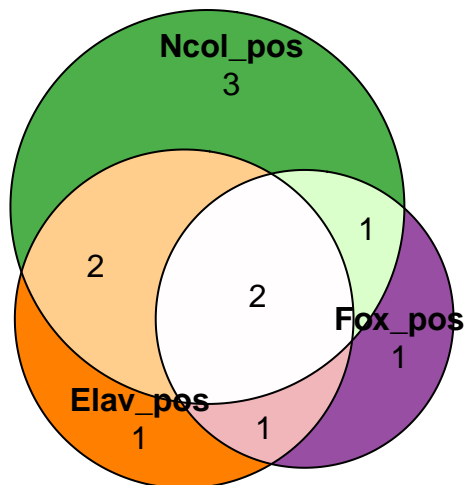


Nvec_vc1.1_XM_048732166.1

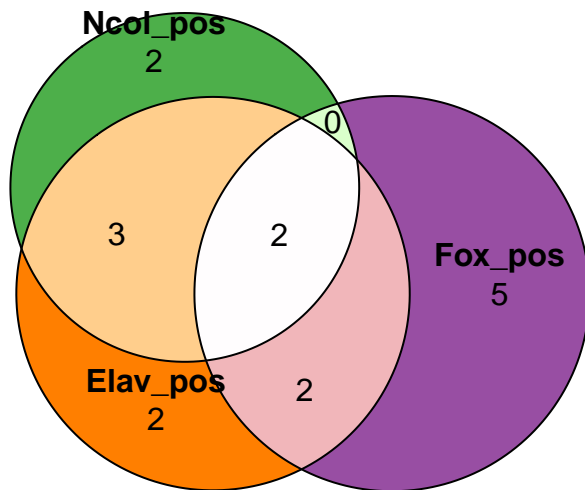
;



Nvec_vc1.1_XM_001624597.3
Ubiquitin C; ubiquitin/ubiquitin/ubiqu

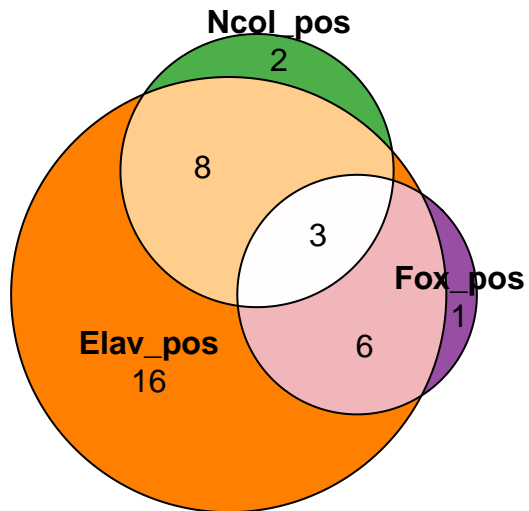


Nvec_vc1.1_XM_048733986.1
BTB:like:AL033529/BACH1/B; BTB/

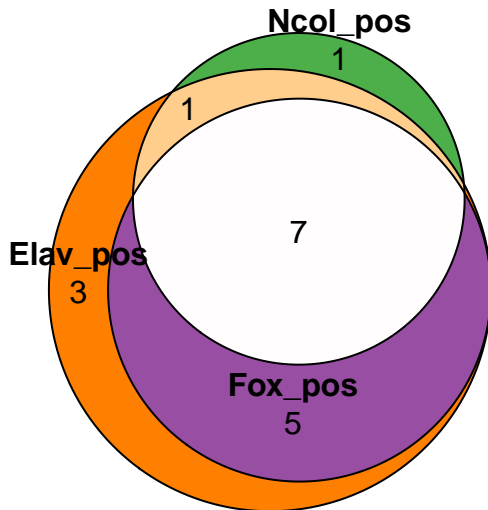


Nvec_vc1.1_XM_032380816.2

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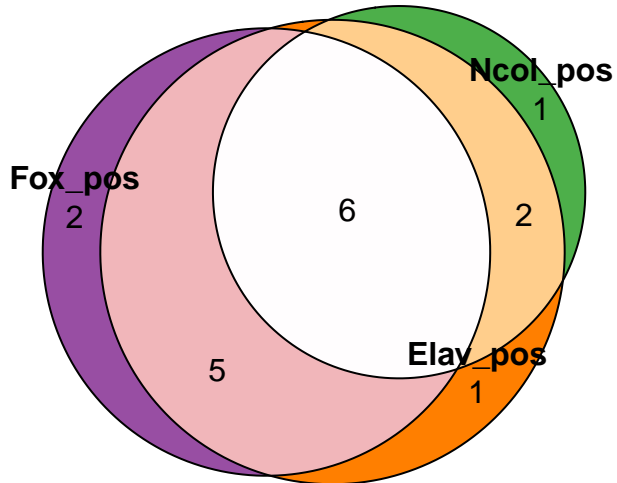


Nvec_vc1.1_XM_001626953.3
; Sel1/



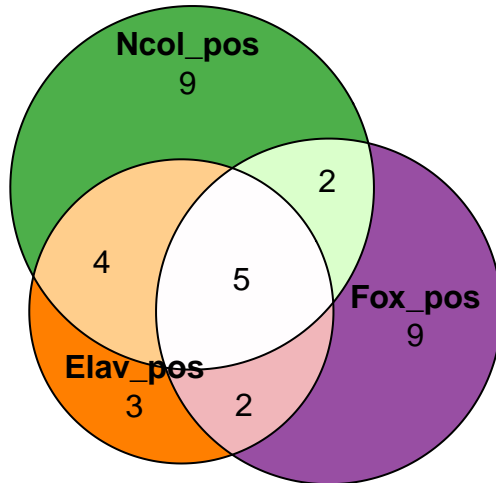
Nvec_vc1.1_XM_032365362.2

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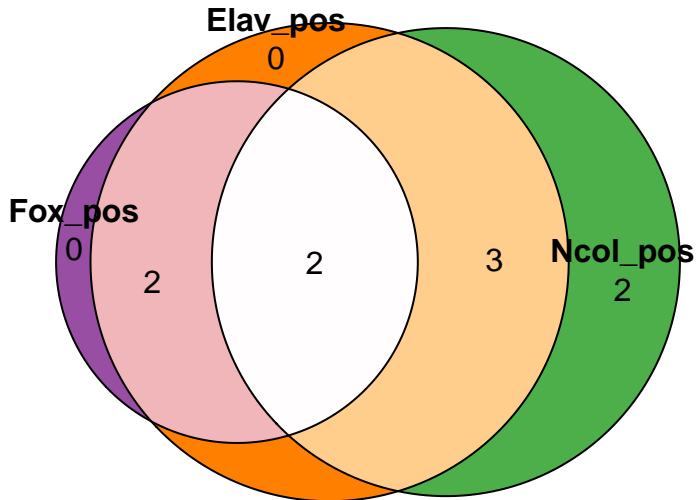


Nvec_vc1.1_XM_001630767.3

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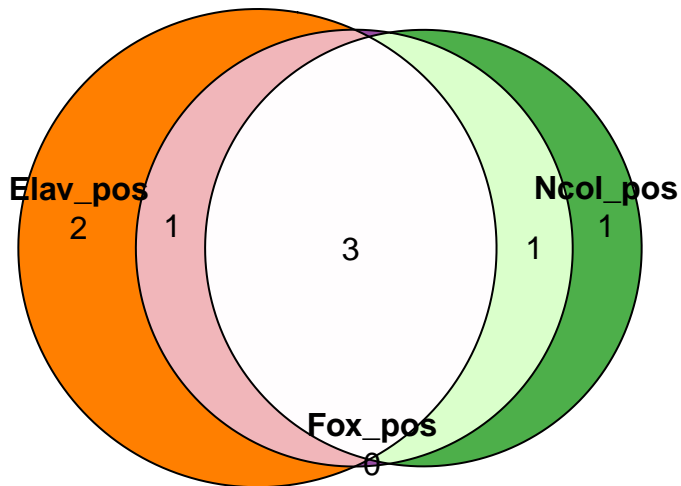


Nvec_vc1.1_XM_001641643.3
cDNA FLJ77442, highly sim;



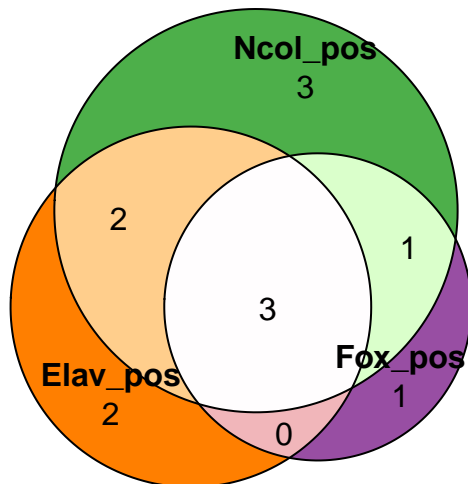
Nvec_vc1.1_XM_001629357.3

Ring finger protein 185; zf-C3HC4_2/



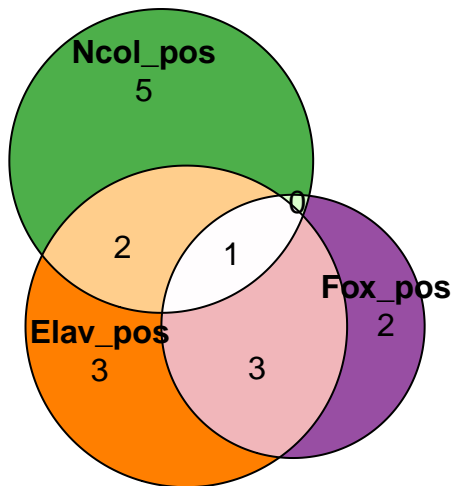
Nvec_vc1.1_XM_001627618.3

cDNA FLJ34613 fis, clone ; Alpha-amylase/



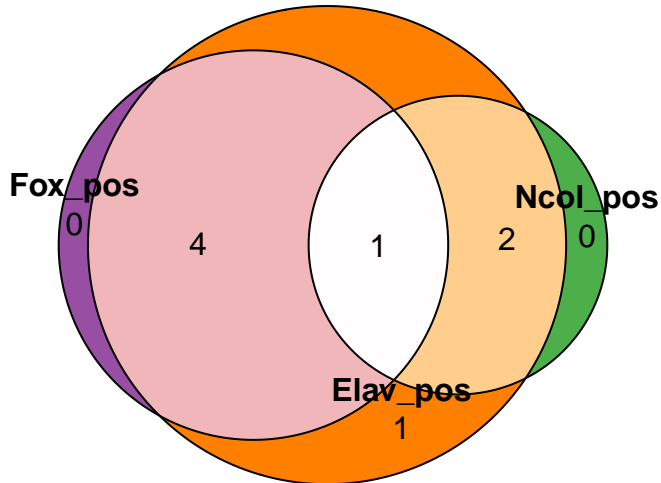
Nvec_vc1.1_XM_048730849.1

E3 ubiquitin-protein liga; zf-B_box/Arf/



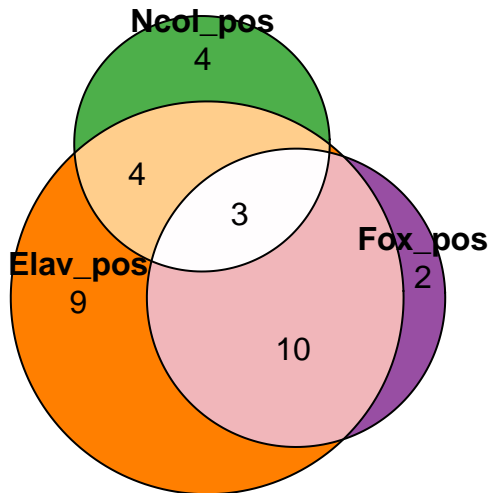
Nvec_vc1.1_XM_001624012.3

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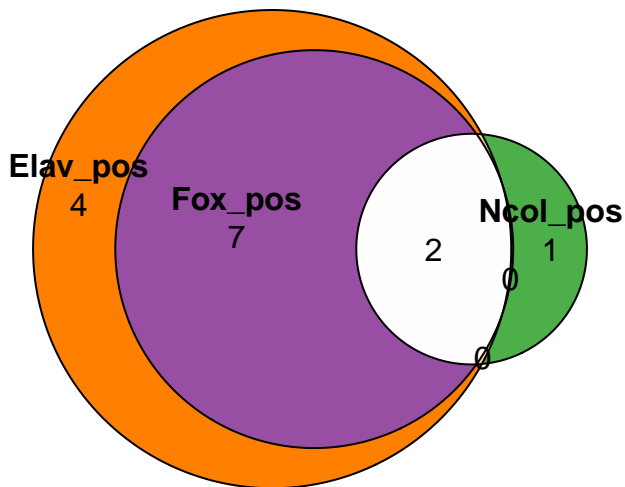
Nvec_vc1.1_XM_032373546.2

Genetic suppressor elemen;

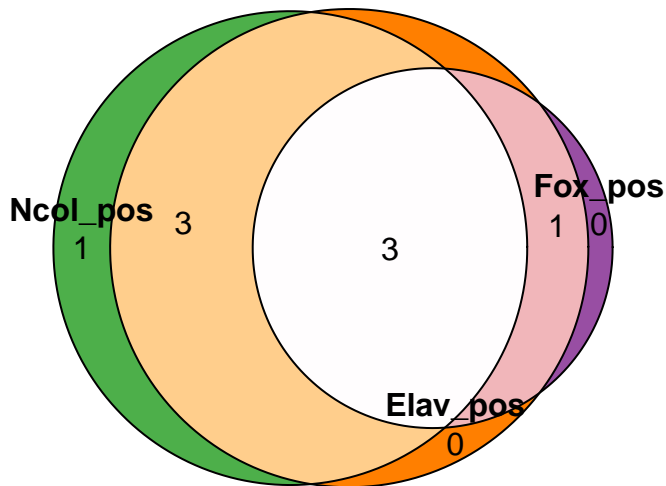


Nvec_vc1.1_XM_001636276.3

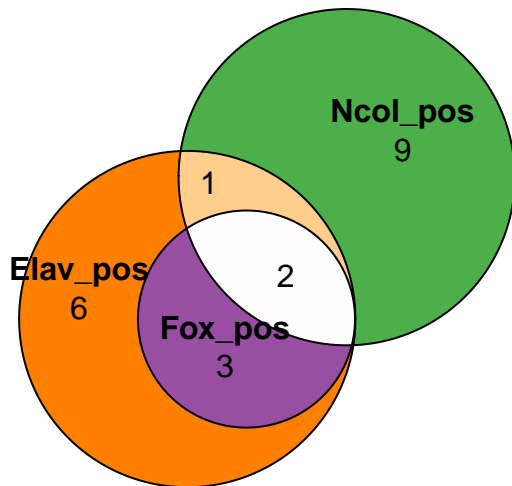
Cytosolic 5'-nucleotidase; 5-nucleotidase/



Nvec_vc1.1_XM_048729416.1
cDNA, FLJ93684, highly si; Mab-21/

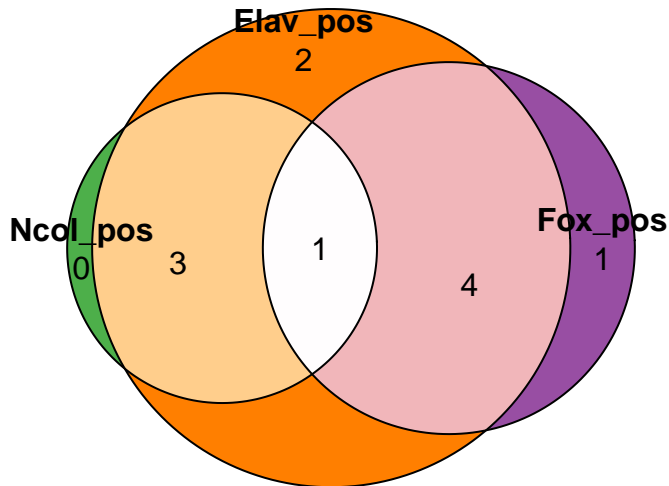


Nvec_vc1.1_XM_032384110.2
AKNA transcript C2;

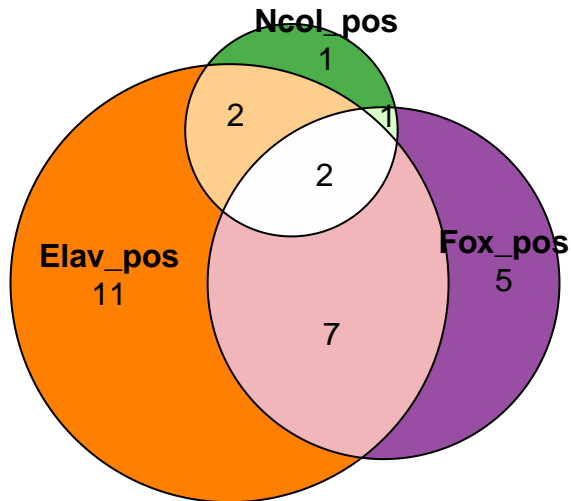


Nvec_vc1.1_XM_001639748.3

Disrupted in renal carcin;

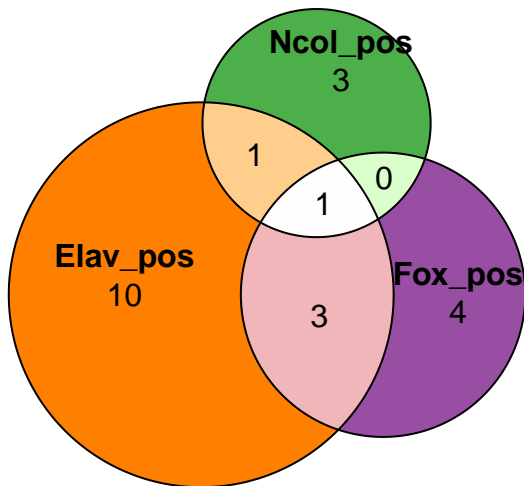


Nvec_vc1.1_XM_001629694.3
XBP1; XBP1



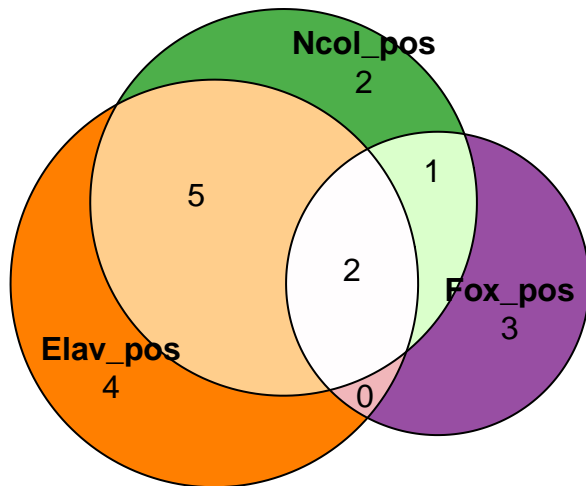
Nvec_vc1.1_XM_032373560.2

cDNA FLJ60383, highly sim; zf-C3HC4/DUF1280/



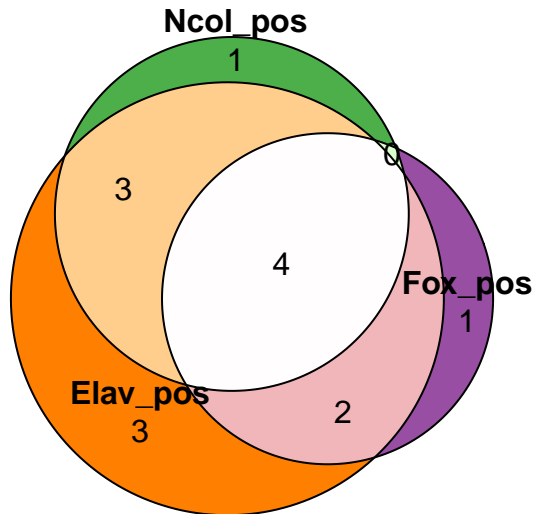
Nvec_vc1.1_XM_032365869.2

E3 ubiquitin-protein liga; FHA/zf-RING_2/



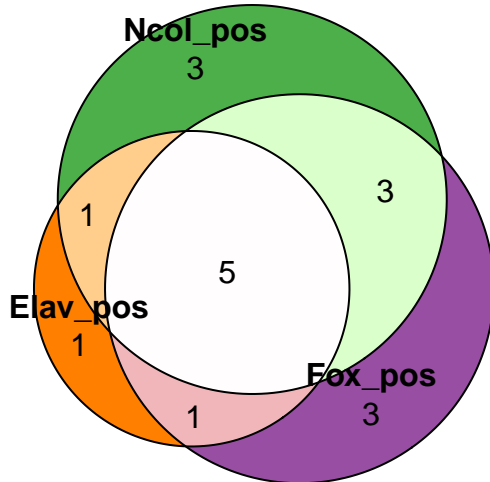
Nvec_vc1.1_XM_048727558.1

Neuroblastoma–amplified s; Nbas_N/Sec39/



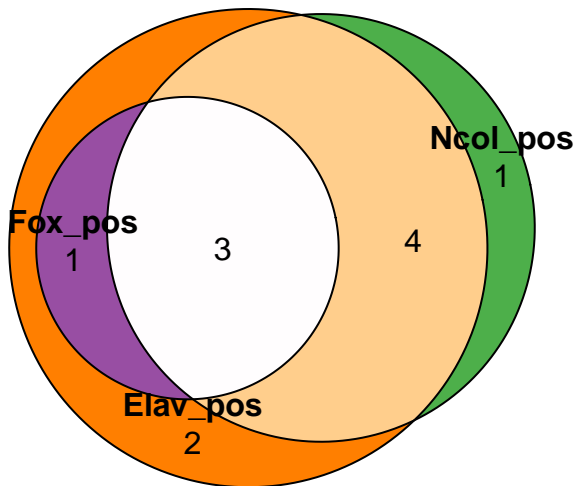
Nvec_vc1.1_XM_032383908.2

V-type proton ATPase subu; V_ATPase_I/V_ATPase_I/V_A



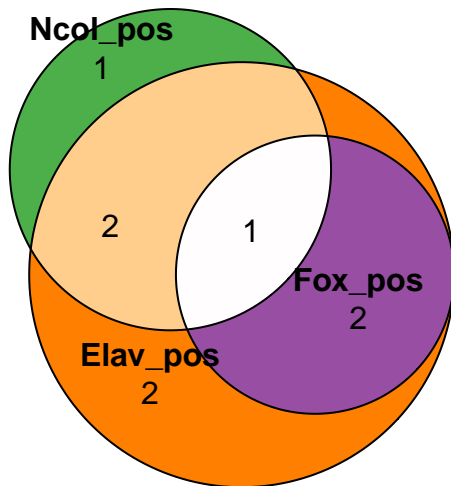
Nvec_vc1.1_XM_001625257.3

Major facilitator superfa; MFS_1_like/MFS_1/



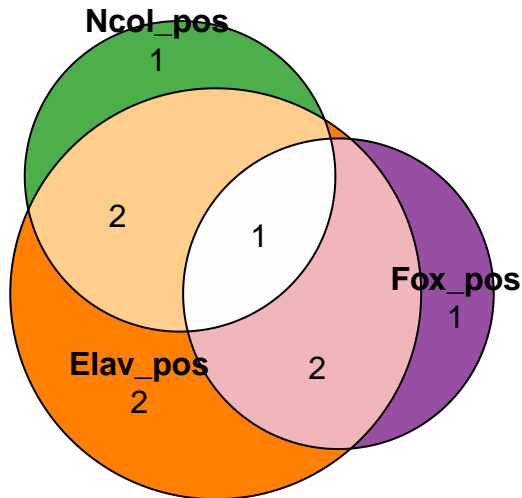
Nvec_vc1.1_XM_048726027.1

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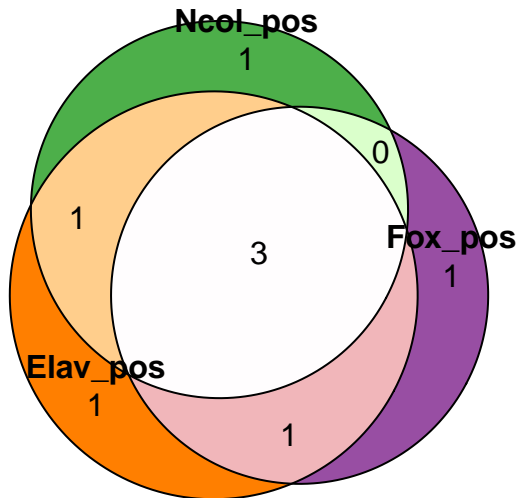


Nvec_vc1.1_XM_032366411.2

Dipeptidyl peptidase 4; DPPIV_N/Peptidase_S9/

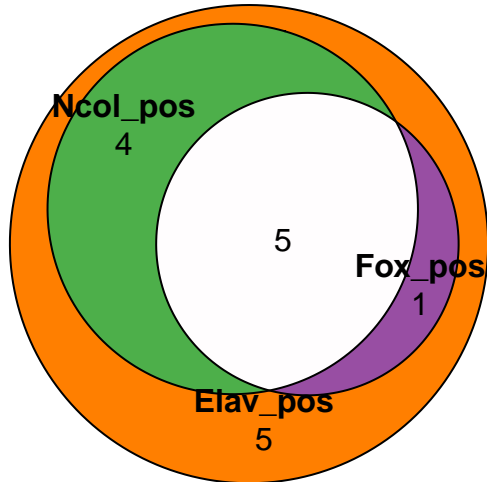


Nvec_vc1.1_XM_048722217.1
Ras-related protein Rab-3; Ras/



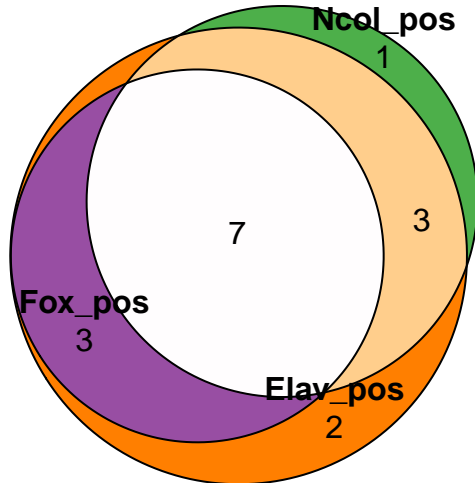
Nvec_vc1.1_XM_048724342.1

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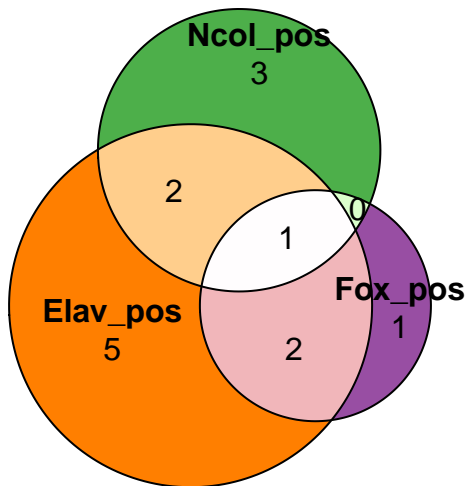


Nvec_vc1.1_XM_032372079.2

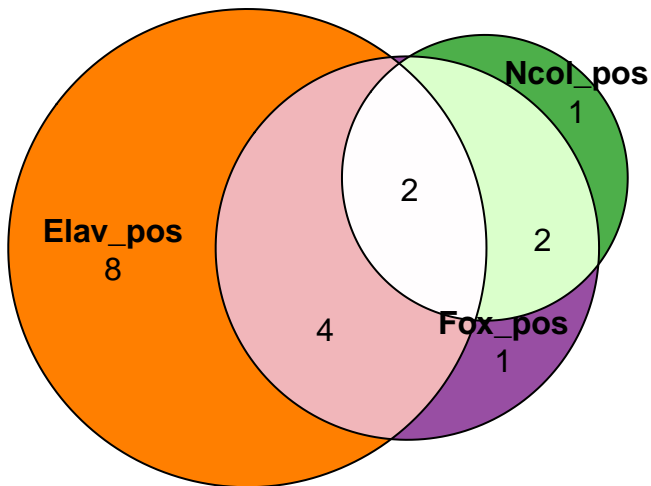
;



Nvec_vc1.1_XM_032374515.2
; Death/

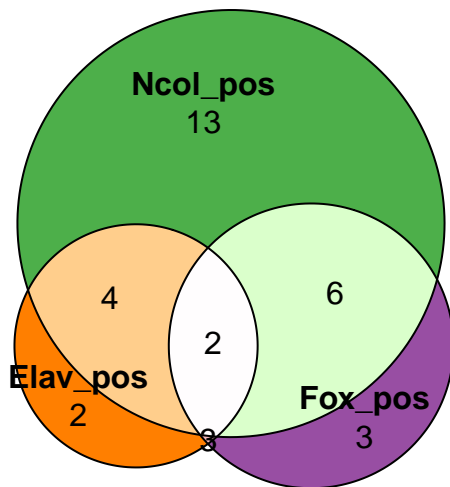


Nvec_vc1.1_XM_001631013.3
START domain containing 1; START/



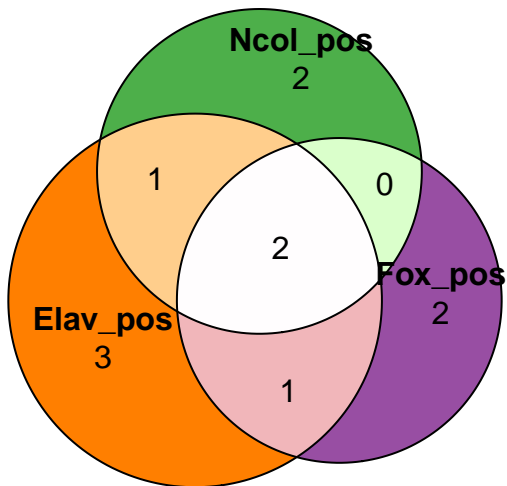
Nvec_vc1.1_XM_048723016.1

Synaptotagmin:like:ESYT1/; DUF2404/C2/



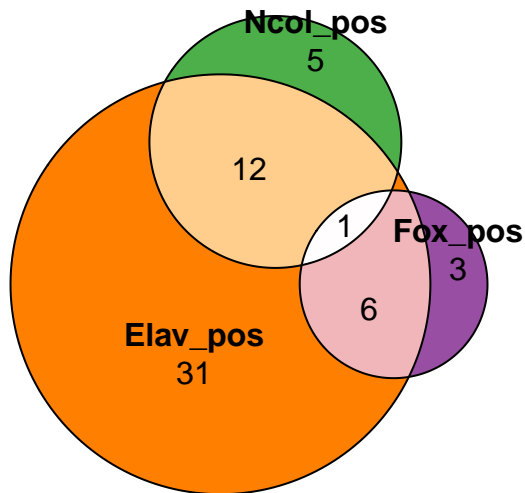
Nvec_vc1.1_XM_032385056.2

Multiple epidermal growth; CUB/Kelch_4/Kelch_3/PSI/E

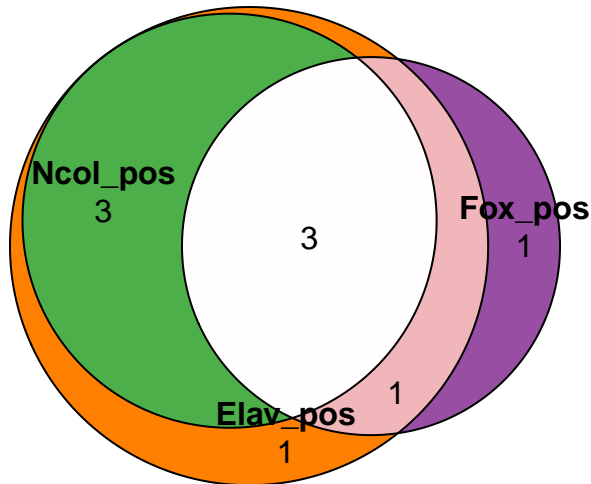


Nvec_vc1.1_XM_048726174.1

zf-C2H2.HG6.2:MYT1/MYT1L/; MYT1/MYT1L/ST18

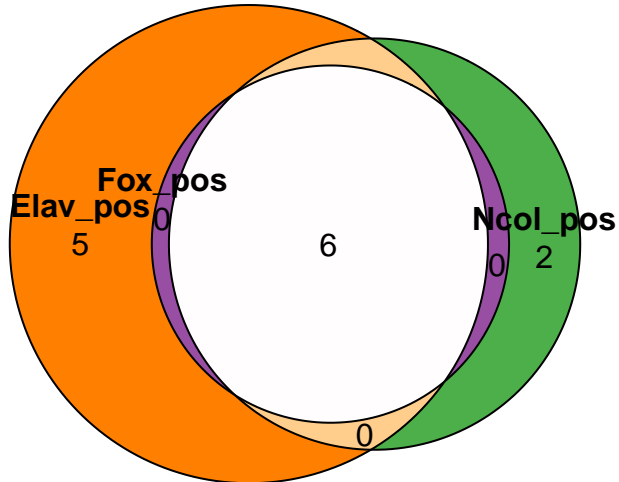


Nvec_vc1.1_XM_048722653.1
cDNA FLJ16032 fis, clone ; Kunitz_BPTI/



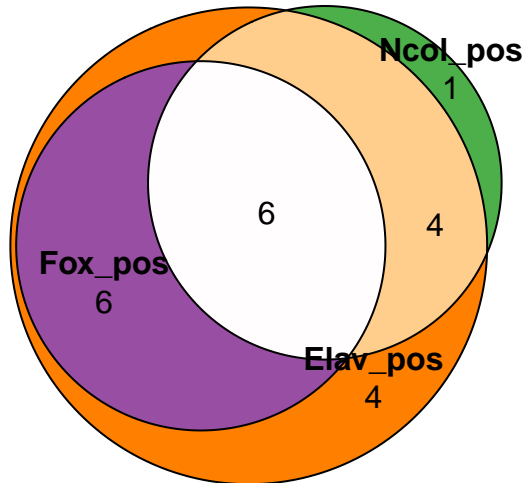
Nvec_vc1.1_XM_048719892.1

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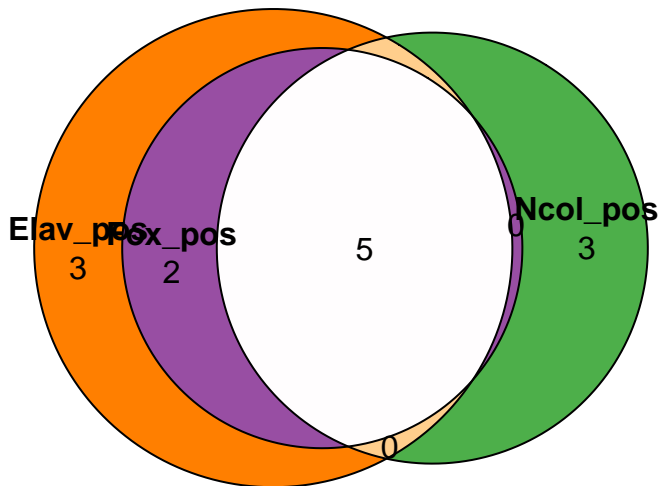


Nvec_vc1.1_XM_048732682.1

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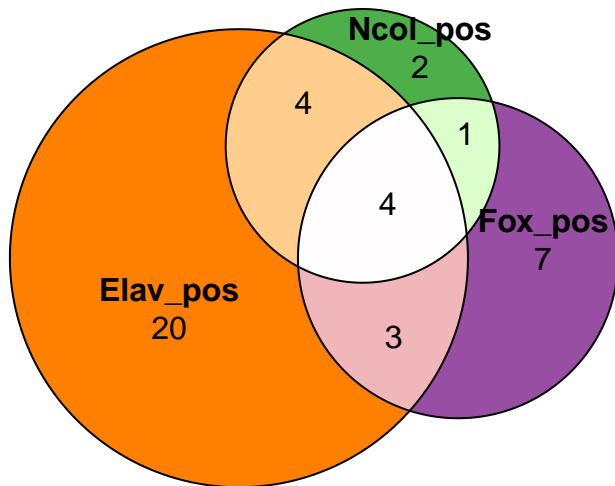


Nvec_vc1.1_XM_032370307.2
Ubiquitin-associated prot; DUF3697/

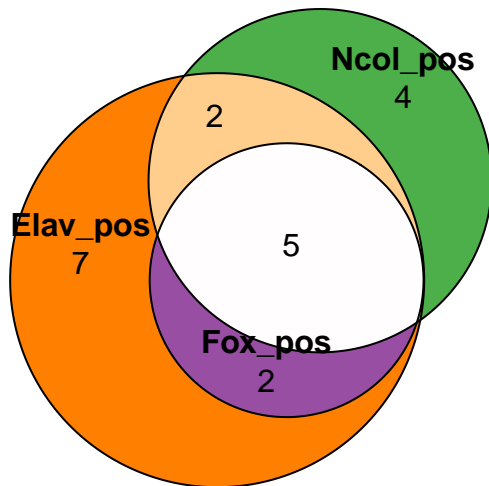


Nvec_vc1.1_XM_032372148.2

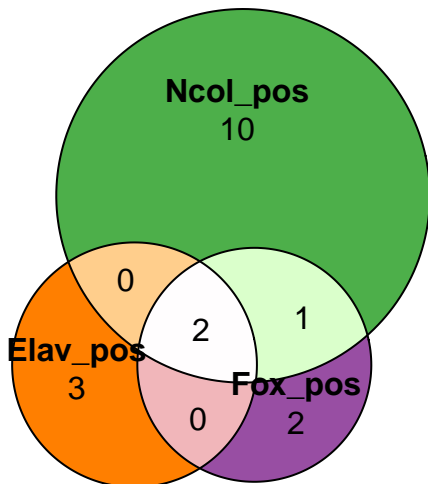
Potassium channel subfami; Ion_trans_2/BK_channel_a/



Nvec_vc1.1_XM_048721594.1
Phosphatidylinositol-bind; ANTH/

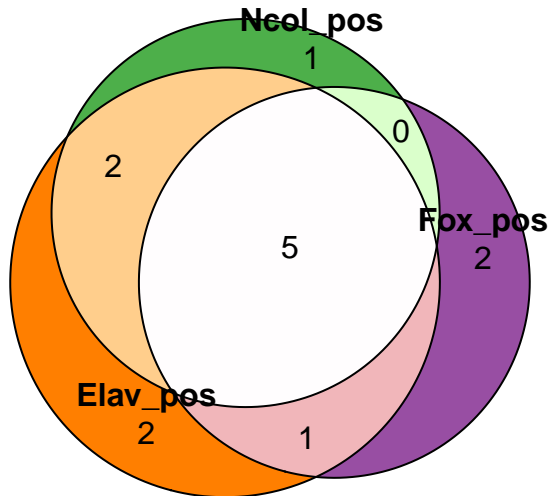


Nvec_vc1.1_XM_032380092.2
Cdh1; Cadherin/Cadherin/Cadheri



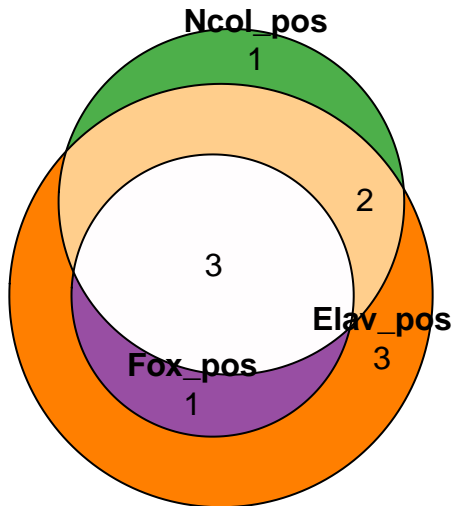
Nvec_vc1.1_XM_032374271.2

Golgin subfamily A member; GOLGA2L5/GOLGA2L5/



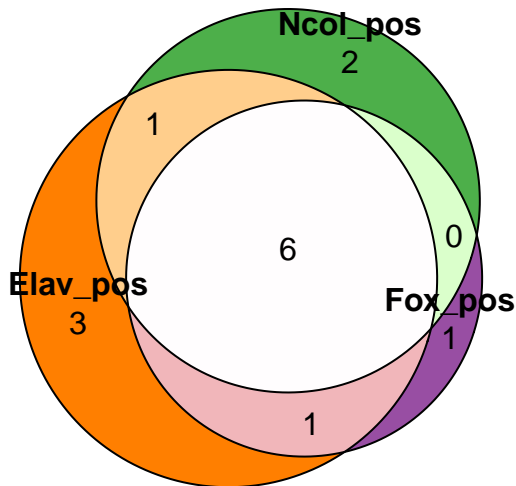
Nvec_vc1.1_XM_001637845.3

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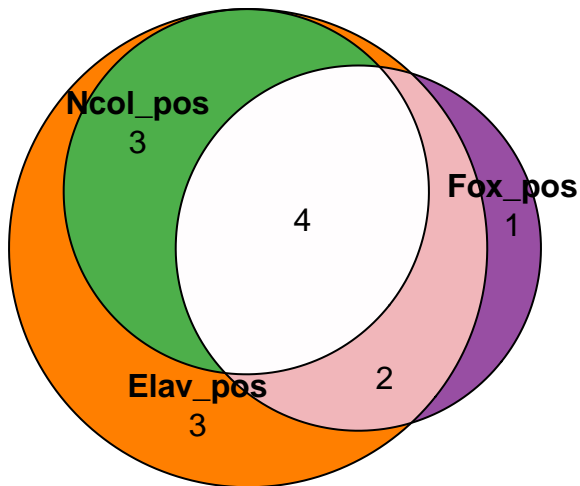


Nvec_vc1.1_XM_001628600.3

Pkinase:SNRK; Pkinase.HG167.0:SNRK

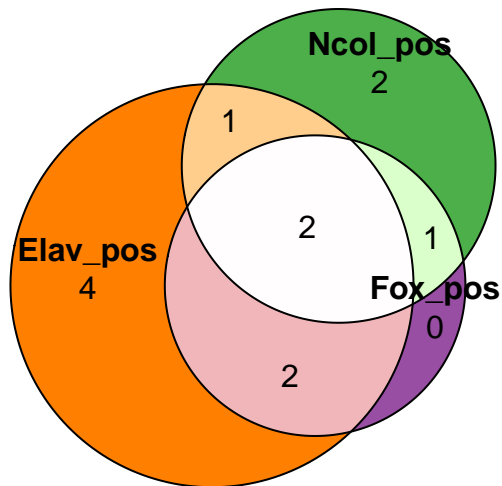


Nvec_vc1.1_XM_032367035.2
FtsH homolog; AAA/Peptidase_M41/



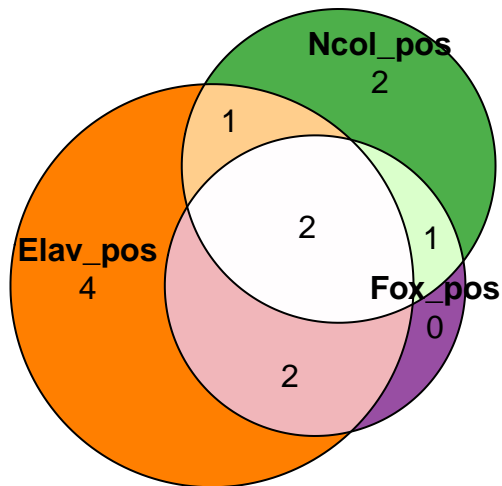
Nvec_vc1.1_XM_048732433.1

WD repeat domain 44; WD40/WD40/WD40/



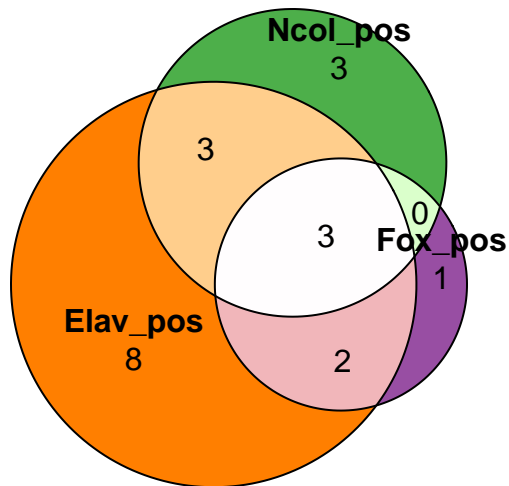
Nvec_vc1.1_XM_048732806.1

Growth hormone-regulated ; RabGAP-TBC/



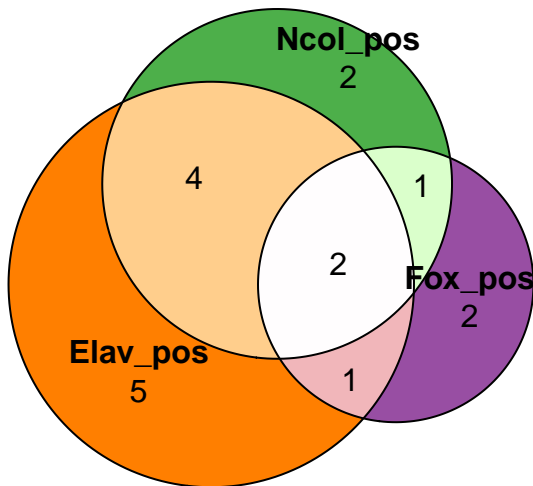
Nvec_vc1.1_XM_001635172.3

HLH.HG1.38:MXD1/MXD3/MXD4; MXD1/MXD3/MXD4/MXI1



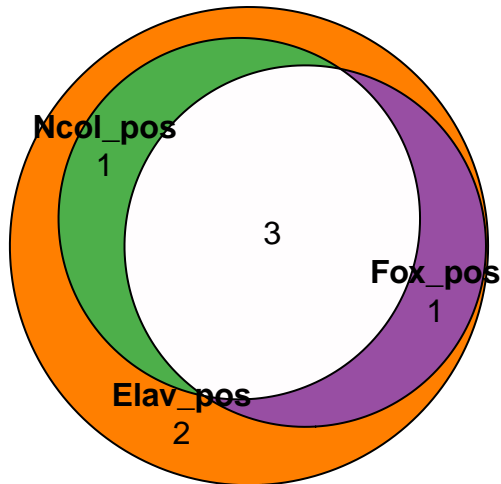
Nvec_v1g176307

p8 protein (Candidate of ; Phospho_p8/



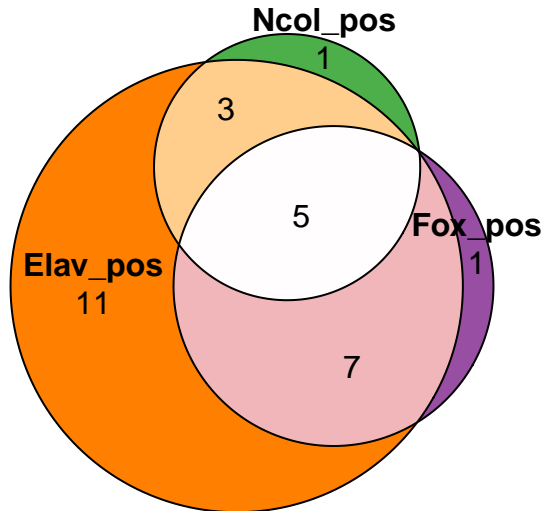
Nvec_vc1.1_XM_032366255.2

INO80 complex subunit D; zf-C3Hc3H/

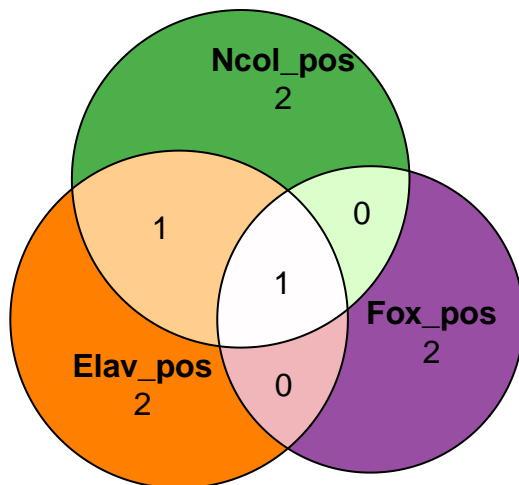


Nvec_vc1.1_XM_001640716.3

Cytoplasmic polyadenylati;

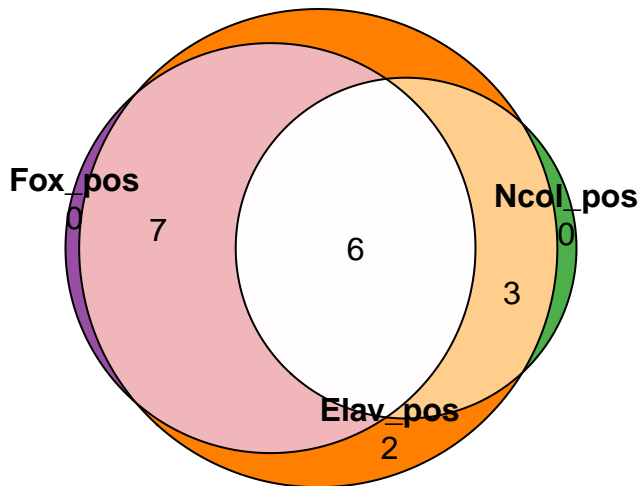


Nvec_vc1.1_XM_032387187.2
B-cell receptor-associate; Bap31/



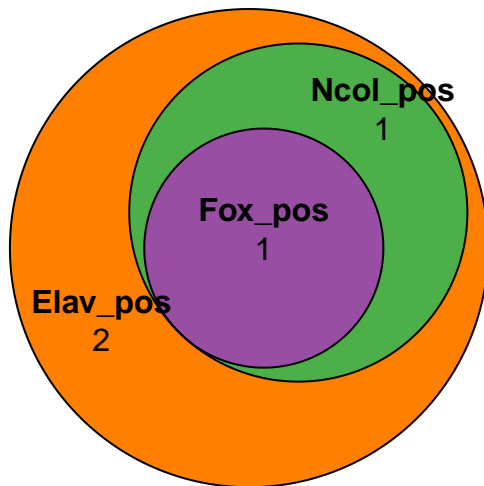
Nvec_vc1.1_XM_032372848.2

Transthyretin; DUF3504/Transthyretin/



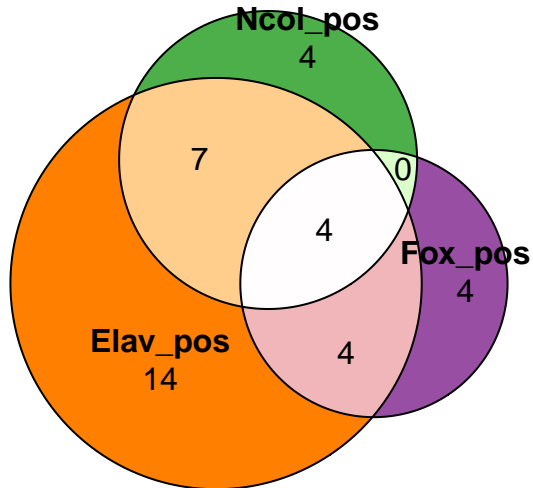
Nvec_vc1.1_XM_032382085.2

Inositol hexaphosphate ki; IPK/



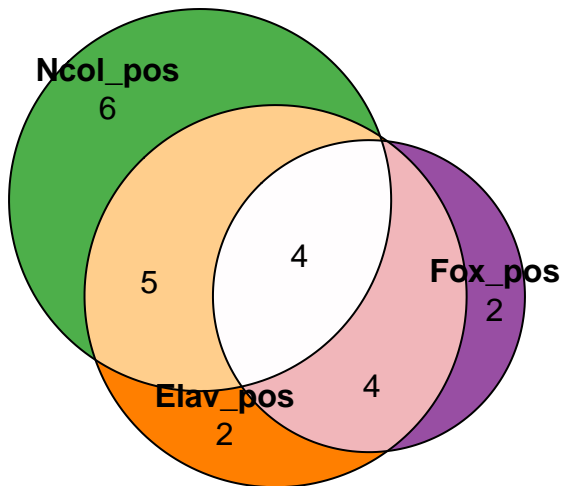
Nvec_vc1.1_XM_001630781.3

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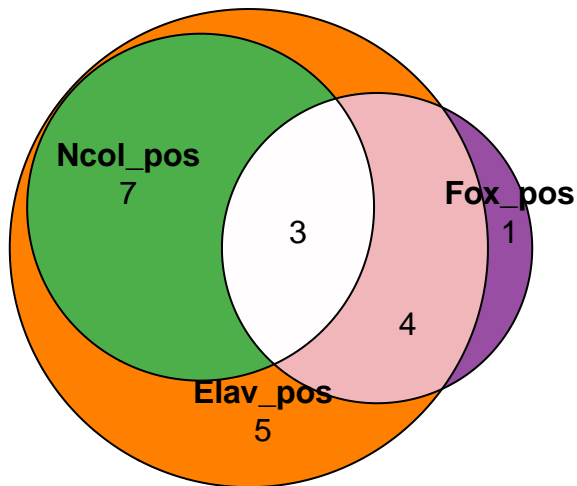
Nvec_vc1.1_XM_032379476.2

DnaJ homolog subfamily B ; DnaJ/CTDII/



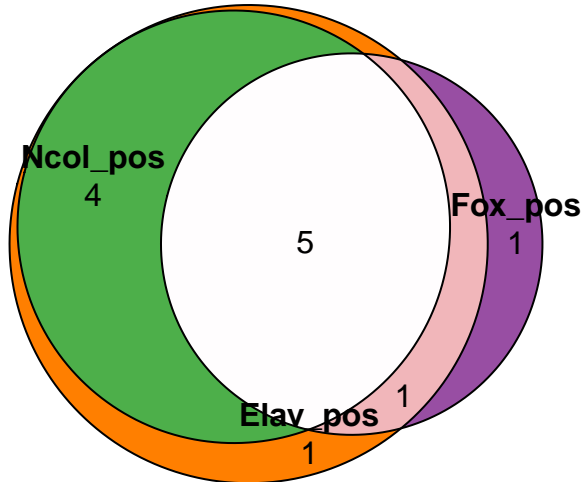
Nvec_vc1.1_XM_032386164.2

Synaptotagmin:CPNE3/CPNE5; Synaptotagmin.HG11.2:CPNE

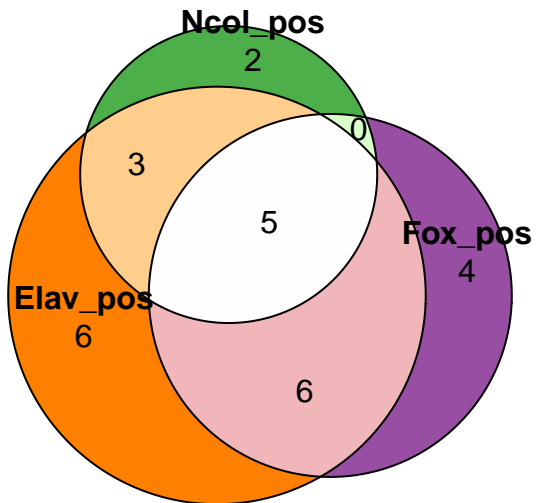


Nvec_vc1.1_XM_032375913.2

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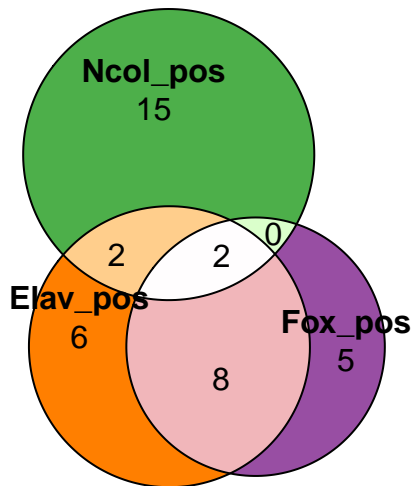


Nvec_vc1.1_XM_032369224.2
; E2R135/



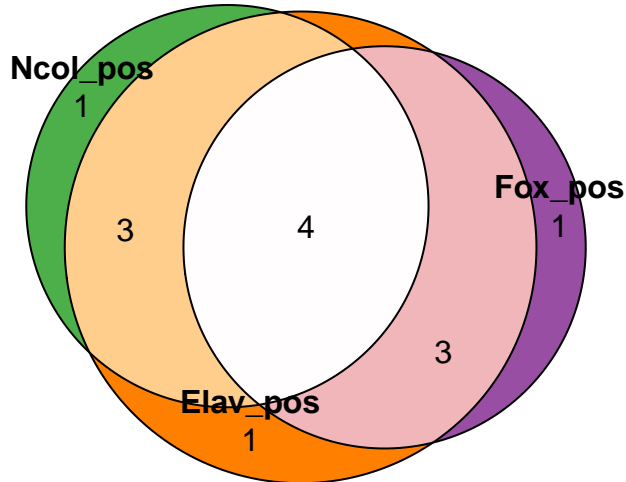
Nvec_vc1.1_XM_032365463.2

ion-NaKCa:like:KCNB1/KCNB; Ion_trans/



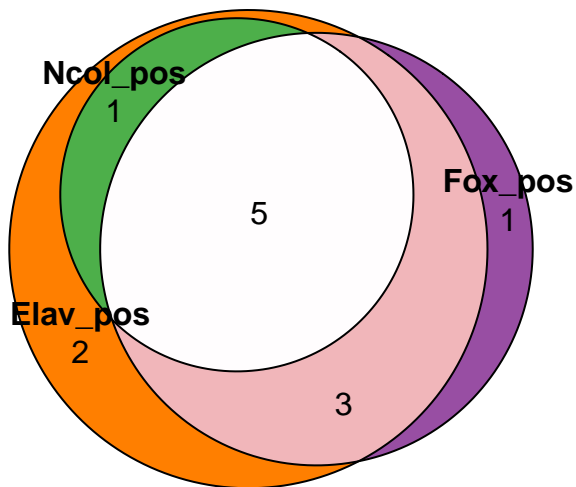
Nvec_vc1.1_XM_048720899.1

NEDD4-like E3 ubiquitin-p; WW/WW/WW/WW/HECT/



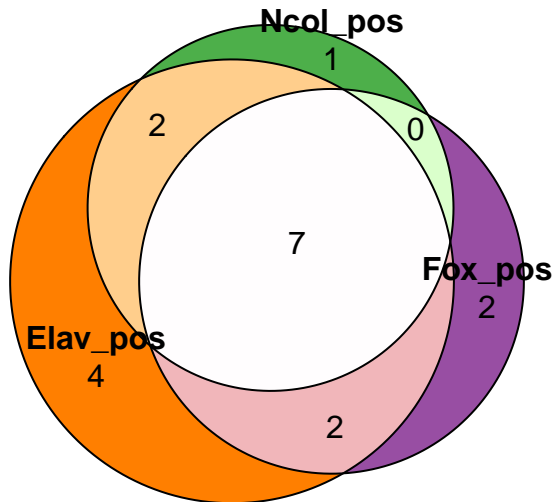
Nvec_vc1.1_XM_032387194.2

EF-hand:DYTN; EF-hand.HG1.4:DYTN

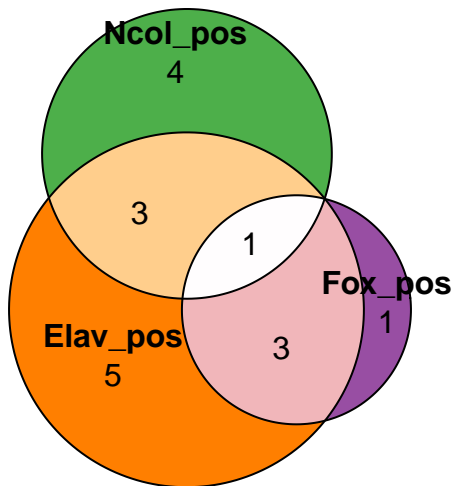


Nvec_vc1.1_XM_048723406.1

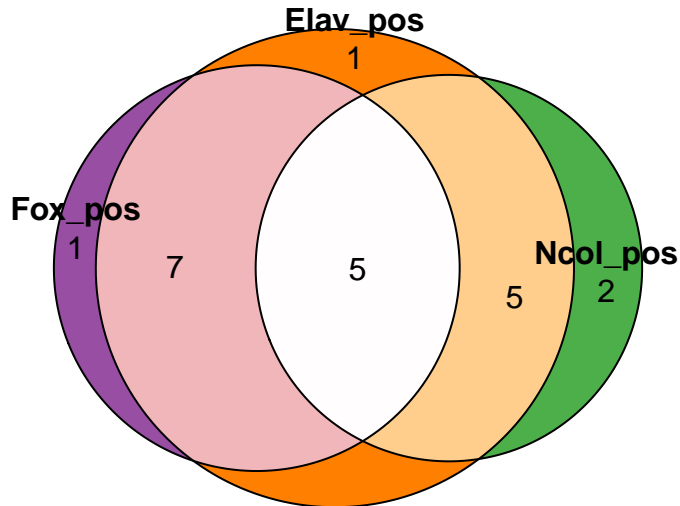
Protein phosphatase 1 reg; KLRAQ/TTKRSYEDQ/



Nvec_vc1.1_XM_001627682.3
Epididymis luminal protei; Serpin/

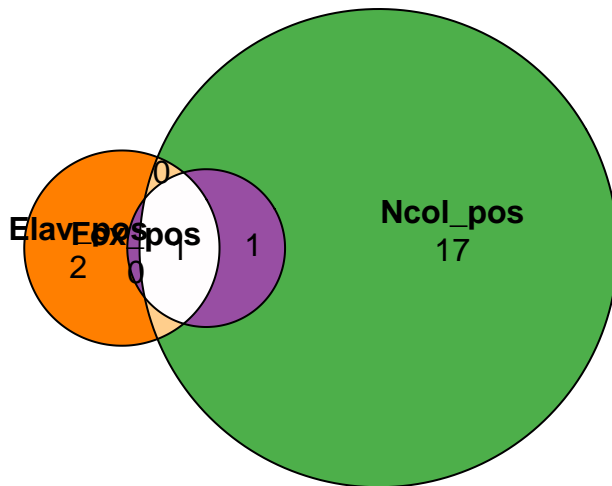


Nvec_vc1.1_XM_048722729.1
ELL-associated factor 2; EAF/



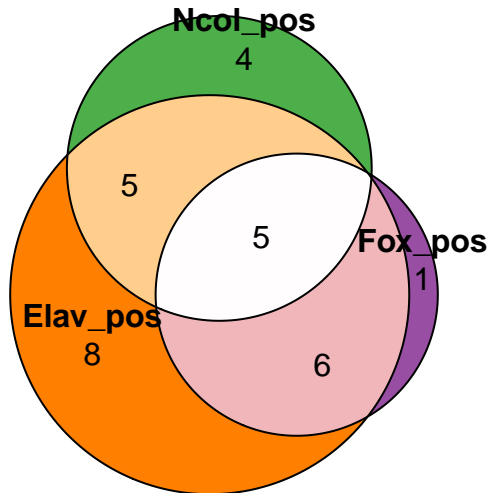
Nvec_vc1.1_XM_048724767.1

F-box/WD repeat-contains; WD40/WD40/WD40/



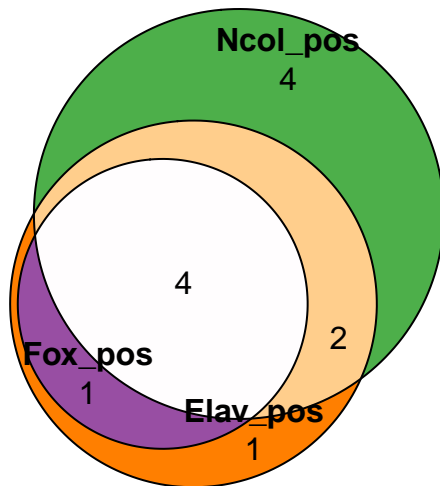
Nvec_vc1.1_XM_032372812.2

Uncharacterized membrane ; DUF1180/

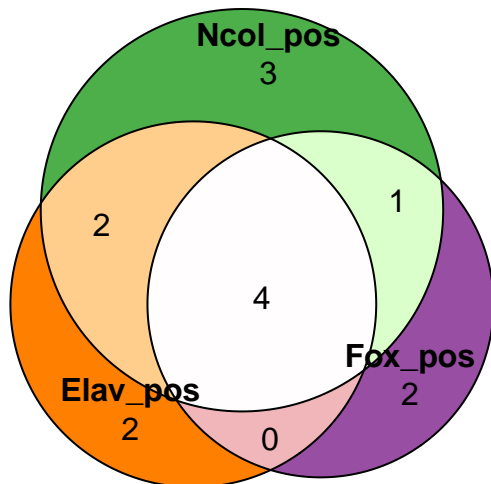


Nvec_vc1.1_XM_048729965.1

Pecanex-like protein 1;

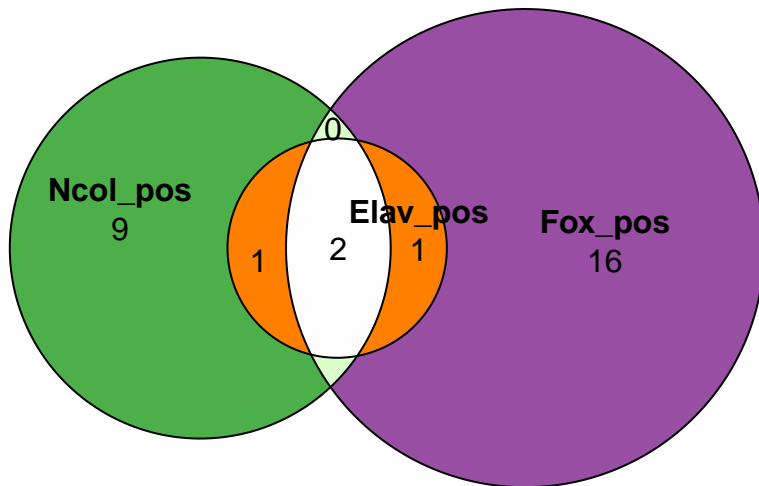


Nvec_vc1.1_XM_032383224.2
Breast carcinoma amplifie; BCAS3/



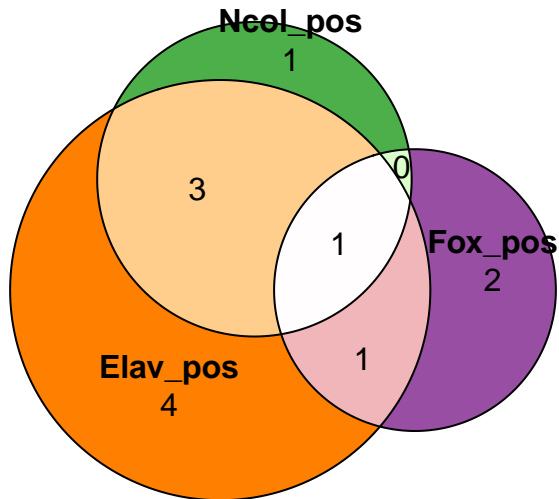
Nvec_vc1.1_XM_032376608.2

Homeodomains.HG3.20:MEIS1; MEIS1/MEIS2/MEIS3



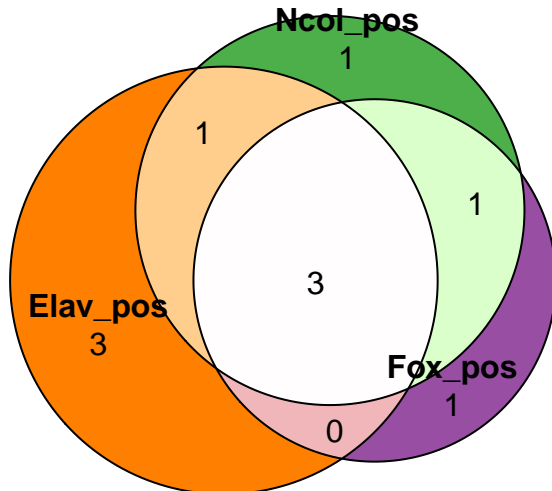
Nvec_vc1.1_XM_032375358.2

LinkerHistone:like:HP1BP3; Linker_histone/



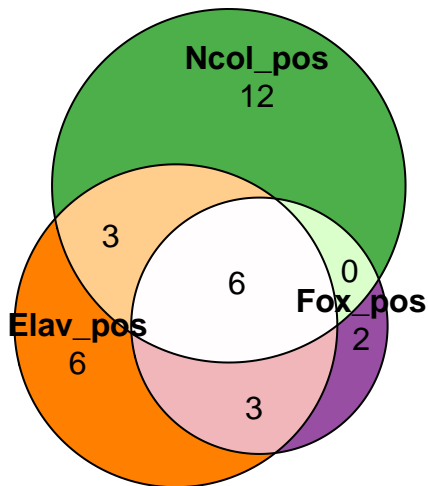
Nvec_vc1.1_XM_001634221.3

Oxidoreductase NAD-binding; FAD_binding_6/NAD_binding



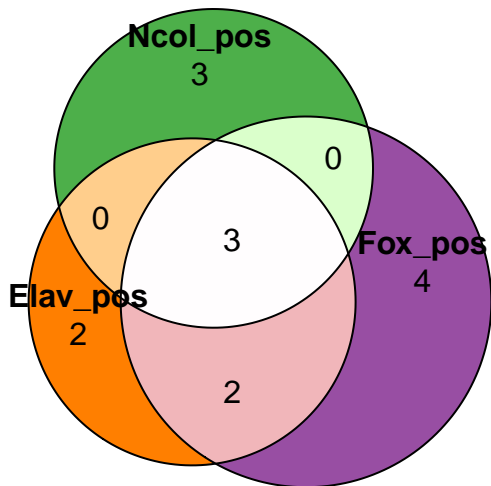
Nvec_vc1.1_XM_032366934.2

WD repeat-containing prot; WD40/WD40/WD40/WD40/WD40/

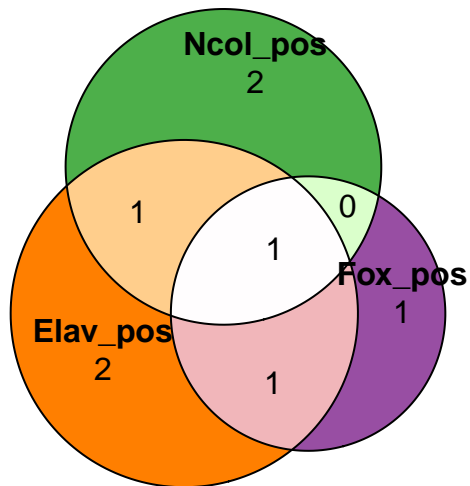


Nvec_vc1.1_XM_032364831.2

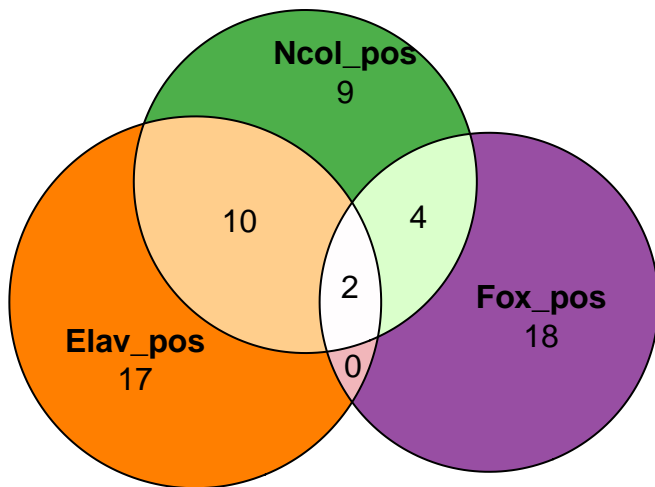
;



Nvec_vc1.1_XM_001640511.3
; Methyltransf_18/

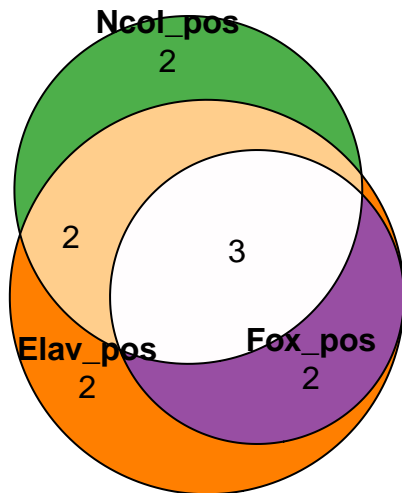


Nvec_vc1.1_XM_048720895.1
Metastasis suppressor pro; IMD/

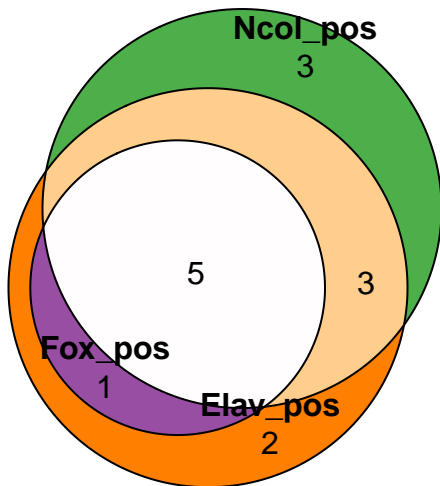


Nvec_vc1.1_XM_048734077.1

Protein scribble homolog; LRR_4/LRR_1/LRR_8/LRR_4/P

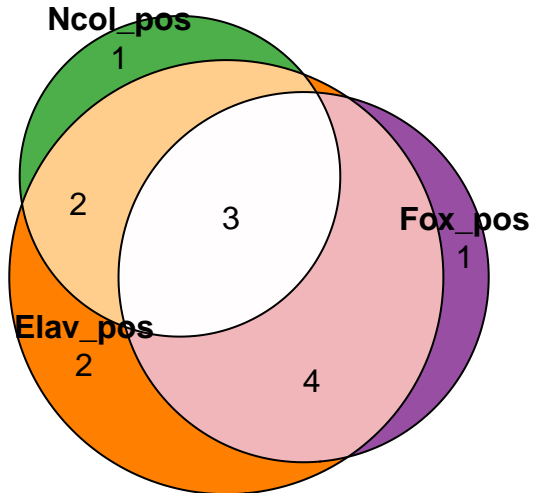


Nvec_vc1.1_XM_032384787.2
Phosphatidylinositide pho; Syja_N/

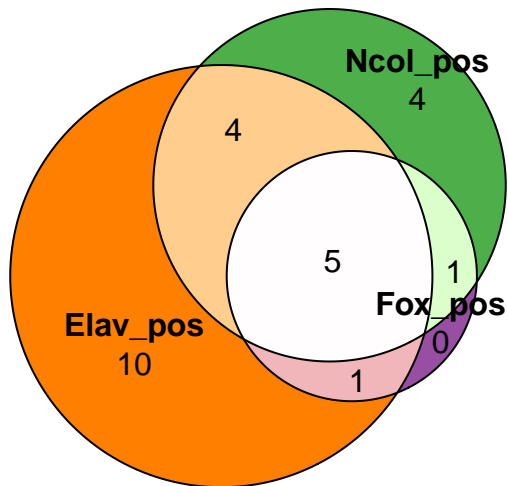


Nvec_vc1.1_XM_032363186.2

Signal transducing adapte; VHS/SH3_1/

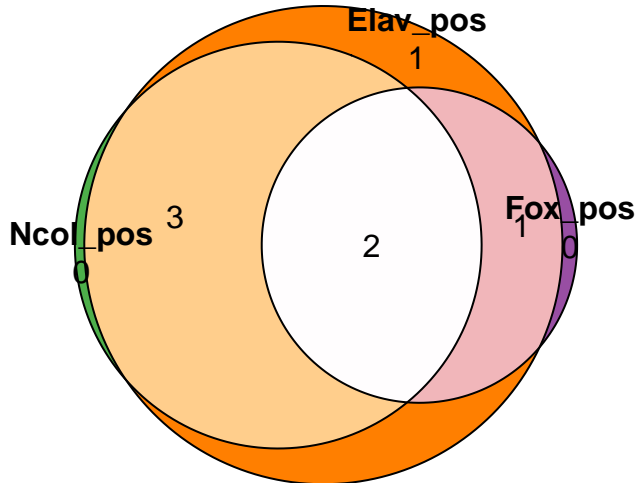


Nvec_vc1.1_XM_048727530.1
E3 ubiquitin-protein liga; IBR/IBR/



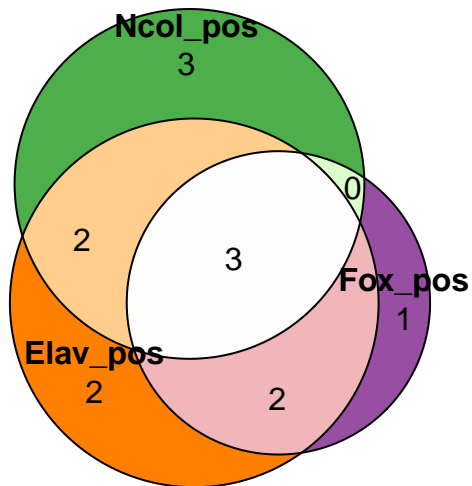
Nvec_vc1.1_XM_001633567.3

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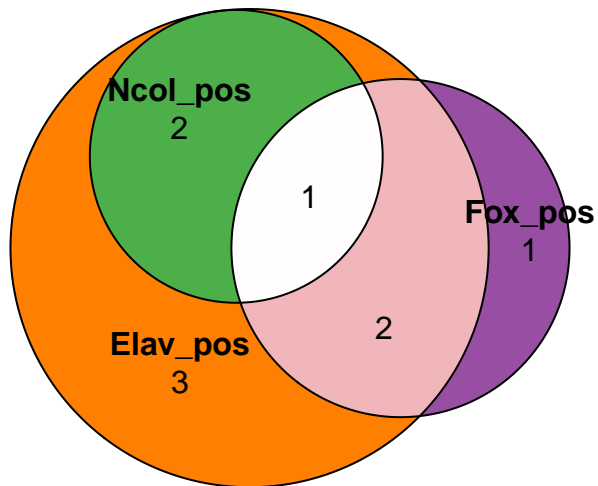
Nvec_vc1.1_XM_048725899.1

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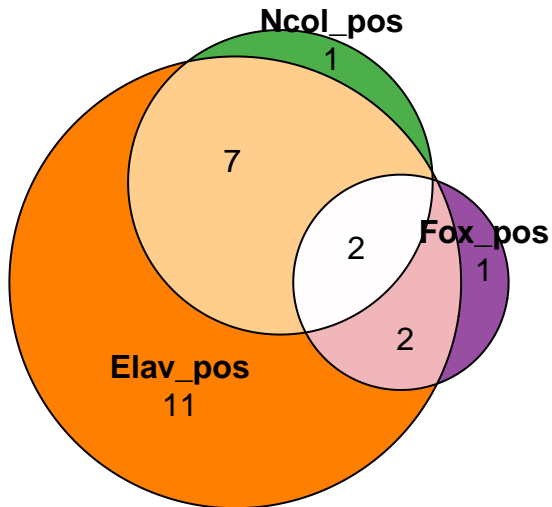


Nvec_vc1.1_XM_001637787.3

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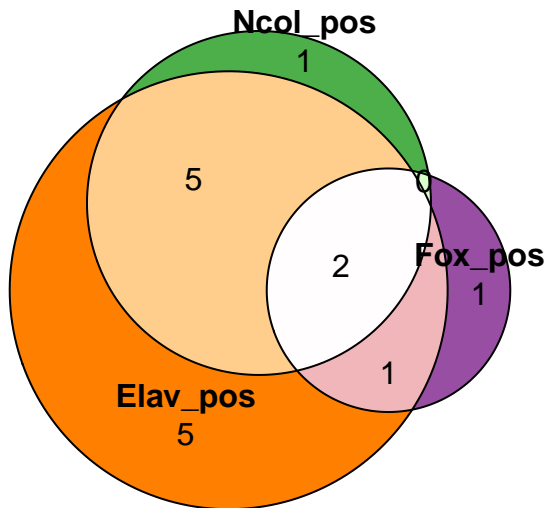


Nvec_vc1.1_XM_001630398.3
; PALP/

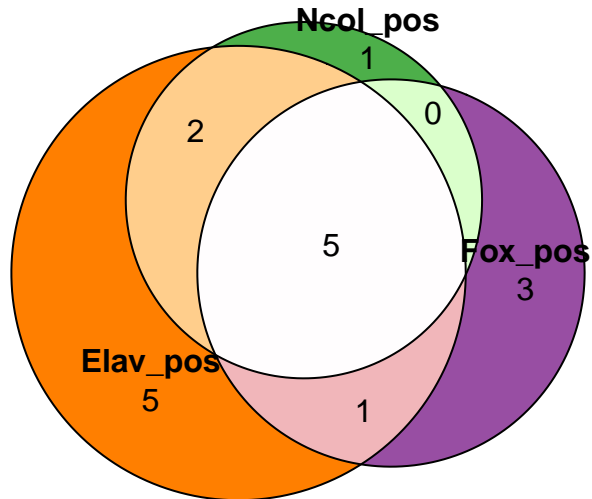


Nvec_vc1.1_XM_001627424.3

Ferritin; Ferritin/

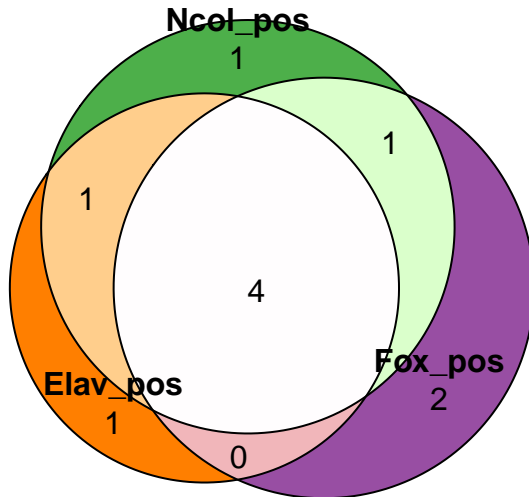


Nvec_vc1.1_XM_001632217.3
; WES_acyltransf/DUF1298/

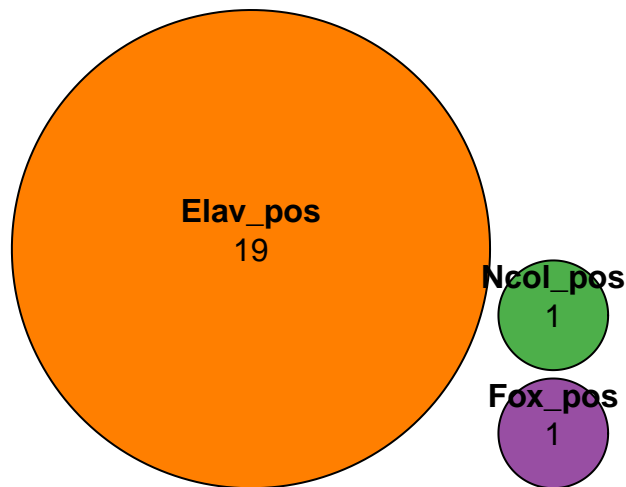


Nvec_vc1.1_XM_032369434.2

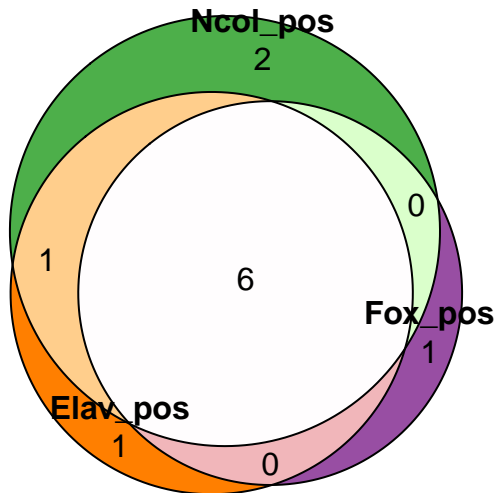
Juxtaposed with another z; zf-C2H2_4/zf-C2H2_4/



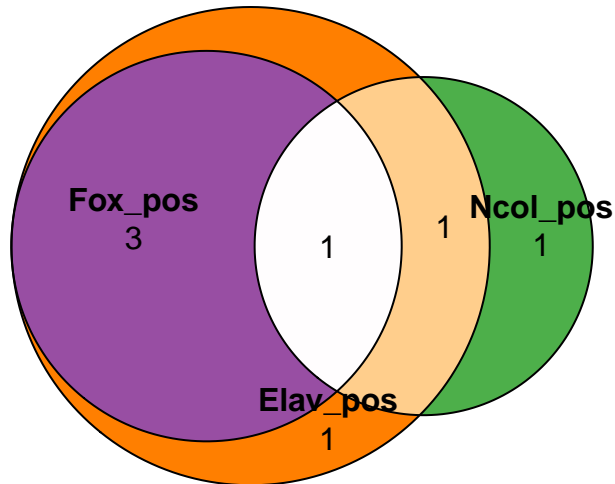
Nvec_vc1.1_XM_032374725.2
cDNA FLJ58014, highly sim; MA3/MA3/



Nvec_vc1.1_XM_001626640.3
HCG1811539; Thioredoxin/

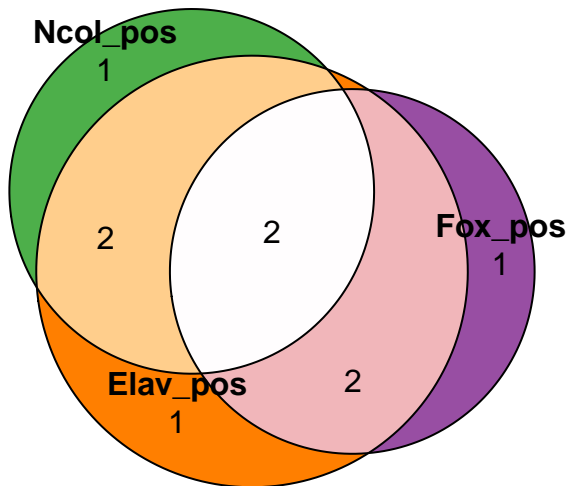


Nvec_vc1.1_XM_001640360.3
SplA/ryanodine receptor d; SPRY/



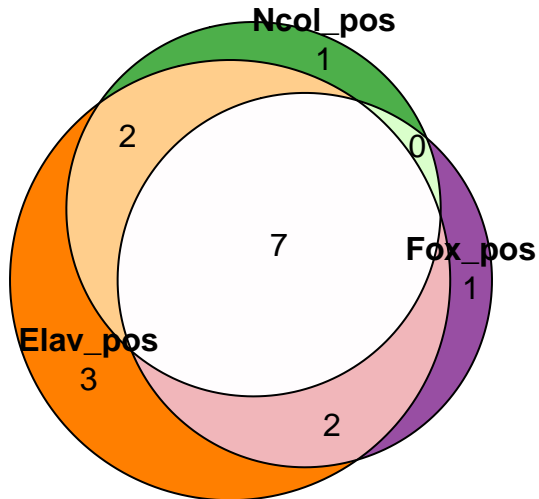
Nvec_vc1.1_XM_032379058.2

RIO kinase 3 (Yeast); RIO1/

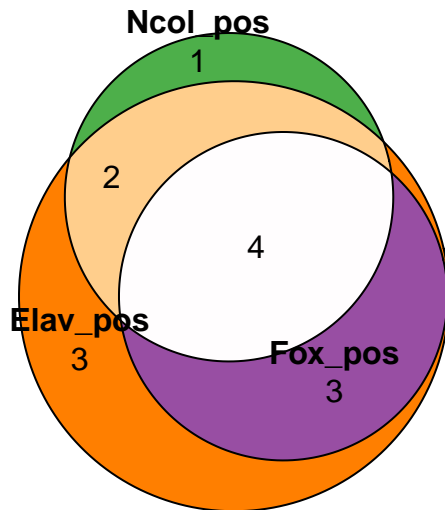


Nvec_vc1.1_XM_032380113.2

SEC14-like 4 (*S. cerevisi*; CRAL_TRIO/GOLD_2/

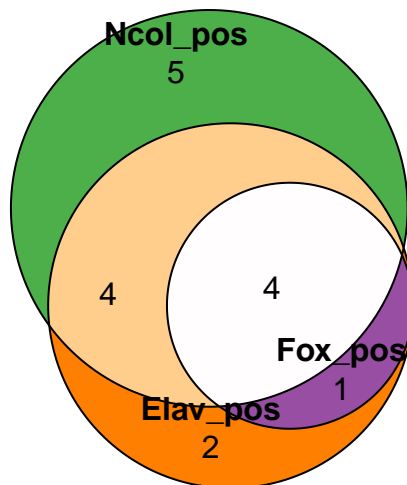


Nvec_vc1.1_XM_048726463.1
cDNA FLJ77587, highly sim;

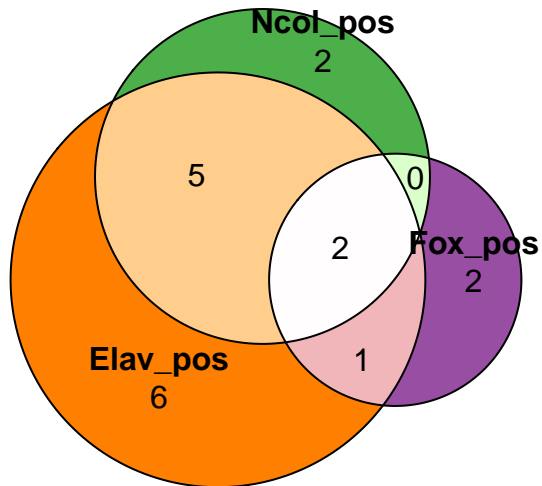


Nvec_vc1.1_XM_048724299.1

Protein DENND6B; SPA/

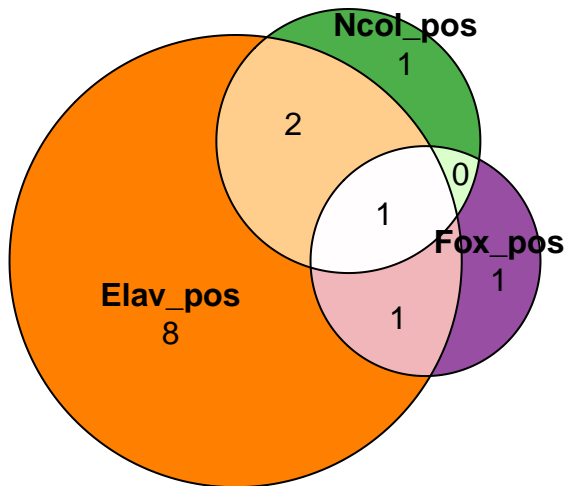


Nvec_vc1.1_XM_001641484.3
cDNA, FLJ94678, highly si; BAG/



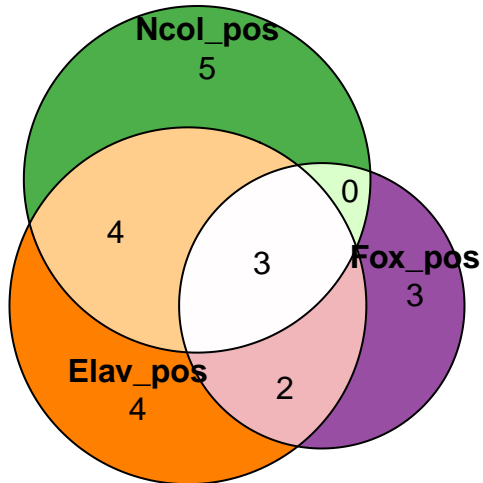
Nvec_vc1.1_XM_032367033.2

DnaJ homolog subfamily B ; DnaJ/CTDII/



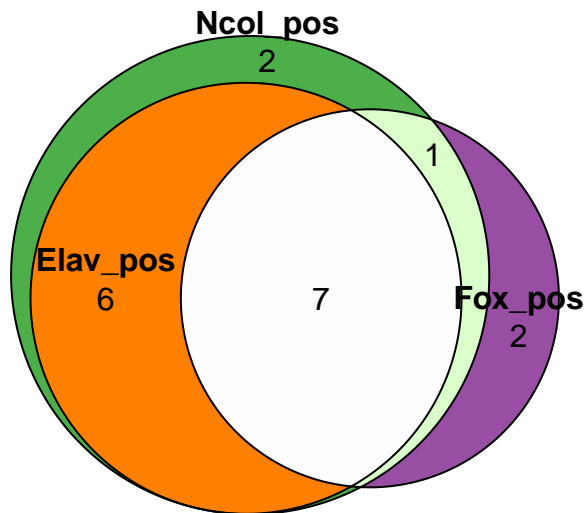
Nvec_vc1.1_XM_001639993.3

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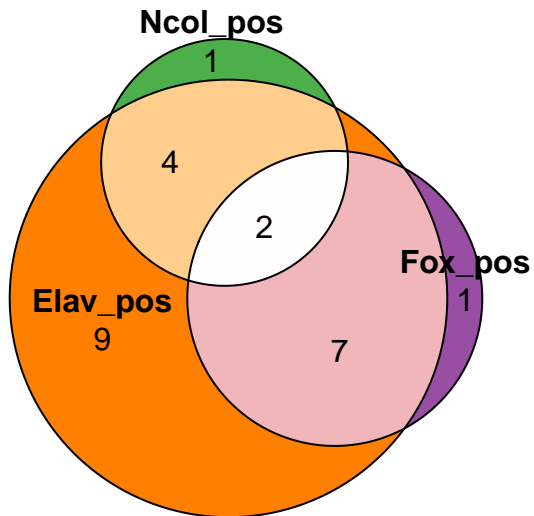
Nvec_vc1.1_XM_001637630.3

HCG37164; LysM/



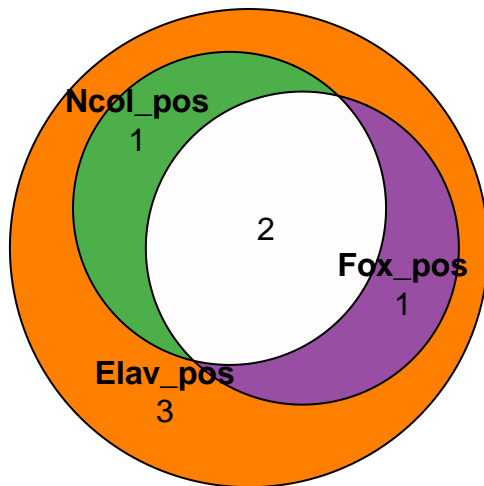
Nvec_vc1.1_XM_001633469.3

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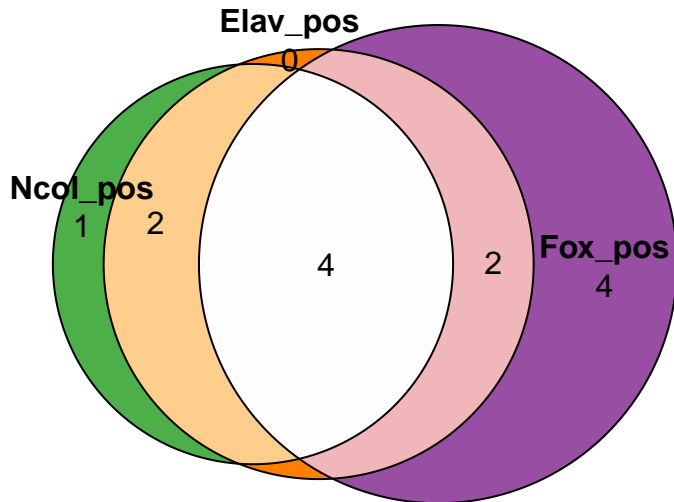
Nvec_vc1.1_XM_032377108.2

Short/branched chain spec; Acyl-CoA_dh_N/Acyl-CoA_dh



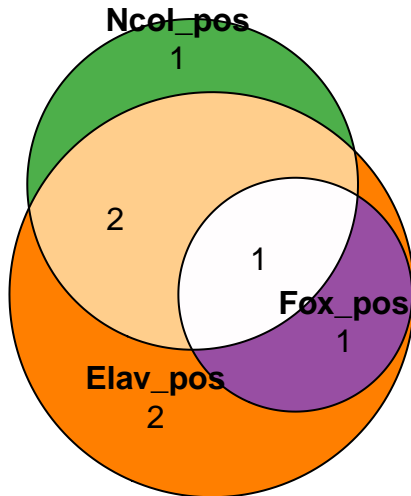
Nvec_vc1.1_XM_048722331.1

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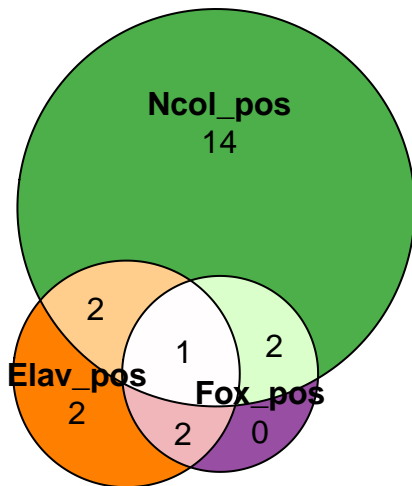
Nvec_vc1.1_XM_032372617.2

Tax1-binding protein 1; CALCOCO1/



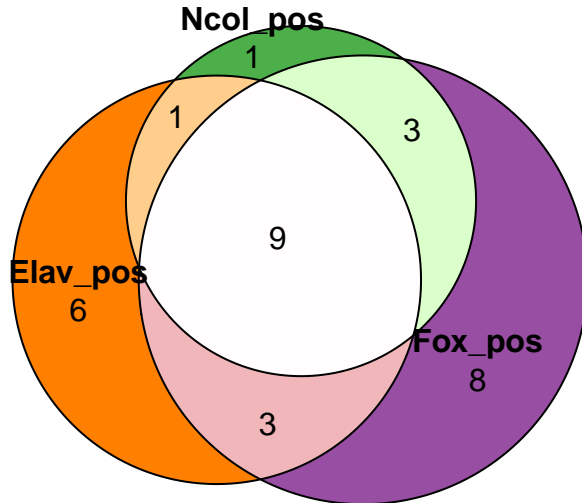
Nvec_vc1.1_XM_048726395.1

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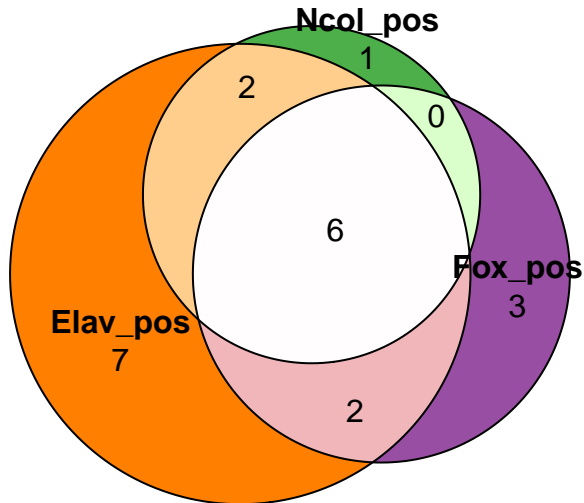
Nvec_vc1.1_XM_048731336.1

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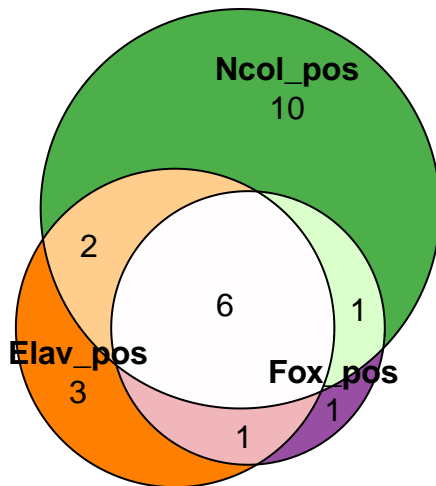


Nvec_vc1.1_XM_032363255.2

T-cell immunomodulatory p; VCBS/

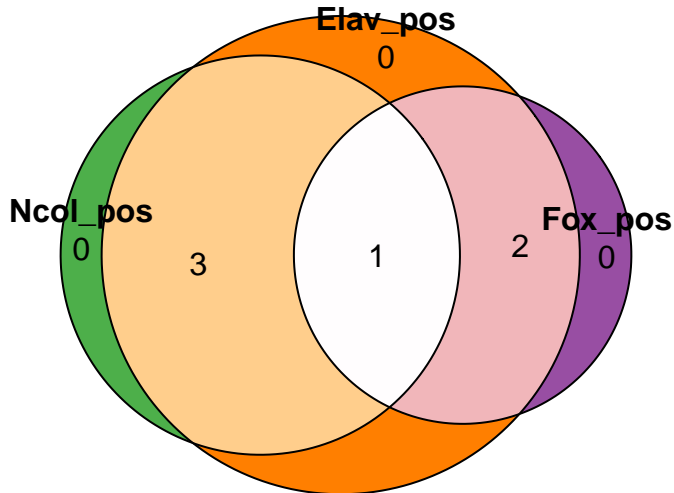


Nvec_vc1.1_XM_048731129.1
cDNA FLJ61192, highly sim; Band_7/



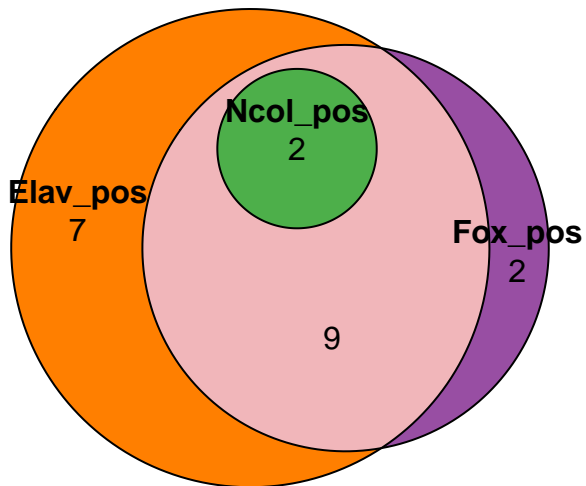
Nvec_vc1.1_XM_032386245.2

Pkinase:DSTYK; Pkinase.HG63.0:DSTYK



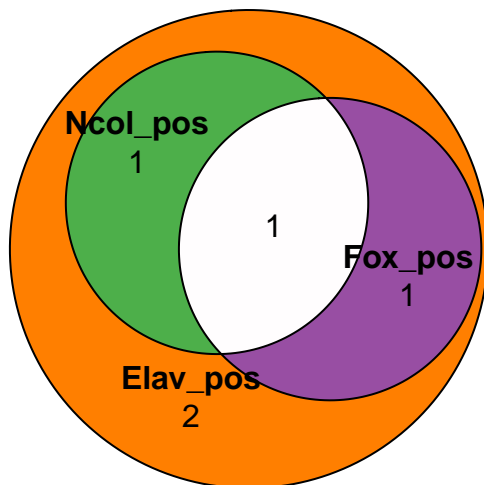
Nvec_vc1.1_XM_032374819.2

;



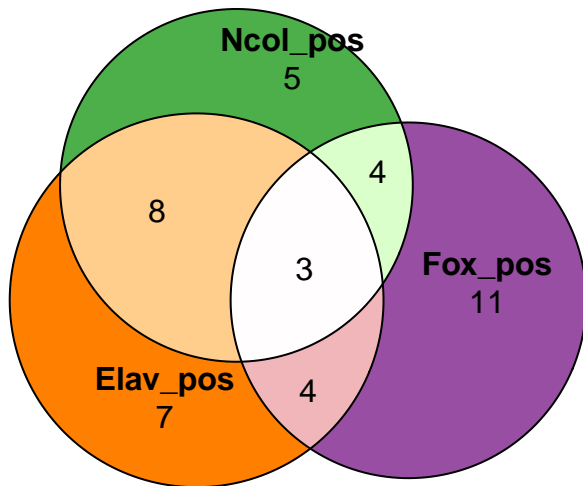
Nvec_vc1.1_XM_001640321.3

Chromo:CBX2/CBX4/CBX6/CBX; Chromo.HG1.28:CBX2/CBX4/C



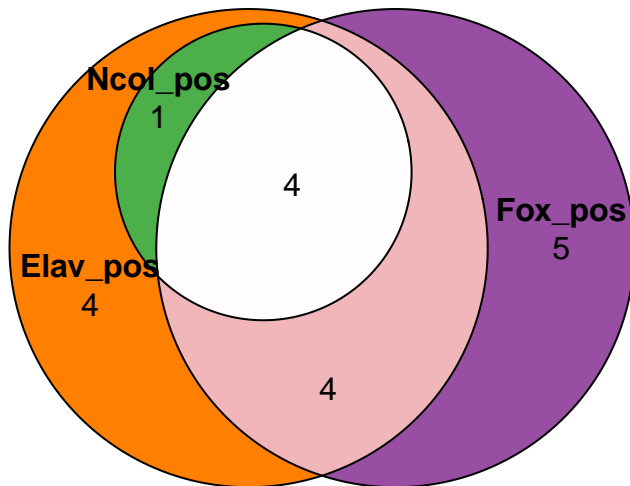
Nvec_v1g196131

Sodium-dependent neutral ; SNF/



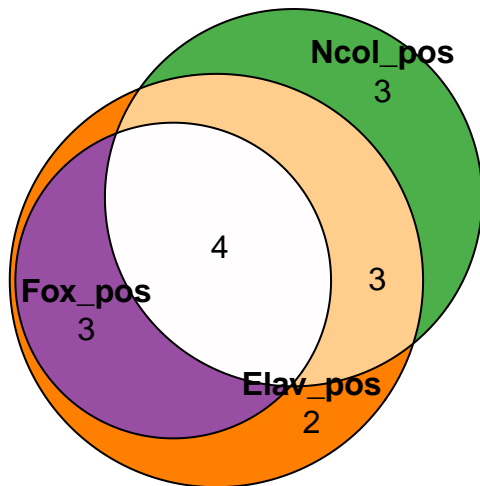
Nvec_NVE15532

Cytochrome c oxidase subu; COX6C/



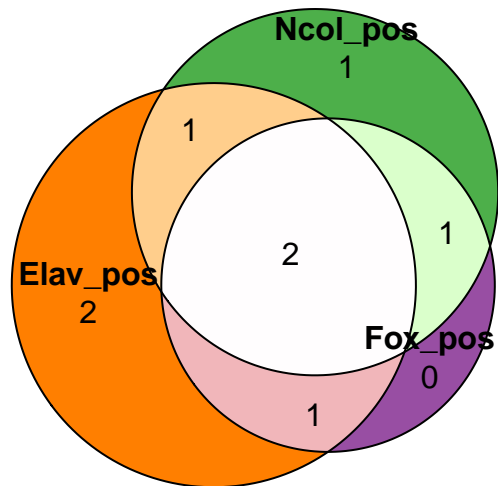
Nvec_vc1.1_XM_032363587.2

RalBP1-associated Eps dom; EF-hand_4/



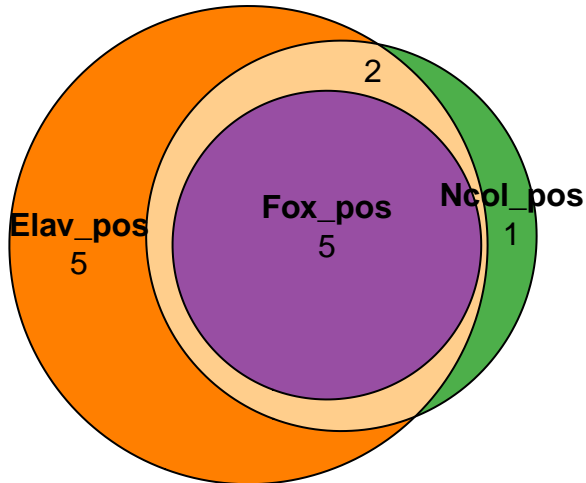
Nvec_vc1.1_XM_032385011.2

Protein FAM73B; DUF2217/



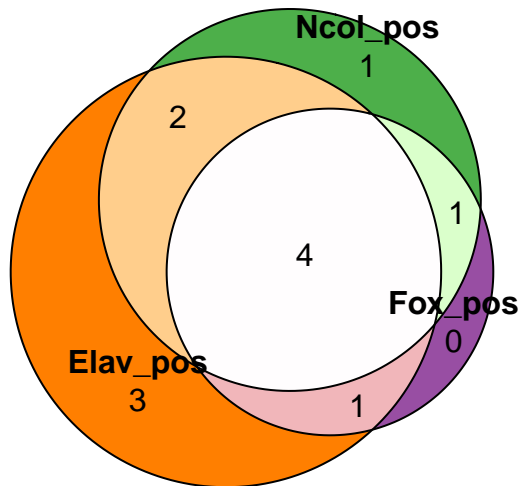
Nvec_vc1.1_XM_032372710.2

Syndecan:like:NRXN1/NRXN3; Syndecan/



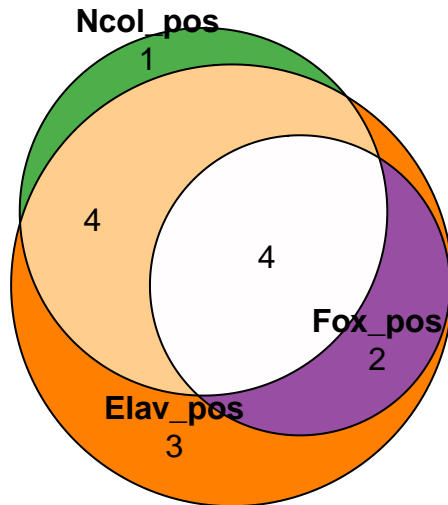
Nvec_vc1.1_XM_001624515.3

UPF0598 protein C8orf82; DUF4505/



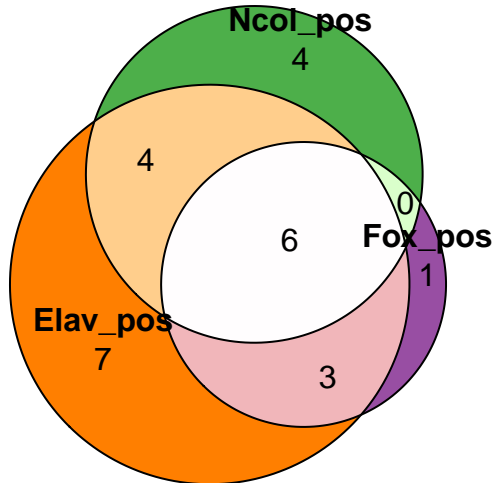
Nvec_vc1.1_XM_001629166.3

Graves disease carrier pr; Mito_carr/Mito_carr/Mito_

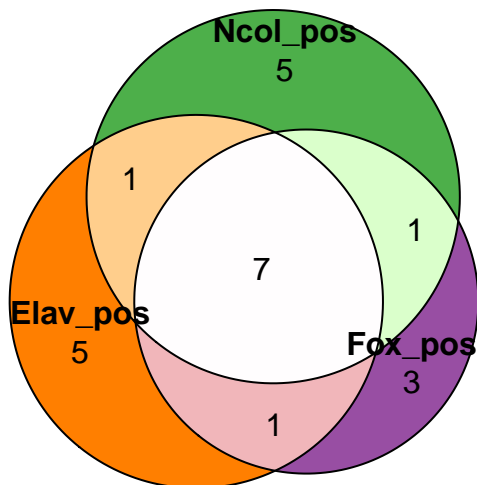


Nvec_vc1.1_XM_001641181.3

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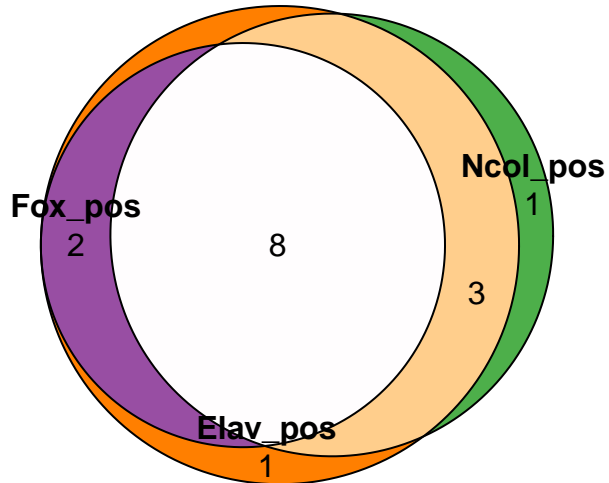


Nvec_vc1.1_XM_001639086.3
cDNA FLJ61388, highly sim; TPR_11/



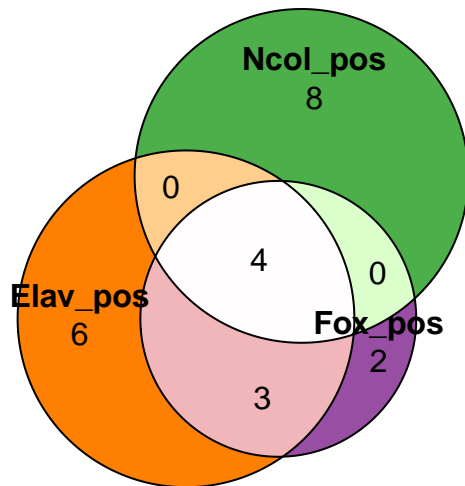
Nvec_vc1.1_XM_001637768.3

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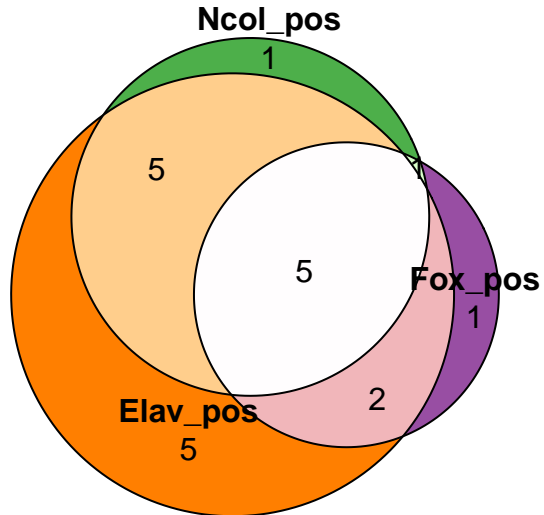
Nvec_vc1.1_XM_048727015.1

DOCK7 protein; Ded_cyto/



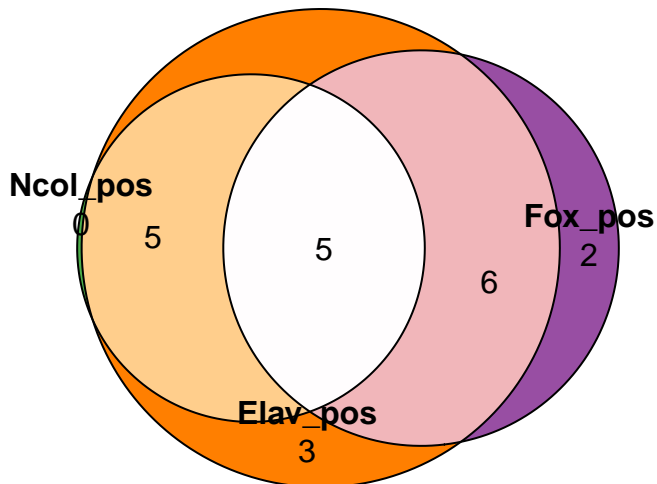
Nvec_vc1.1_XM_032375591.2

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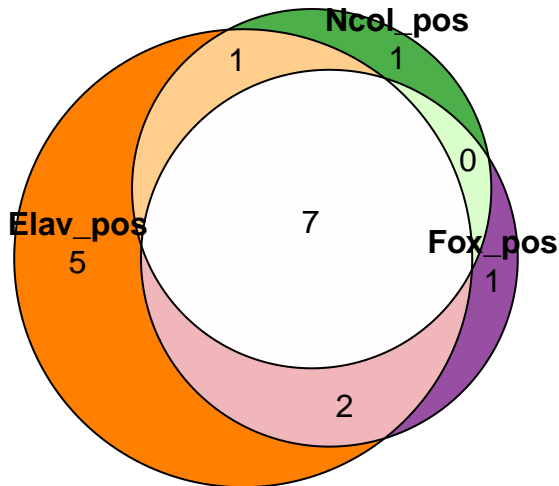
Nvec_vc1.1_XM_001634737.3

ion-NaKCa:like:KCNB1/KCNB; BTB_2/Ion_trans/



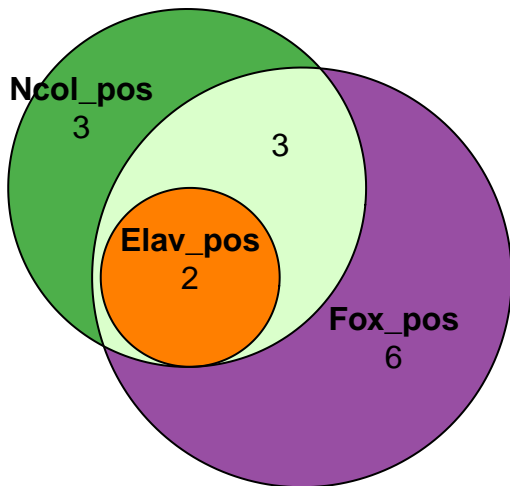
Nvec_vc1.1_XM_048734049.1

Vesicle docking protein p; Uso1_p115_head/



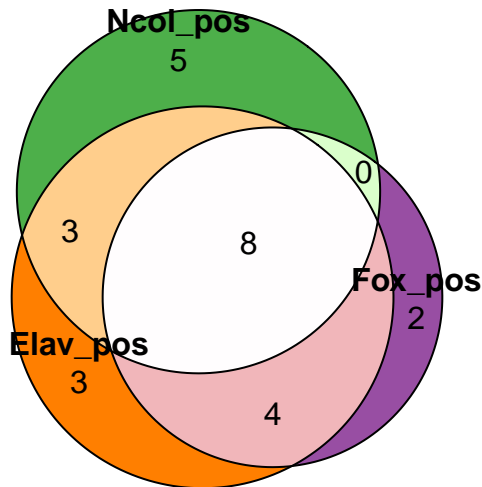
Nvec_vc1.1_XM_032386956.2

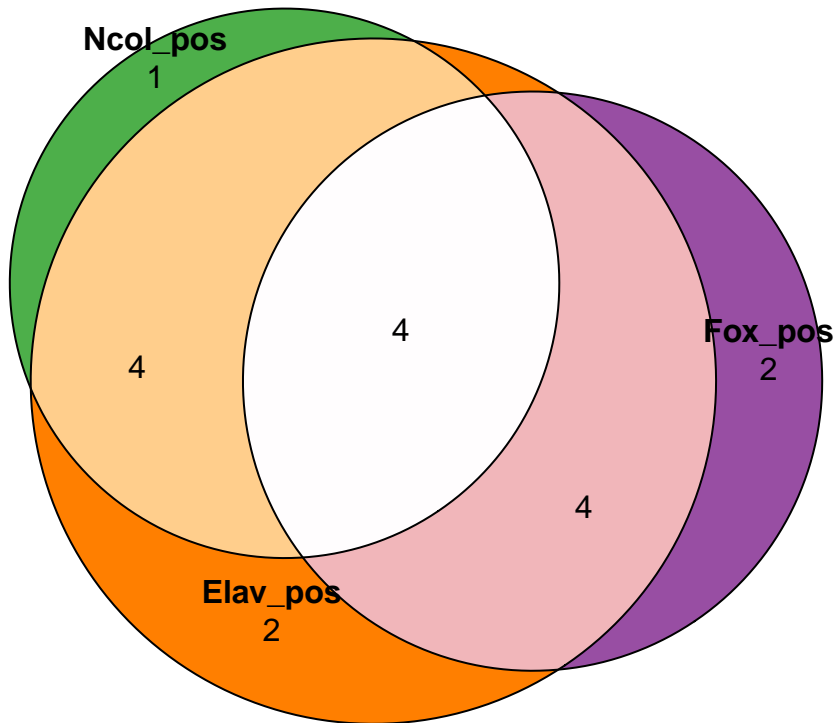
Tetraspanin-33; Tetraspannin/



Nvec_vc1.1_XM_001628870.3

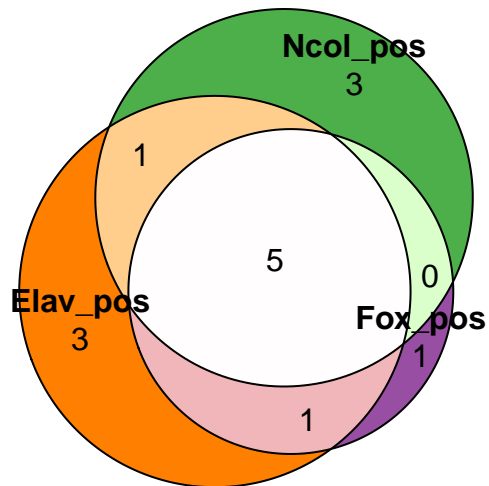
Tripartite motif-containing; zf-C3HC4_2/zf-B_box/zf-B_



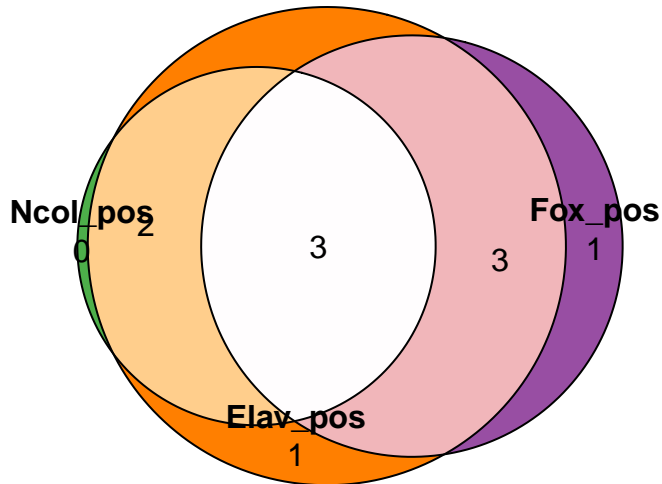


Nvec_vc1.1_XM_048731183.1

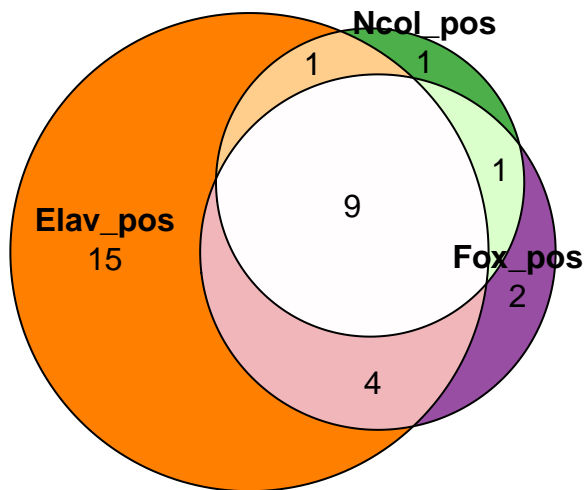
;



Nvec_vc1.1_XM_001636230.3
; Death/Death/

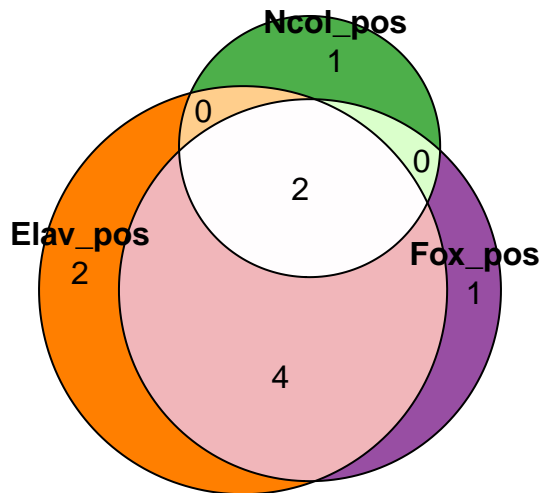


Nvec_vc1.1_XM_032385897.2
; zf-MYND/

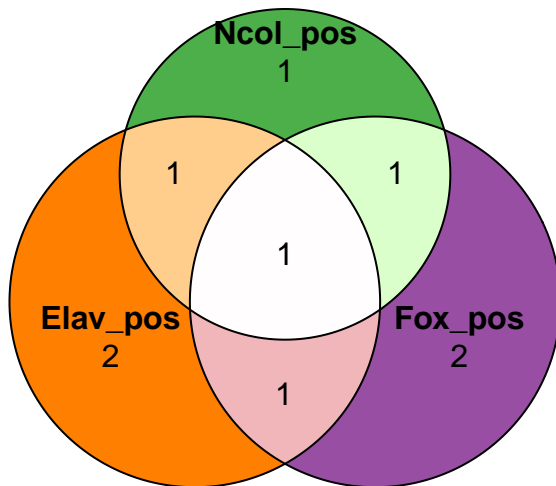


Nvec_vc1.1_XM_032384325.2

Synaptotagmin:RIMS1/RIMS2; Synaptotagmin.HG1.33:RIMS

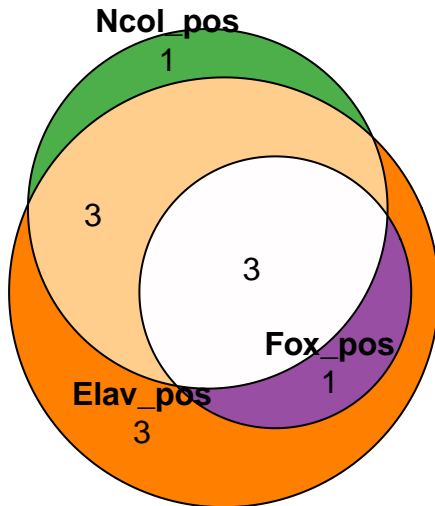


Nvec_vc1.1_XM_032362659.2
; SRP-alpha_N/



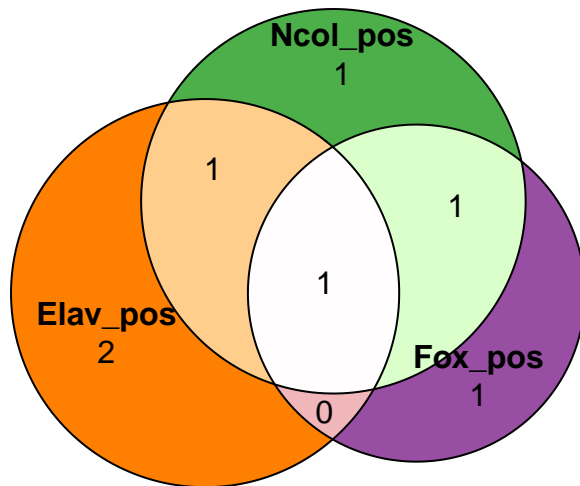
Nvec_vc1.1_XM_001641251.3

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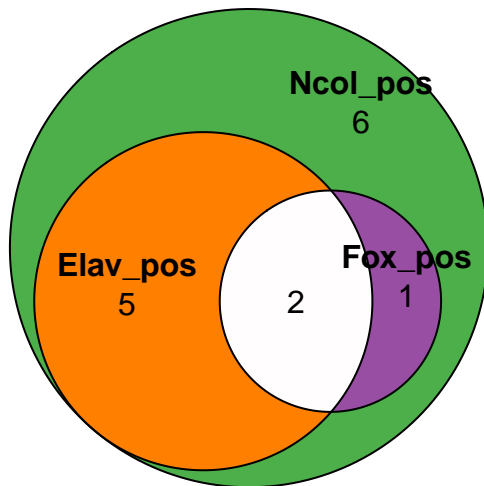
Nvec_vc1.1_XM_001632360.3

Receptor expression-enhanc



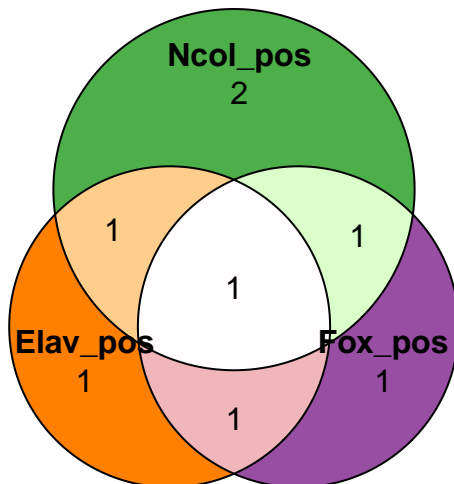
Nvec_vc1.1_XM_032375518.2

PMCA:ATP2C1/ATP2C2; PMCA.HG2.2:ATP2C1/ATP2C2

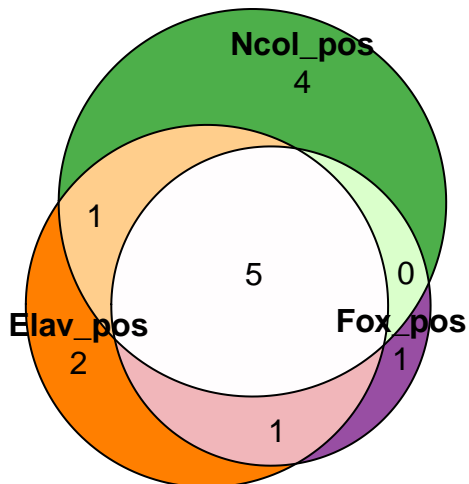


Nvec_vc1.1_XM_001633809.3

Pkinase:TNK1/TNK2; Pkinase.HG78.0:TNK1/TNK2

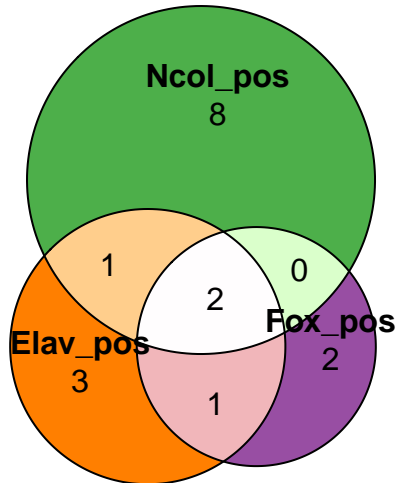


Nvec_vc1.1_XM_001633805.3
cDNA FLJ37604 fis, clone ; Frag1/



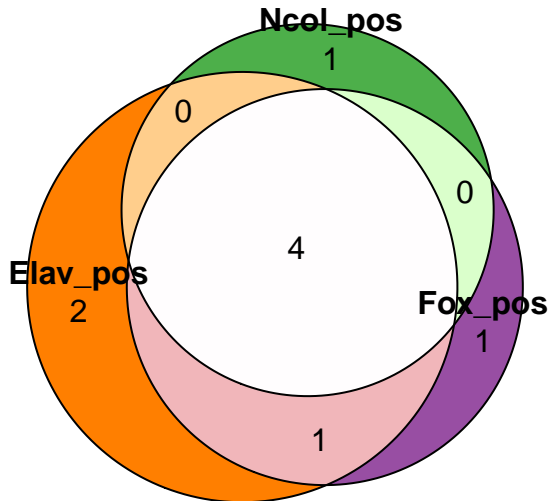
Nvec_vc1.1_XM_001632635.3

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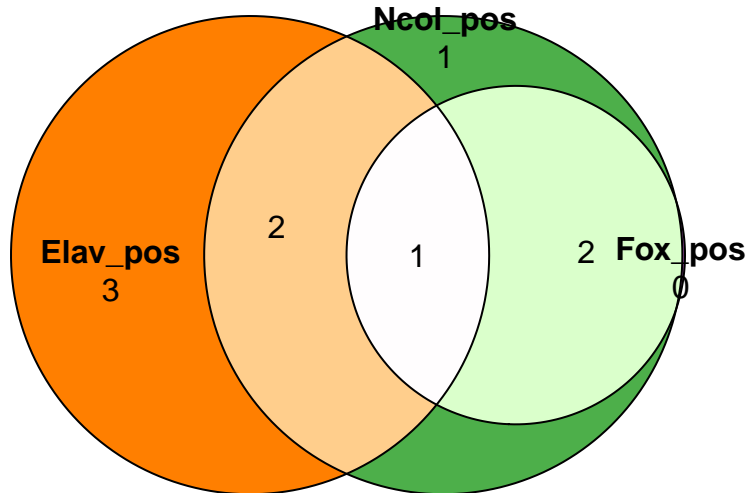


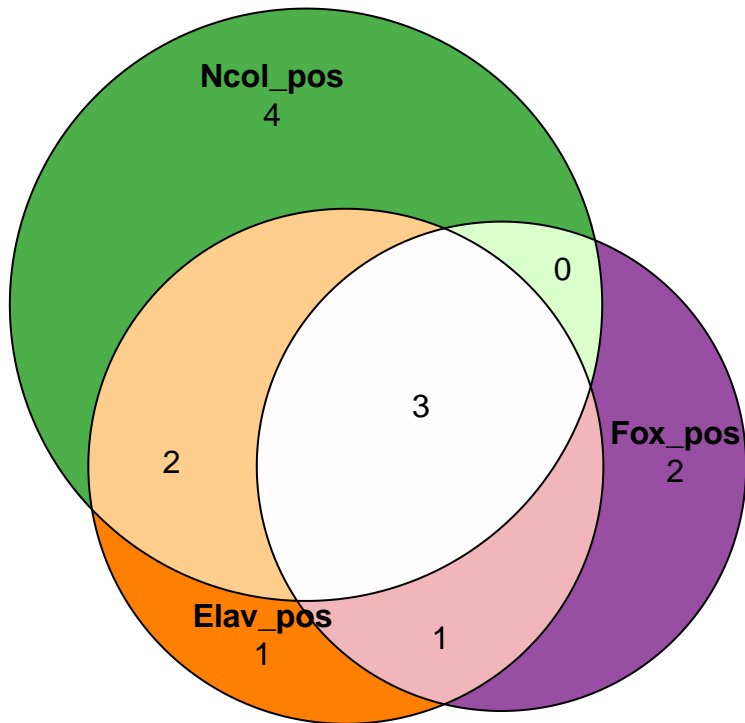
Nvec_vc1.1_XM_001627934.3

Mediator complex subunit ; Med25_VWA/Med25/Med25_NR-

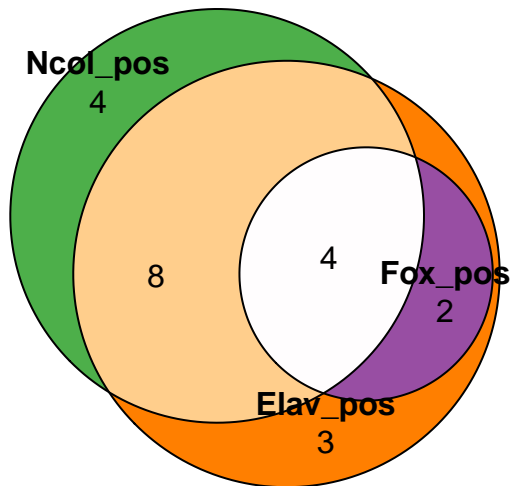


Nvec_vc1.1_XM_048733597.1
; DUF640/



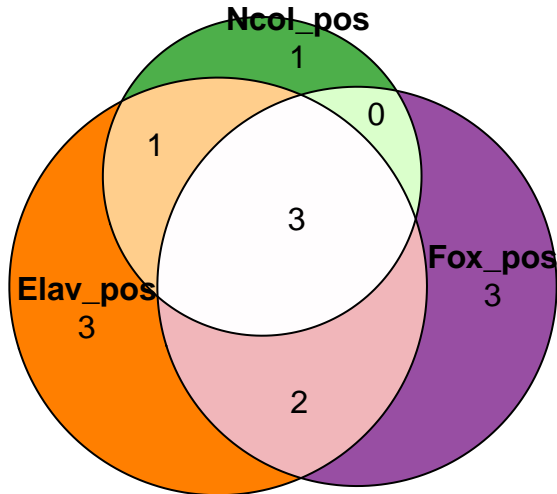


Nvec_vc1.1_XM_048733384.1
cDNA, FLJ93684, highly si; Mab-21/



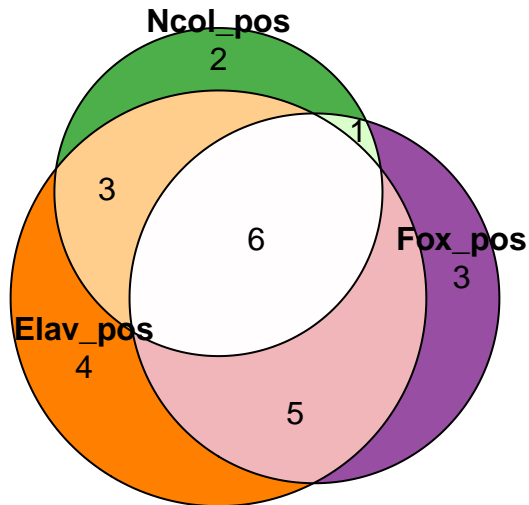
Nvec_vc1.1_XM_001635751.3

Synaptobrevin-like 1; Longin/Synaptobrevin/



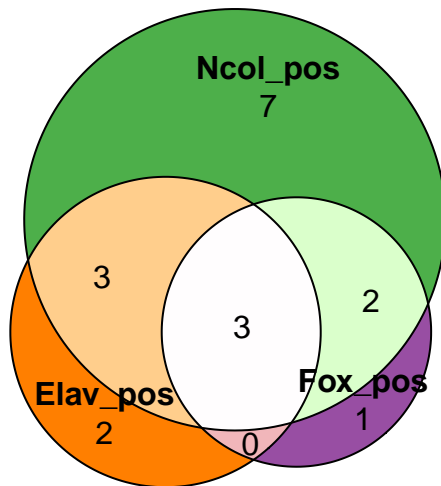
Nvec_vc1.1_XM_001636259.3

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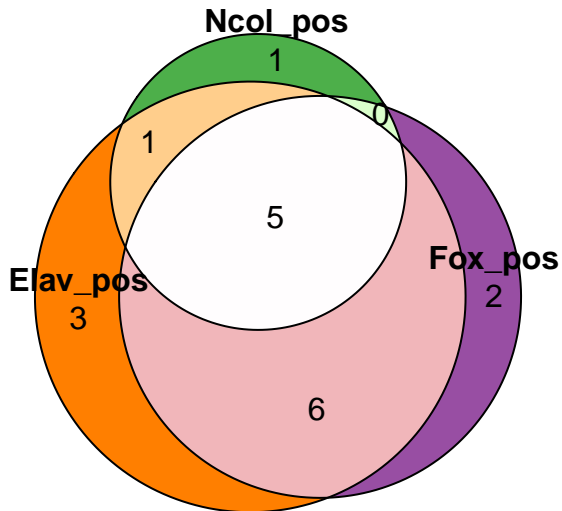
Nvec_vc1.1_XM_048721067.1

Vesicle-associated membra; Motile_Sperm/



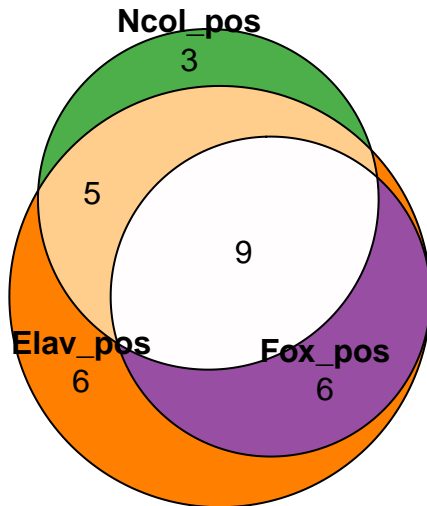
Nvec_vc1.1_XM_048734307.1

Phosphatidate phosphatase; Lipin_N/



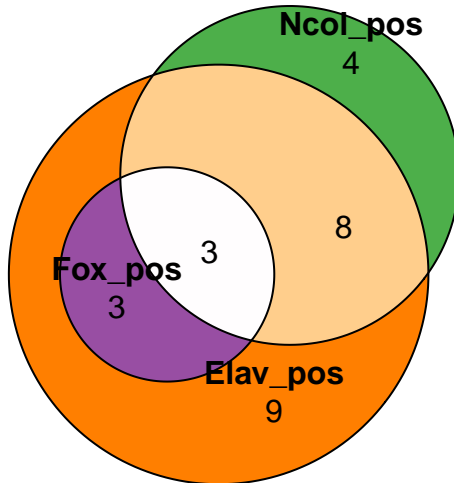
Nvec_vc1.1_XM_032386897.2

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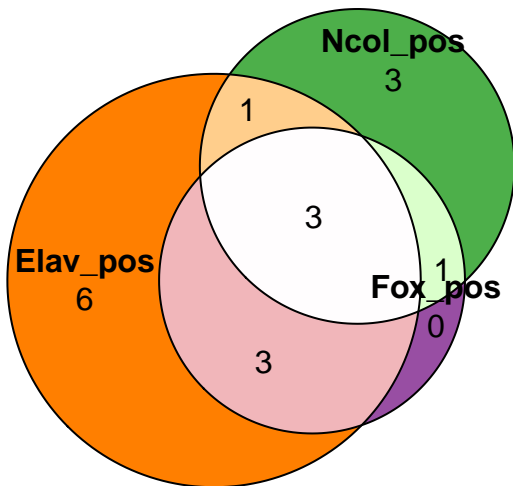
Nvec_vc1.1_XM_048721230.1

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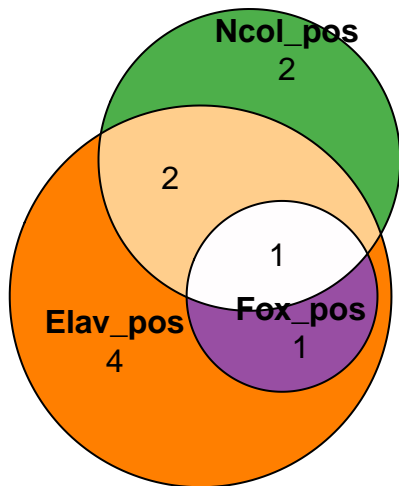
Nvec_vc1.1_XM_032373632.2

Putative uncharacterized ; TPR_11/TPR_11/TPR_1/

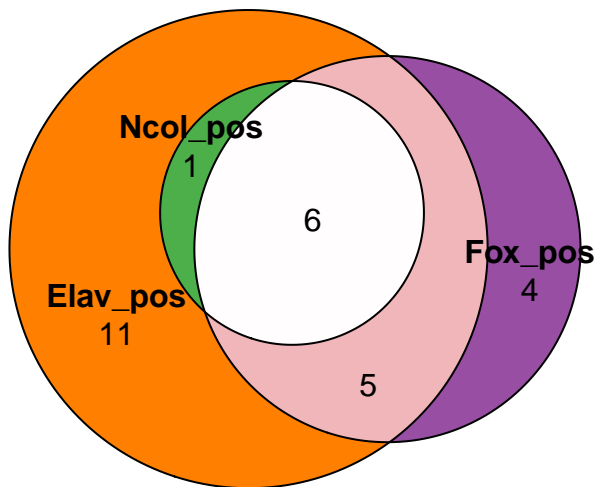


Nvec_vc1.1_XM_001629070.3

N-acetylglucosamine-1-pho; DUF2233/

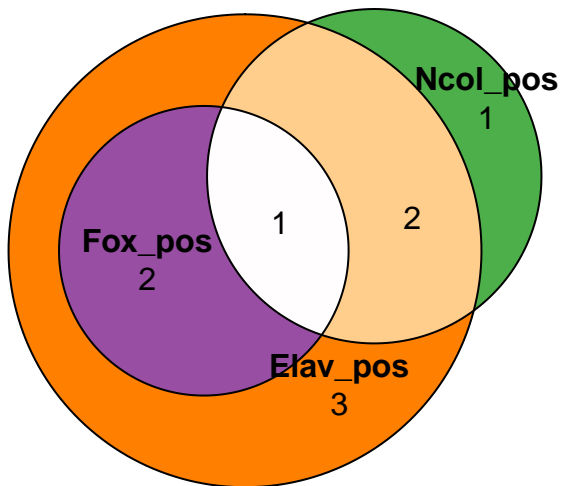


Nvec_vc1.1_XM_032371145.2
NEDD4 family-interacting ; DUF2370/



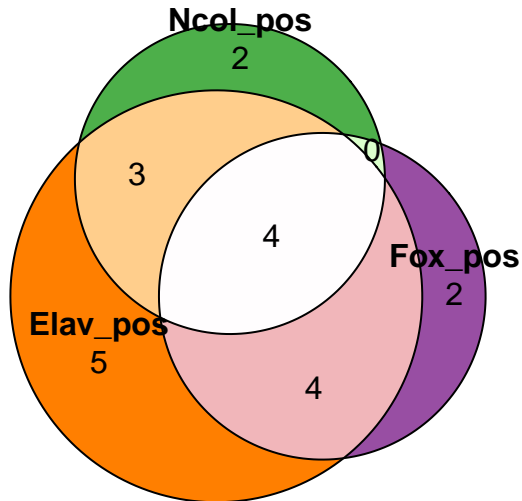
Nvec_NVE25390

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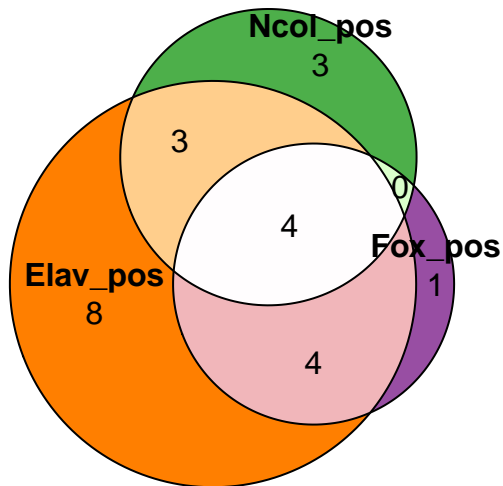


Nvec_vc1.1_XM_032375428.2

Glutamate receptor ionotr; CaM_bdg_C0/

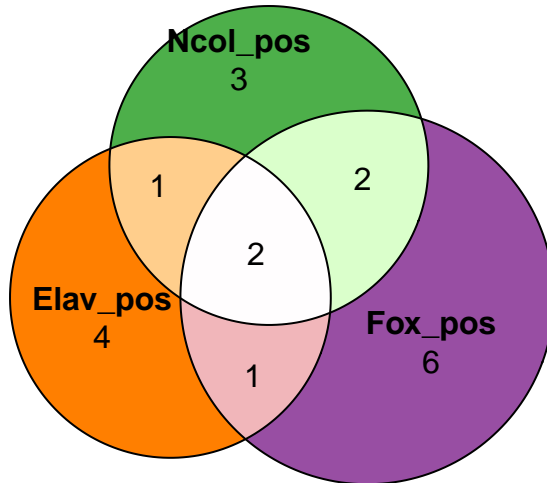


Nvec_vc1.1_XM_001629013.3
Y_phosphatase:like:DUSP1/; DSPc/



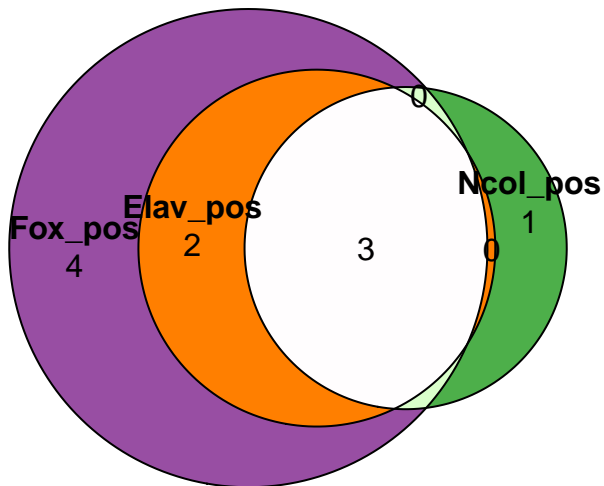
Nvec_vc1.1_XM_048726200.1

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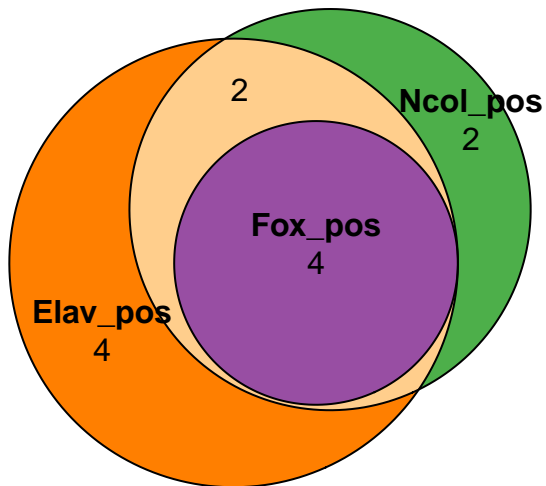
Nvec_vc1.1_XM_048732973.1

Phospholipid-transporting;



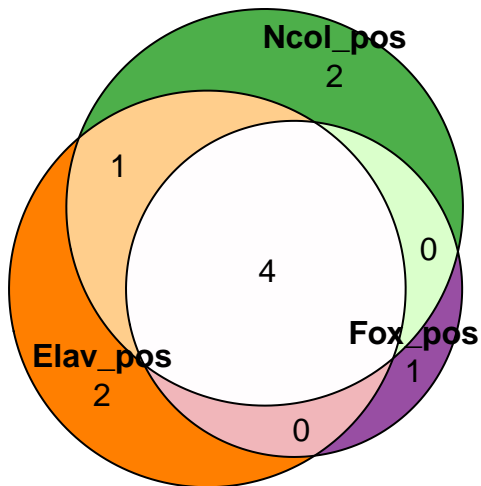
Nvec_vc1.1_XM_032363357.2

RRM:A1CF/RBM46/RBM47; RRM.HG11.4:A1CF/RBM46/RBM



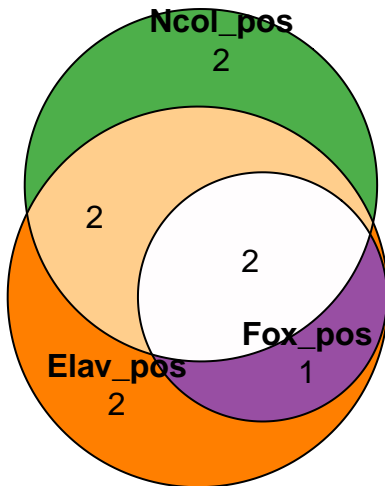
Nvec_vc1.1_XM_048731571.1

Pkinase:PINK1; Pkinase.HG67.1:PINK1

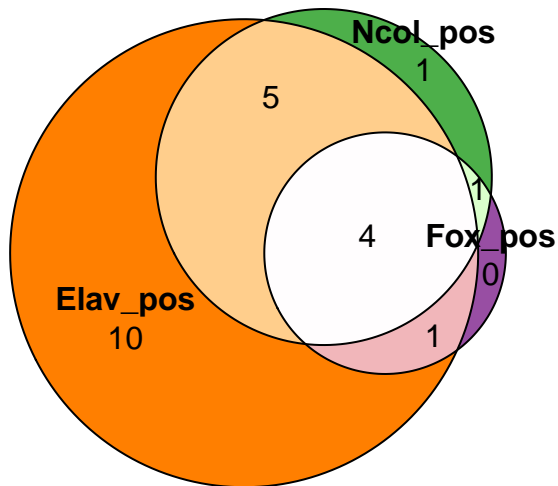


Nvec_vc1.1_XM_048730105.1

TRAF-type zinc finger dom;

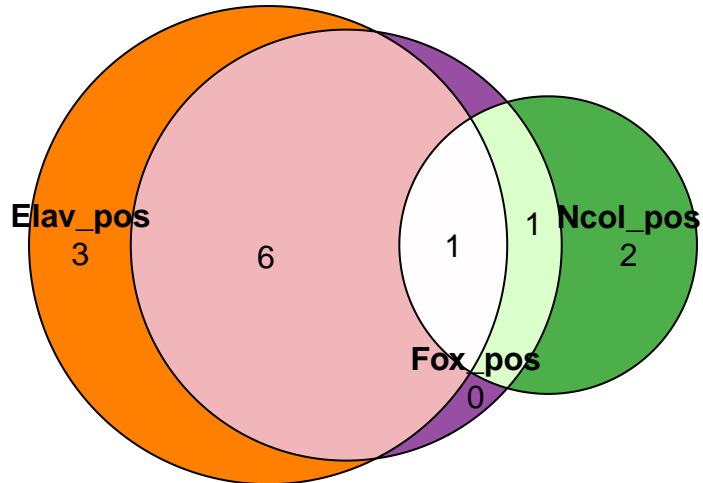


Nvec_vc1.1_XM_032384320.2
cDNA FLJ52785, highly sim;



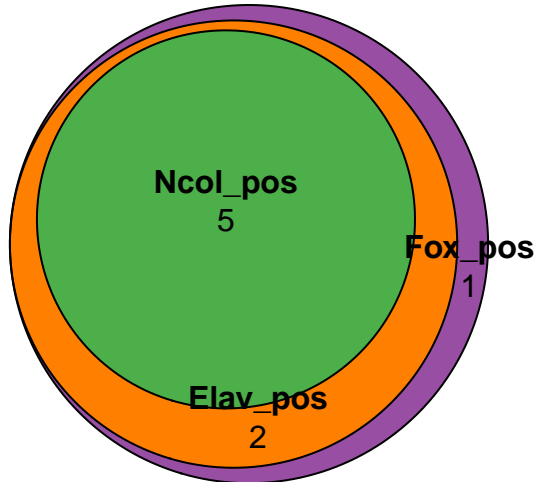
Nvec_vc1.1_XM_032366272.2

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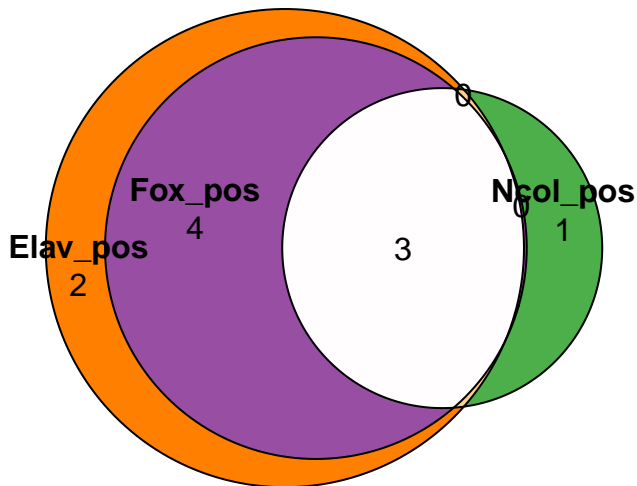
Nvec_vc1.1_XM_032378434.2

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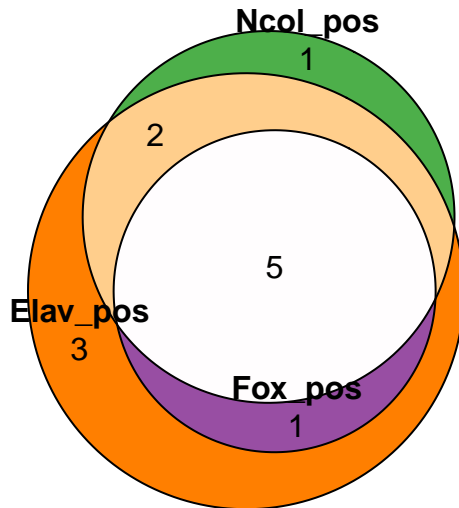
Nvec_vc1.1_XM_048726086.1

Pkinase:MAPKAPK5; Pkinase.HG178.0:MAPKAPK5

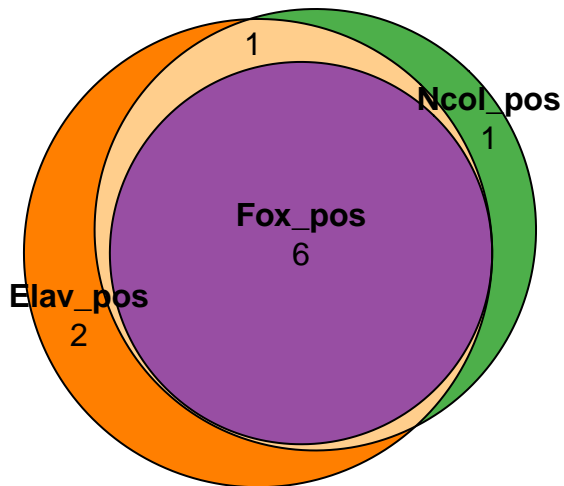


Nvec_vc1.1_XM_032380416.2

DDB1– and CUL4–associated; WD40/

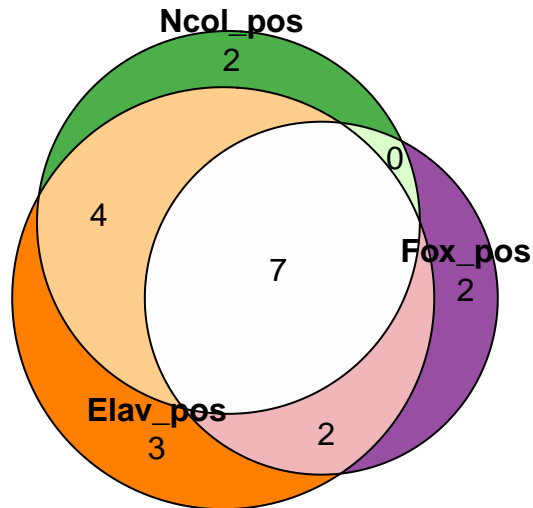


Nvec_vc1.1_XM_032364659.2
cDNA FLJ55491; DUF2045/



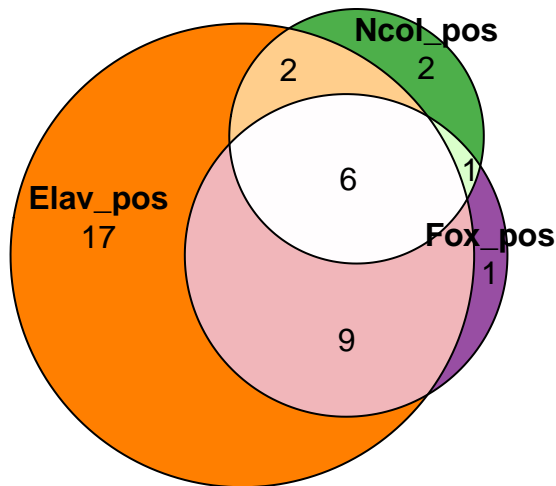
Nvec_vc1.1_XM_001621986.3

WD and tetratricopeptide ; WD40/WD40/WD40/TPR_11/WD4



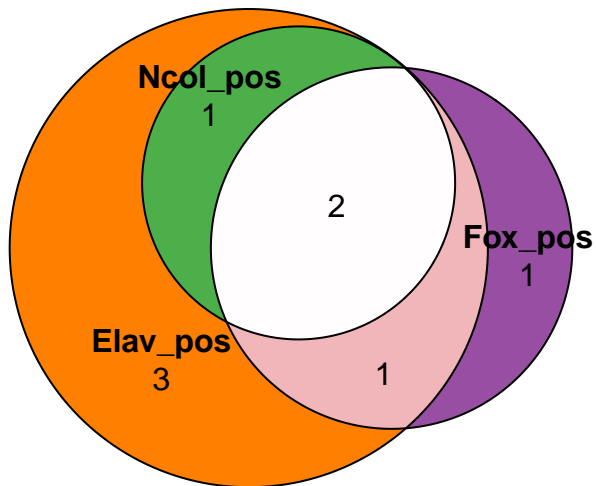
Nvec_vc1.1_XM_001633176.3

Protein kinase, cAMP-depe; RIIa/cNMP_binding/cNMP_bi



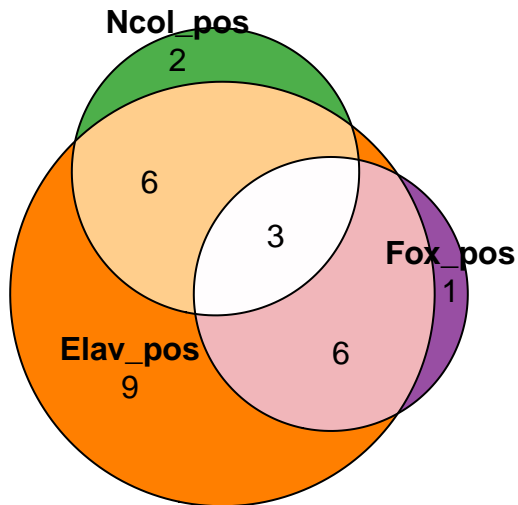
Nvec_vc1.1_XM_001624272.3

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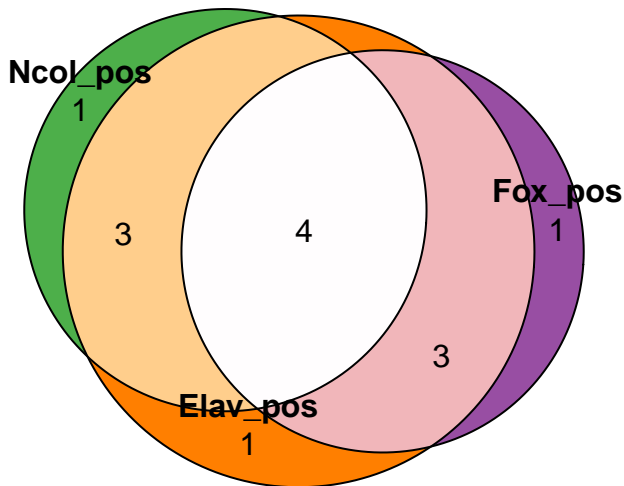
Nvec_v1g241236

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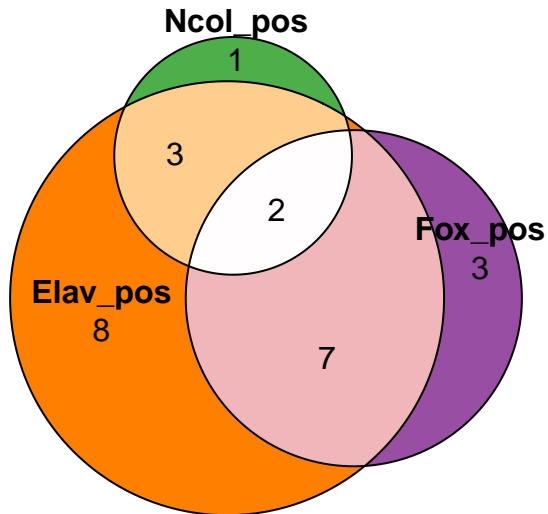
Nvec_vc1.1_XM_032362879.2

F-box/LRR-repeat protein ; Hemerythrin/F-box-like/LR



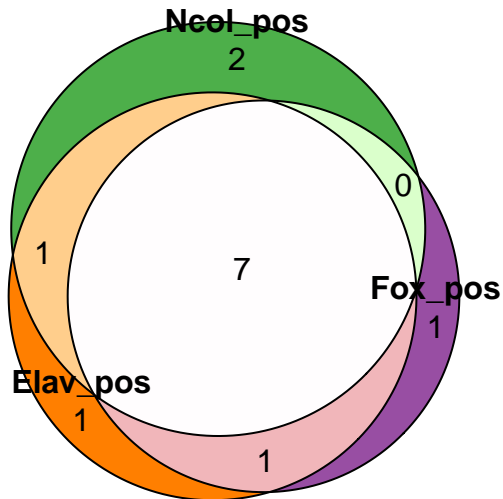
Nvec_vc1.1_XM_032374324.2

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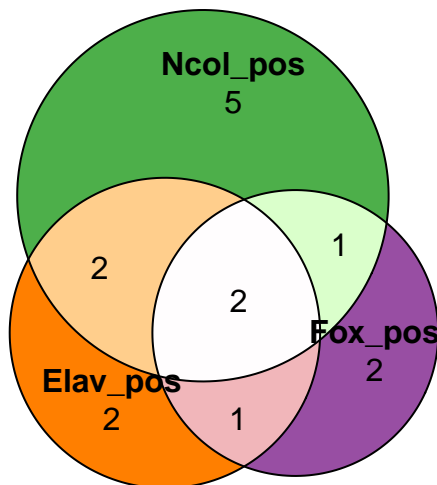


Nvec_vc1.1_XM_032382893.2

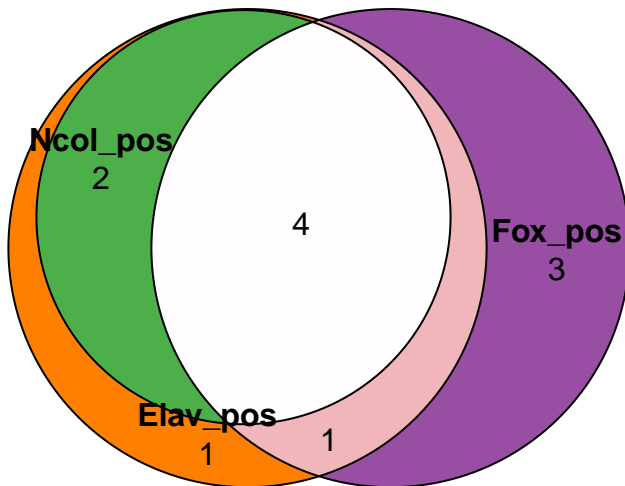
Pkinase:CSNK2A1/CSNK2A2/C; Pkinase.HG104.1:CSNK2A1/C



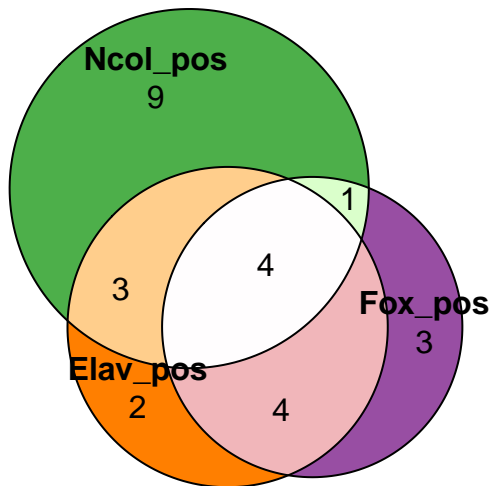
Nvec_vc1.1_XM_032372600.2
RING finger protein 32; zf-RING_2/



Nvec_vc1.1_XM_001638643.3
Cytochrome c oxidase VIIc; COX7C/

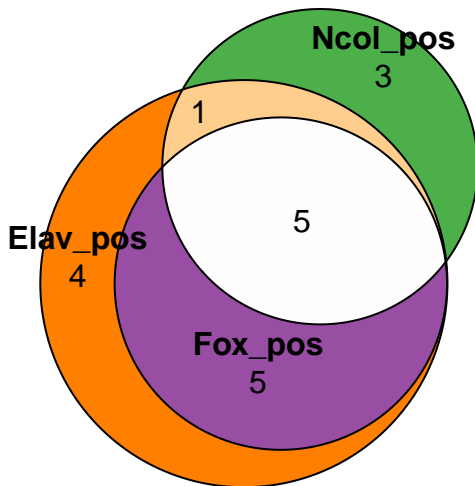


Nvec_vc1.1_XM_032371787.2
Putative uncharacterized ; AAA_2/

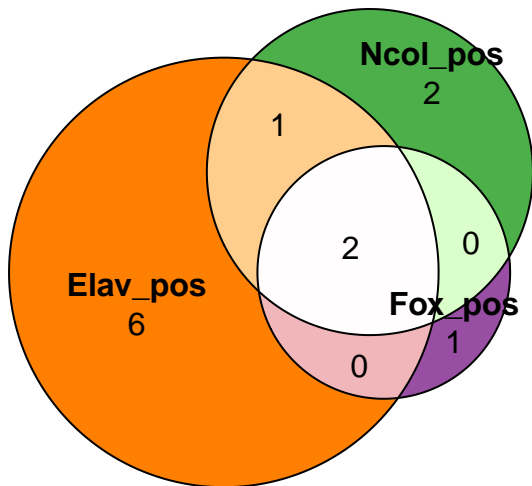


Nvec_vc1.1_XM_001637304.3

Amyloid-like protein 1; APP_N/APP_Cu_bd/APP_E2/AP

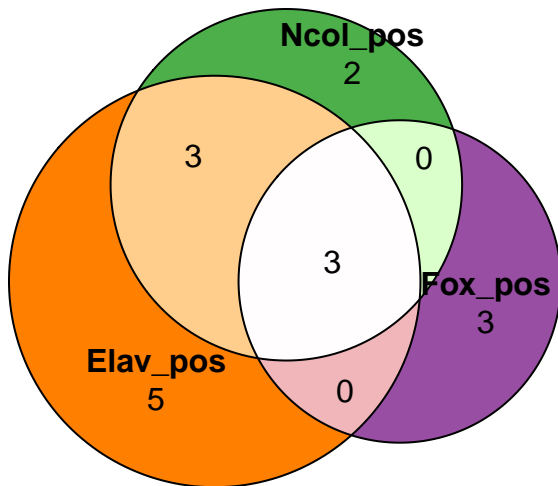


Nvec_vc1.1_XM_001629883.3
Membrane-associated ring ; RINGv/



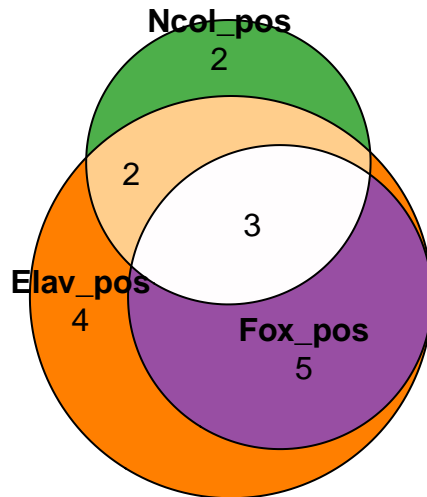
Nvec_vc1.1_XM_001641834.3

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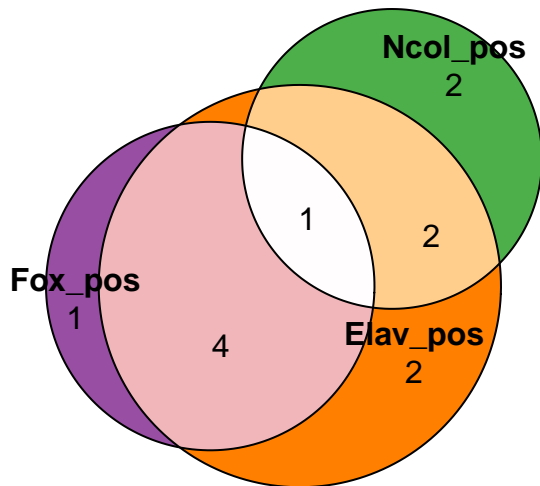
Nvec_vc1.1_XM_032372158.2

Protein FAM102B; NT-C2/



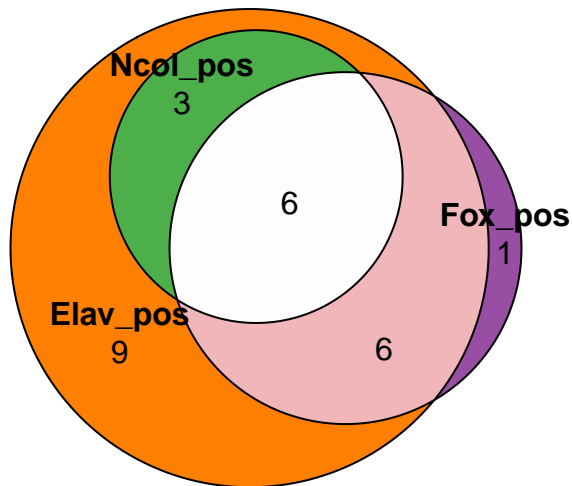
Nvec_vc1.1_XM_048731911.1

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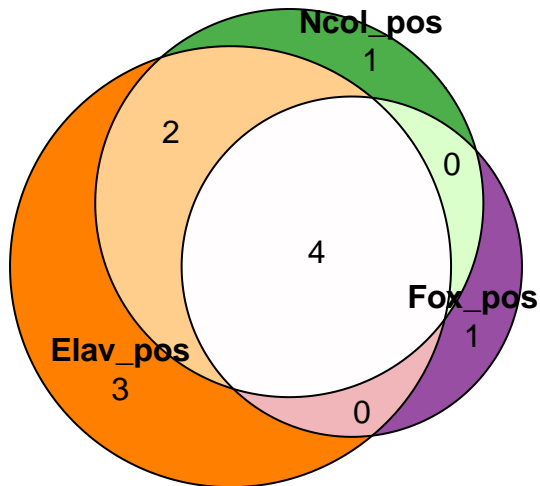
Nvec_vc1.1_XM_032374809.2

Sugar_tr:NA; MFS_1/



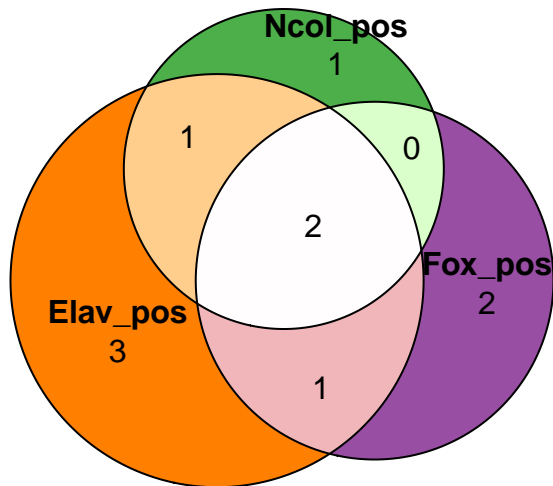
Nvec_vc1.1_XM_032382247.2

Protein ripply2; Ripply/



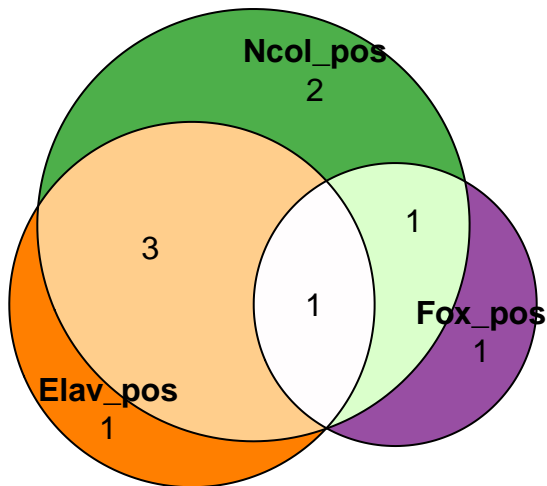
Nvec_vc1.1_XM_032384537.2

Chromosome 1 open reading; SprT-like/



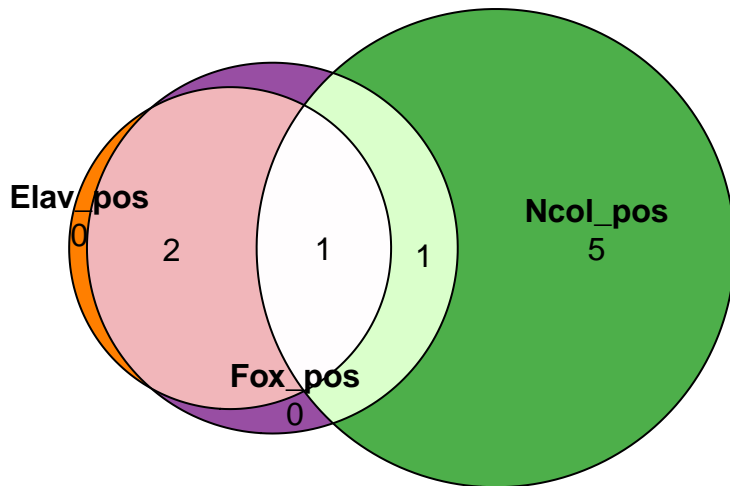
Nvec_vc1.1_XM_032383201.2

AN1-type zinc finger prot; zf-AN1/zf-AN1/



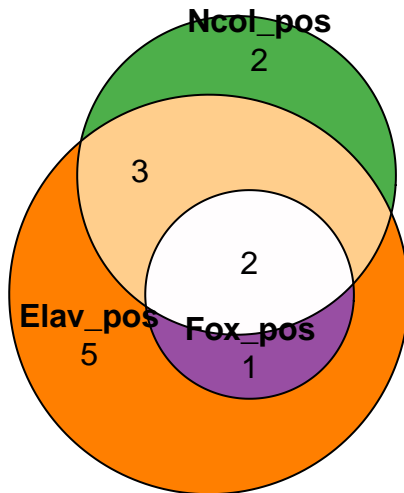
Nvec_vc1.1_XM_001621929.3

Long-chain-fatty-acid--Co; AMP-binding/



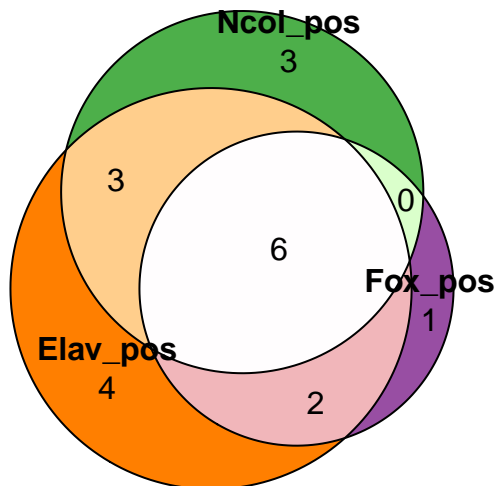
Nvec_vc1.1_XM_048730775.1

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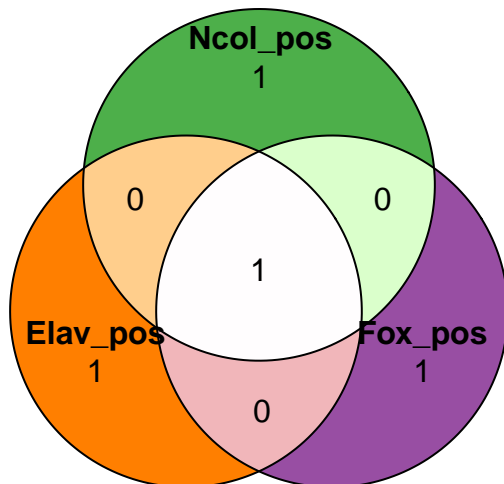
Nvec_vc1.1_XM_032376354.2

RasGAP:RASA1; RasGAP.HG1.5:RASA1

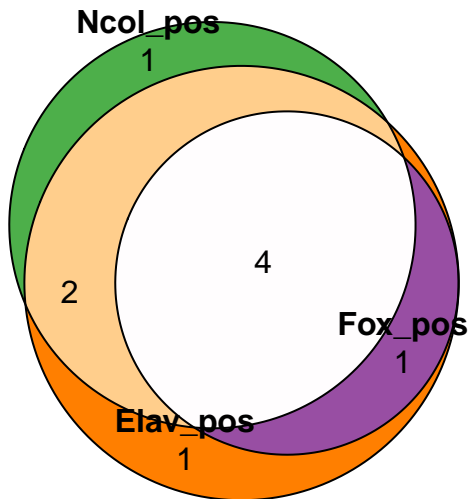


Nvec_vc1.1_XM_001626510.3

Caspase 8 isoform C; Peptidase_C14/

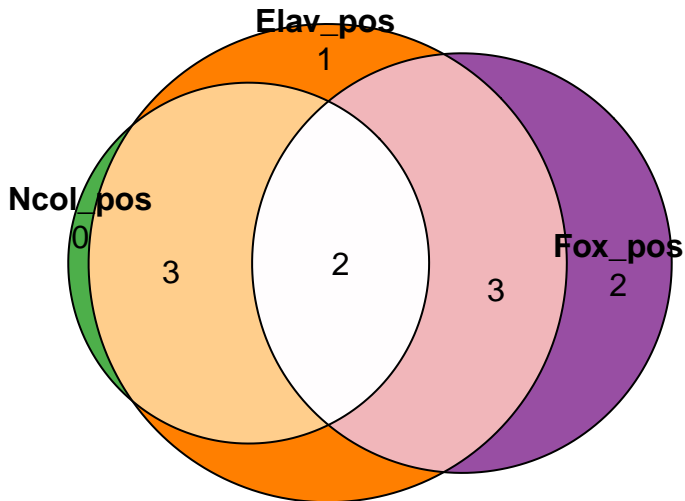


Nvec_vc1.1_XM_032386887.2
cDNA FLJ61530, highly sim; VHS/FYVE/



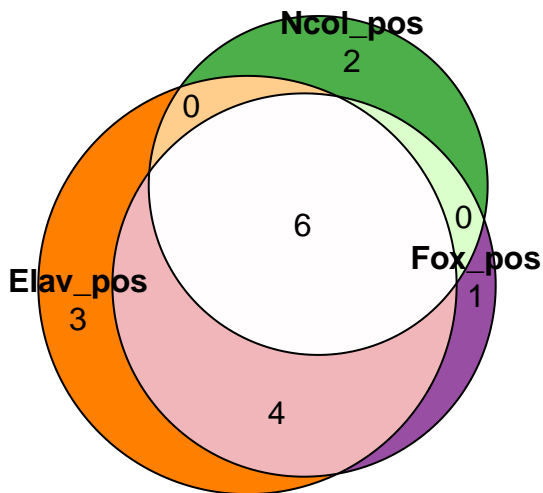
Nvec_vc1.1_XM_048734130.1

Monocarboxylate transport;



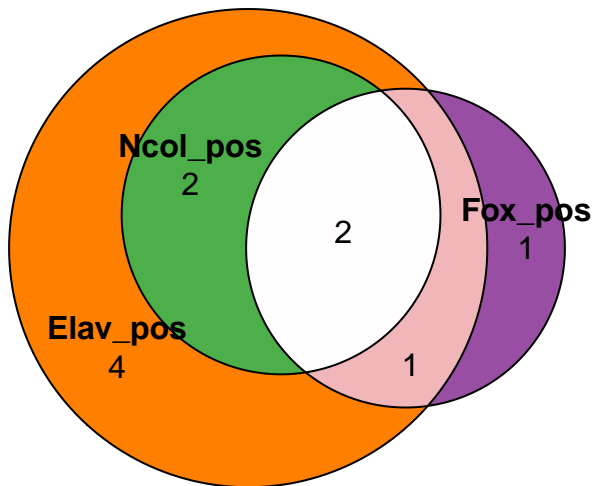
Nvec_vc1.1_XM_032379588.2

RRM:NONO/PSPC1/SFPQ; RRM.HG29.2:NONO/PSPC1/SFP



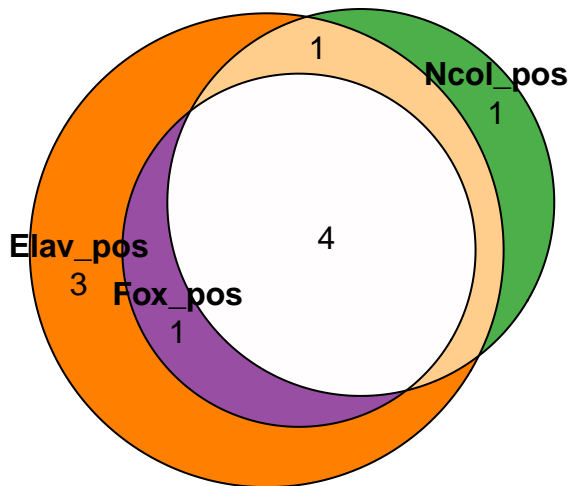
Nvec_vc1.1_XM_048721290.1

Dihydropyrimidine dehydro; GATase_2/Glu_syn_central/

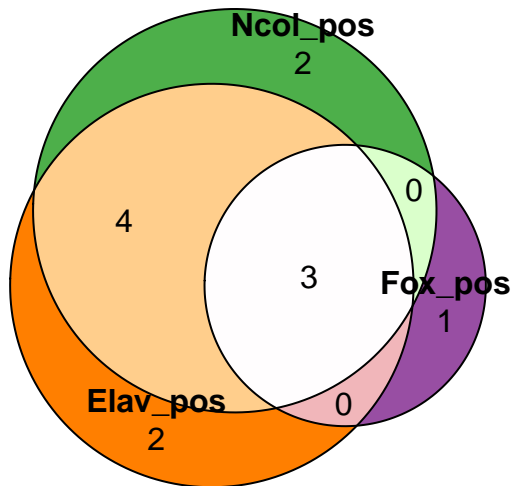


Nvec_vc1.1_XM_048723209.1

cDNA FLJ53444, highly sim; BRO1/ALIX_LYPXL_bnd/ALIX_

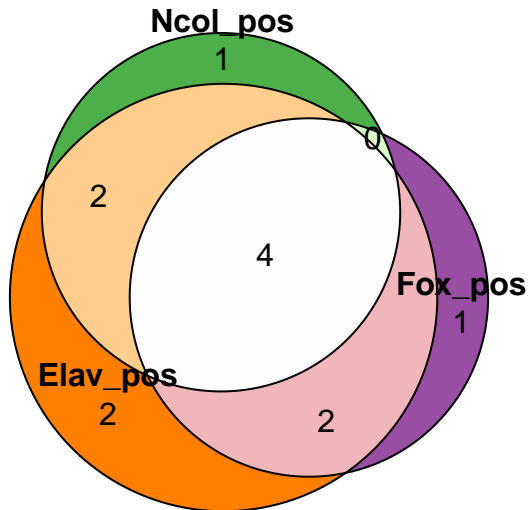


Nvec_vc1.1_XM_048722635.1
zf-C2H2.HG6.0:like:AL0316;



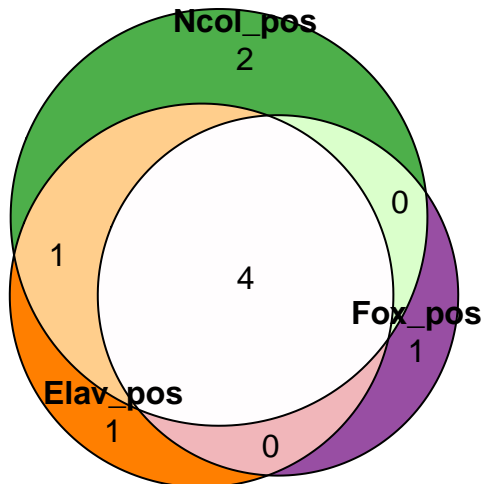
Nvec_vc1.1_XM_001635503.3

Membrane magnesium transp; MMgT/



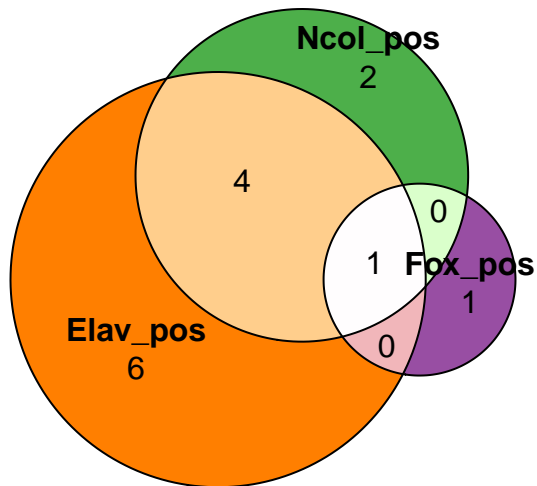
Nvec_vc1.1_XM_032383649.2

E3 ubiquitin-protein liga; zf-C3HC4_3/



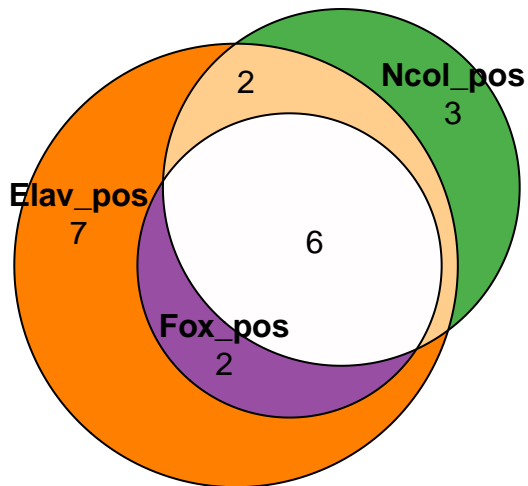
Nvec_vc1.1_XM_032365442.2

Epididymis secretory sper; AhpC-TSA/1-cysPrx_C/



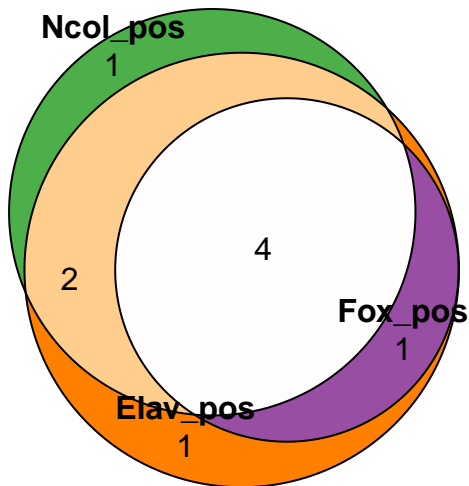
Nvec_vc1.1_XM_032377918.2

Pkinase:MAP3K12/MAP3K13; Pkinase.HG42.4:MAP3K12/MA



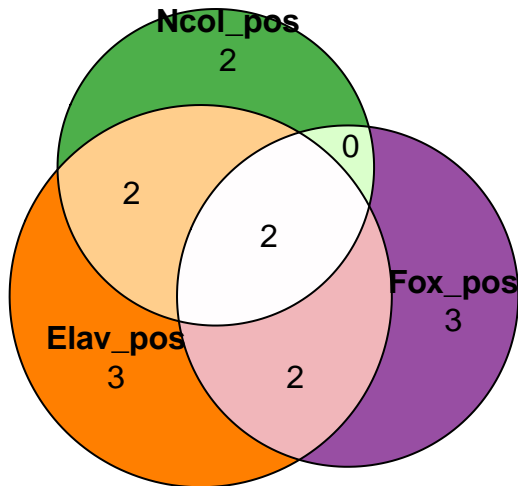
Nvec_vc1.1_XM_001627411.3

Cytochrome c oxidase asse; Pet191_N/

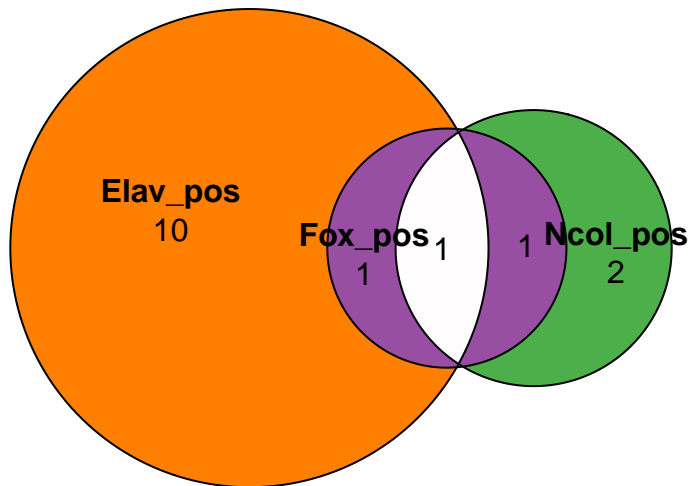


Nvec_vc1.1_XM_048725694.1

WD repeat-containing prot; WD40/WD40/WD40/WD40/

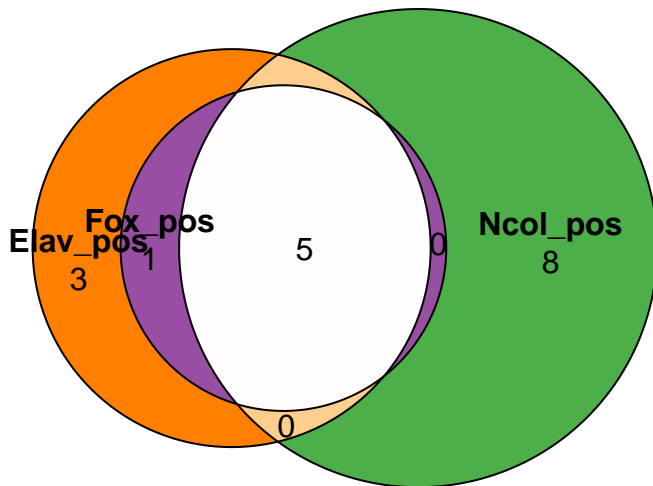


Nvec_vc1.1_XM_048720219.1
zf-C2H2.HG1.36:like:AC008;



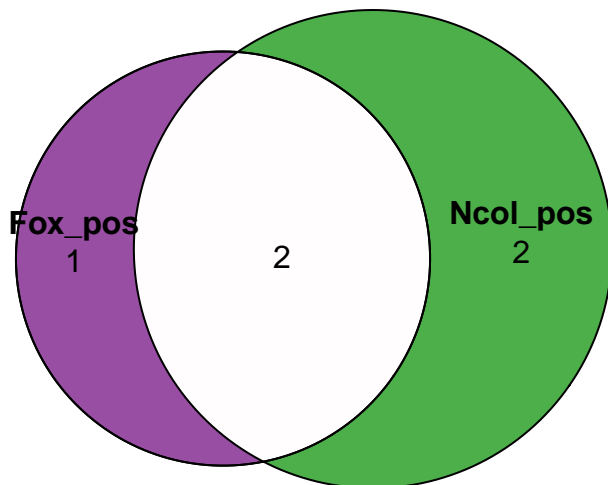
Nvec_vc1.1_XM_032381208.2

Actin-related protein 10; Actin/



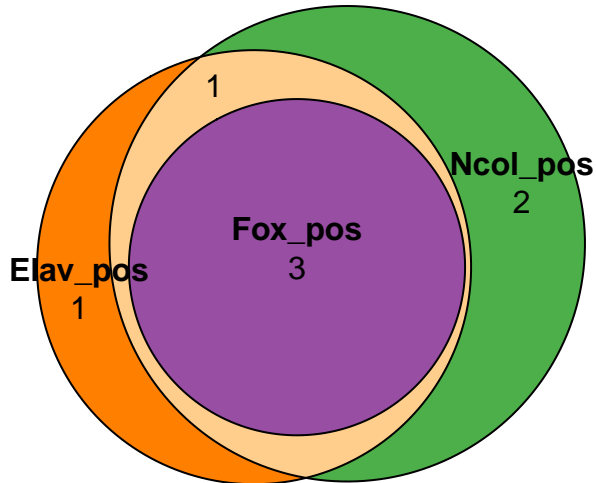
Nvec_vc1.1_XM_001626835.3

LAMB1 protein; Laminin_EGF/Laminin_EGF/



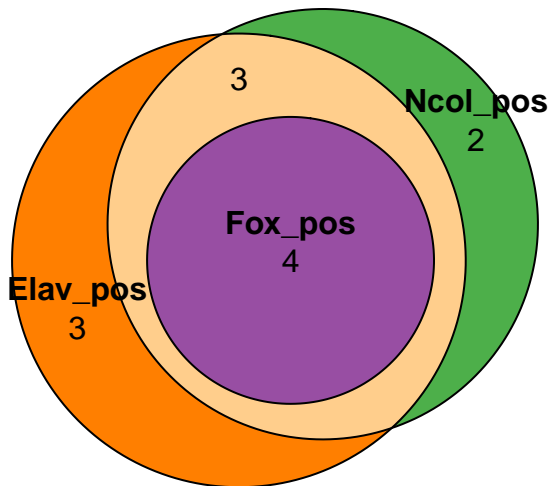
Nvec_vc1.1_XM_032363509.2

Centrosomal protein of 16;



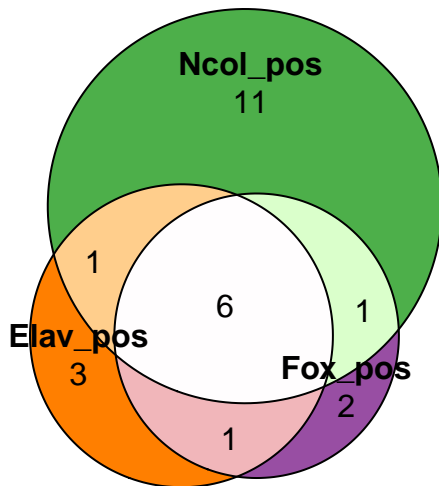
Nvec_vc1.1_XM_048733482.1

ERM_M:FRMPD3/FRMPD4; FERM_M.HG6.2:FRMPD3/FRMPD4

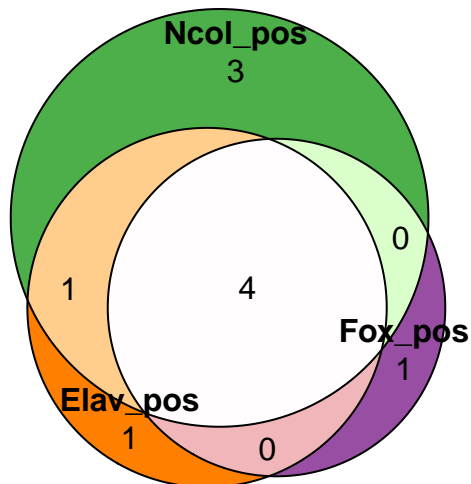


Nvec_vc1.1_XM_001640186.3

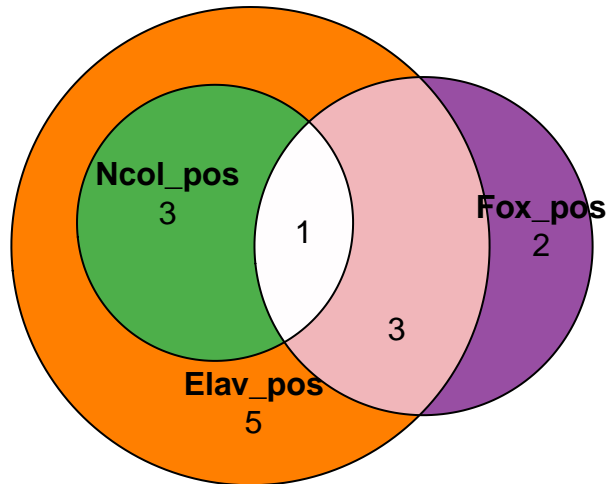
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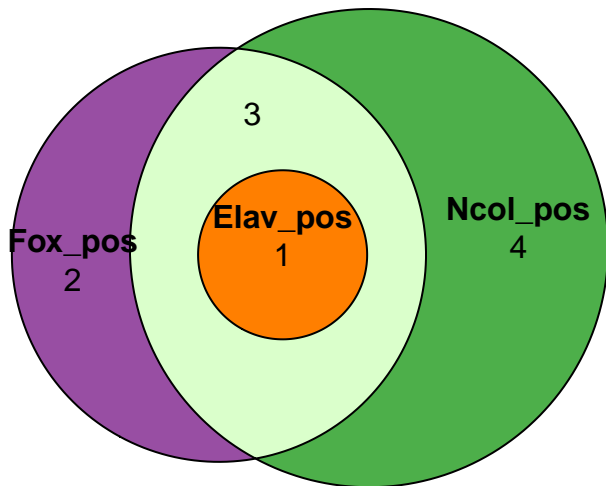
Nvec_vc1.1_XM_032361818.2
BTB:ABTB1; BTB.HG16.0:ABTB1



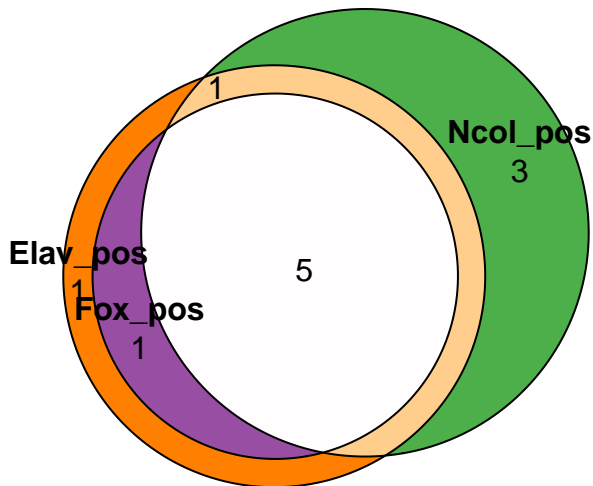
Nvec_vc1.1_XM_032375471.2
Calcium-activated potassi; CaKB/



Nvec_vc1.1_XM_032382866.2
Repressor of RNA polymera; Maf1/

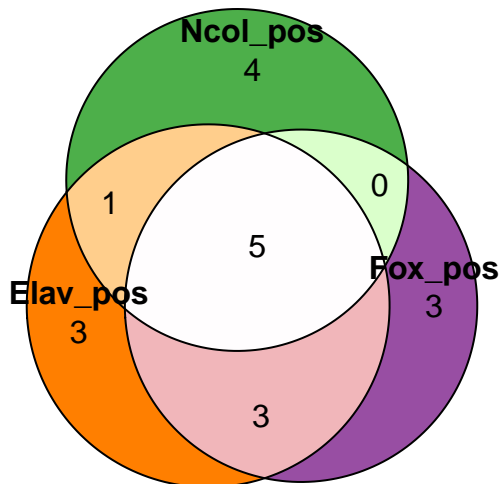


Nvec_vc1.1_XM_001628113.3
Amyloid beta protein (Cyt; TPR_10/



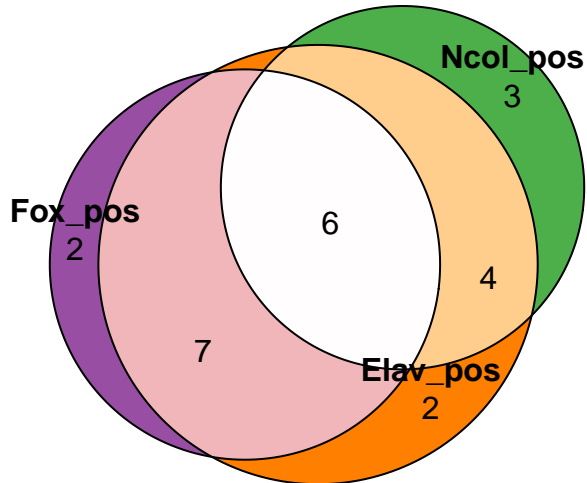
Nvec_vc1.1_XM_048734449.1

Copper chaperone for supe; Sod_Cu/

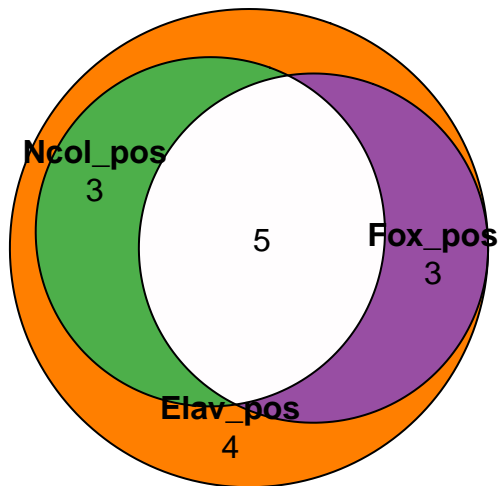


Nvec_vc1.1_XM_048731043.1

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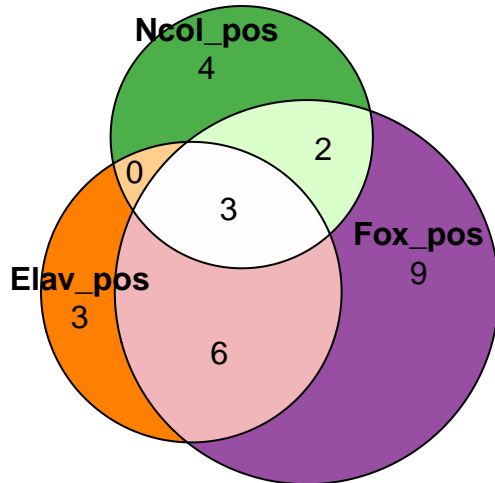


Nvec_vc1.1_XM_032364400.2
Ankyrin repeat and SOCS b; Ank_2/



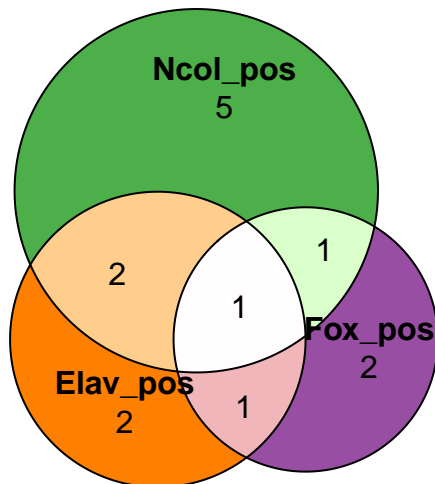
Nvec_vc1.1_XM_032363714.2

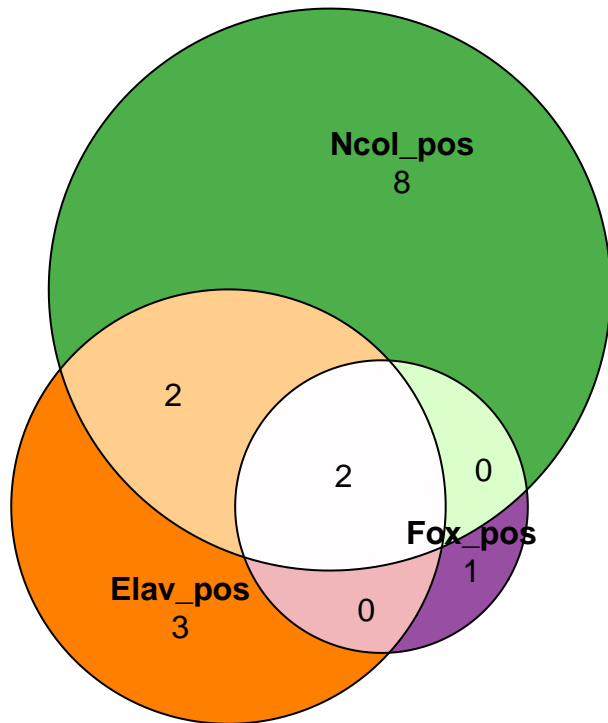
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Nvec_vc1.1_XM_032366925.2

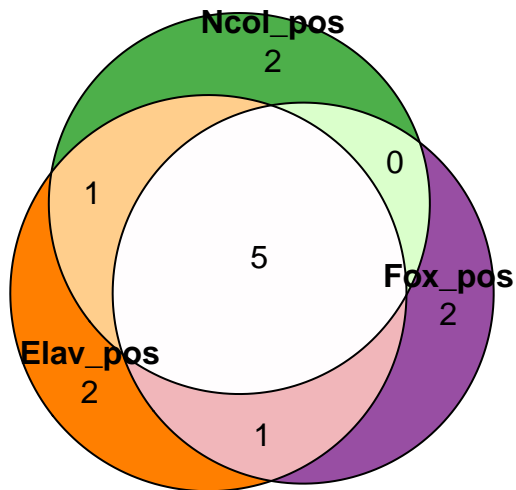
Potassium channel tetrame; BTB_2/





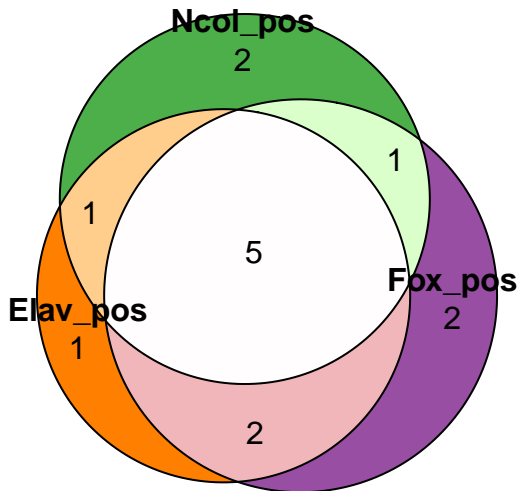
Nvec_vc1.1_XM_032367682.2

Mediator of DNA damage ch;

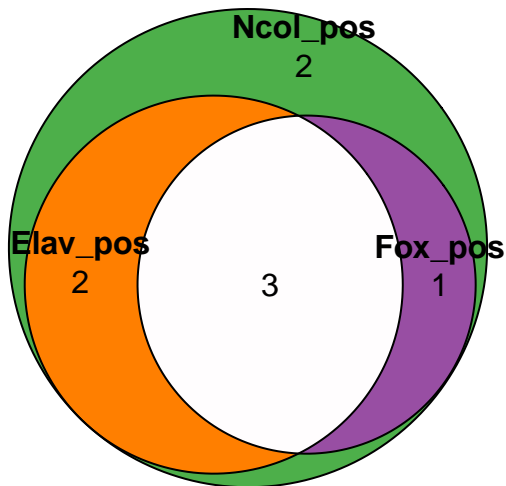


Nvec_vc1.1_XM_048721598.1

Dopey family member 1;

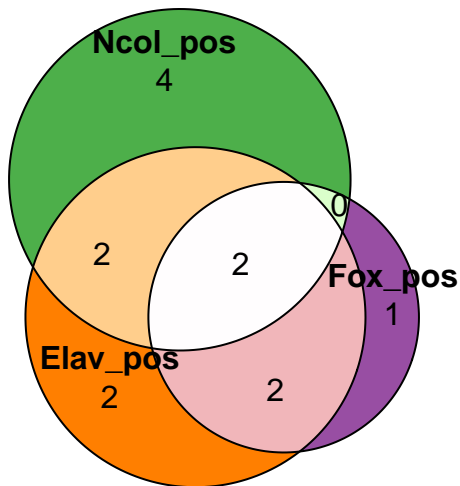


Nvec_vc1.1_XM_001628125.3
Protein-L-isoaspartate O⁻; PCMT/



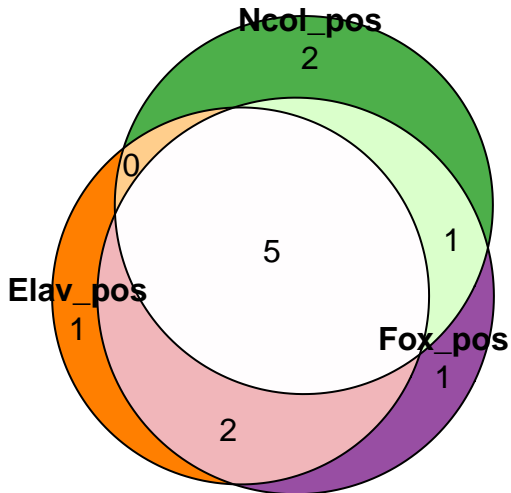
Nvec_vc1.1_XM_001639692.3

HLH.HG1.0:AC019117.4/AHR/; CLOCK/NPAS2

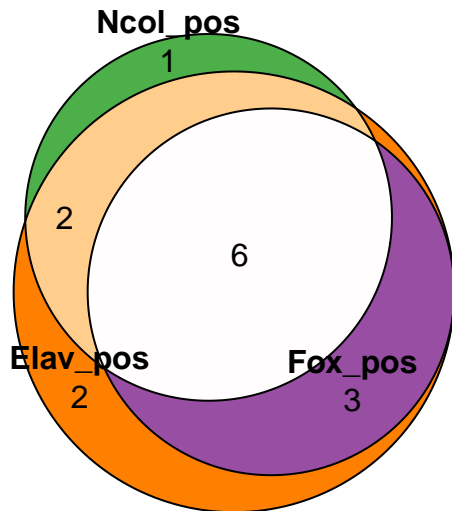


Nvec_vc1.1_XM_032362486.2

Formimidoyltransferase-cy; FTCD_N/FTCD/

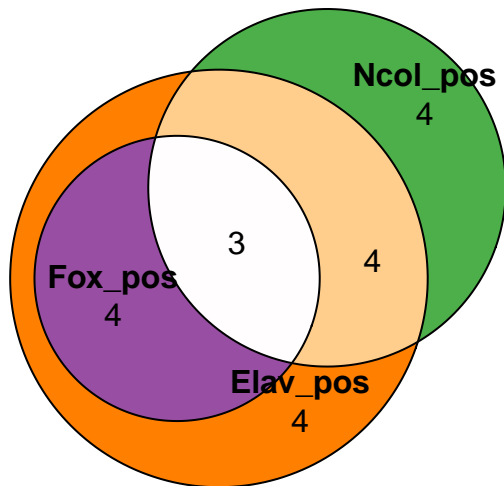


Nvec_vc1.1_XM_032365793.2
cTAGE family member 5; SH3_2/Trefoil/



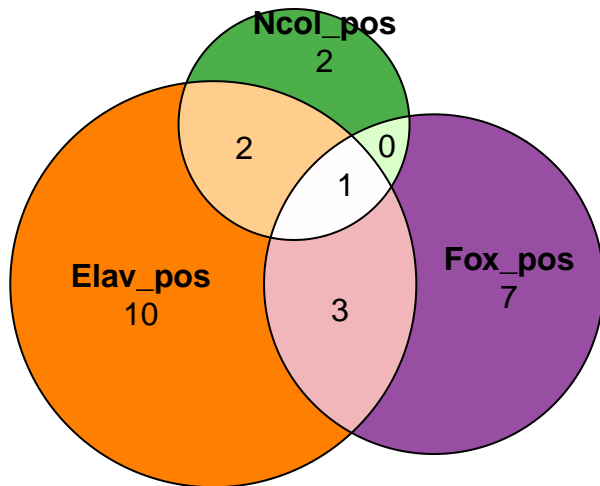
Nvec_vc1.1_XM_032384169.2

E3 ubiquitin-protein liga; zf-C3HC4_3/



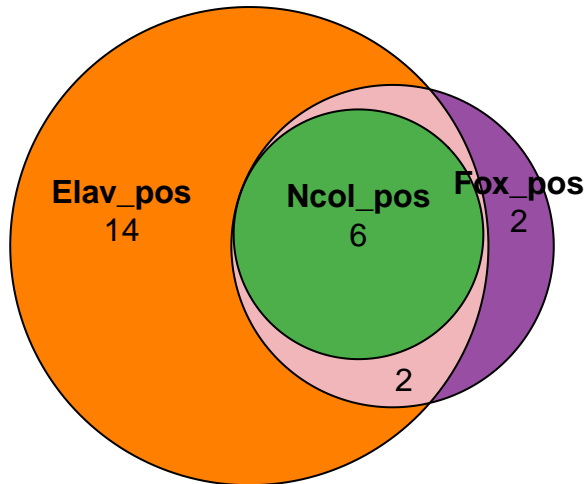
Nvec_vc1.1_XM_048734181.1

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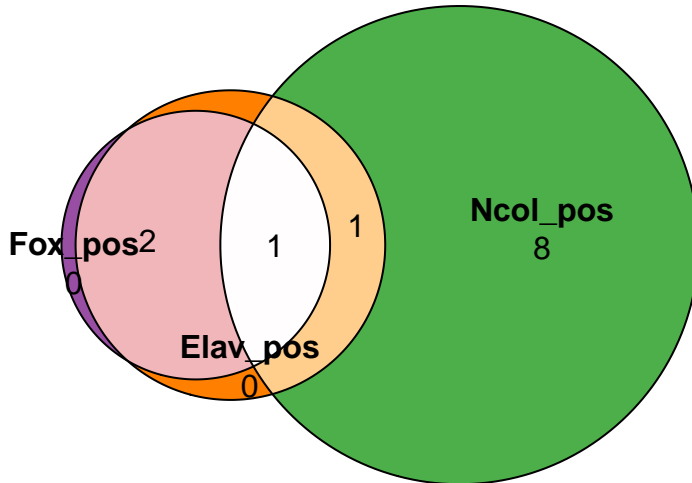


Nvec_vc1.1_XM_032374073.2

Ubiquitin-conjugating enz; UQ_con/

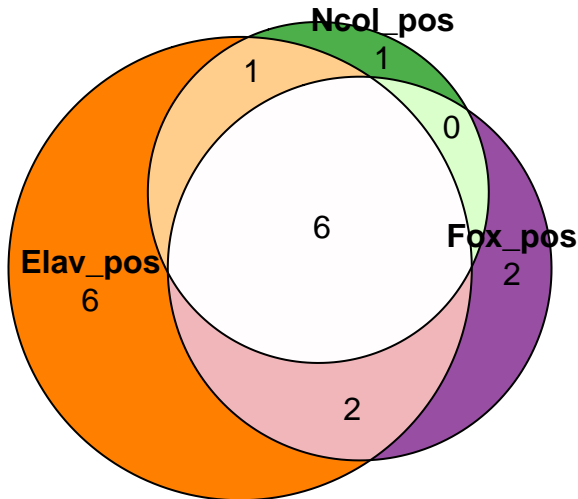


Nvec_vc1.1_XM_001639281.3
; UPAR_LY6/



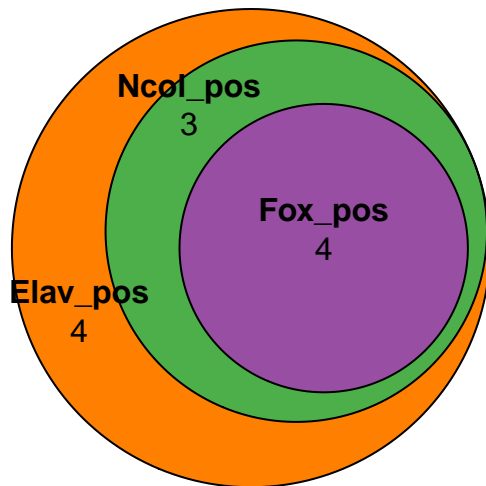
Nvec_vc1.1_XM_001638048.3

Pkinase:SIK1/SIK1B/SIK2/S; Pkinase.HG60.2:SIK1/SIK1B



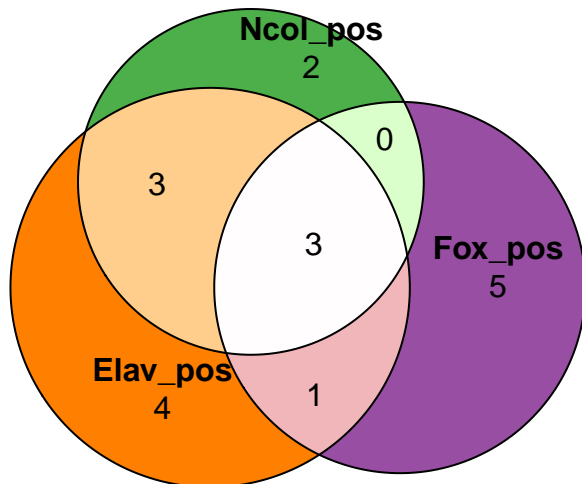
Nvec_vc1.1_XM_001633600.3

Phosphoenolpyruvate carbo; PEPCK/

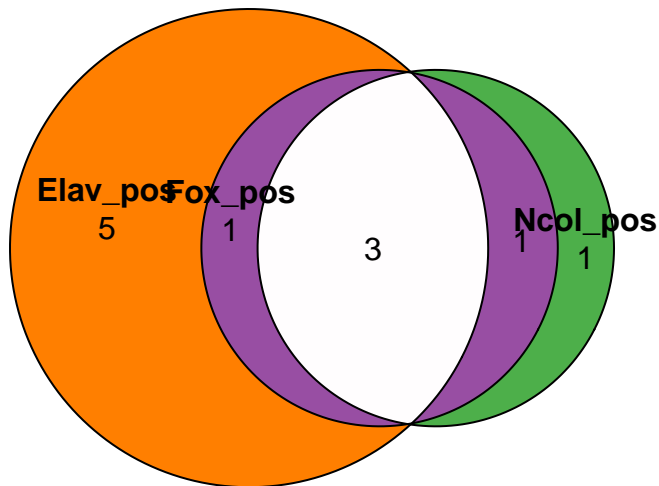


Nvec_vc1.1_XM_032374041.2

Homeodomains.HG2.9:LHX6/L; LHX6/LHX8

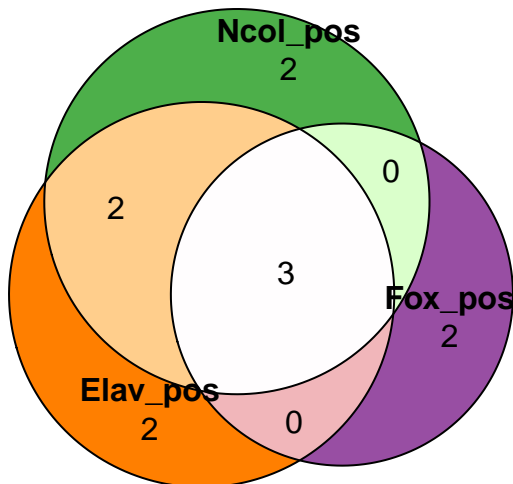


Nvec_vc1.1_XM_048726441.1
HSPC276; HBS1_N/

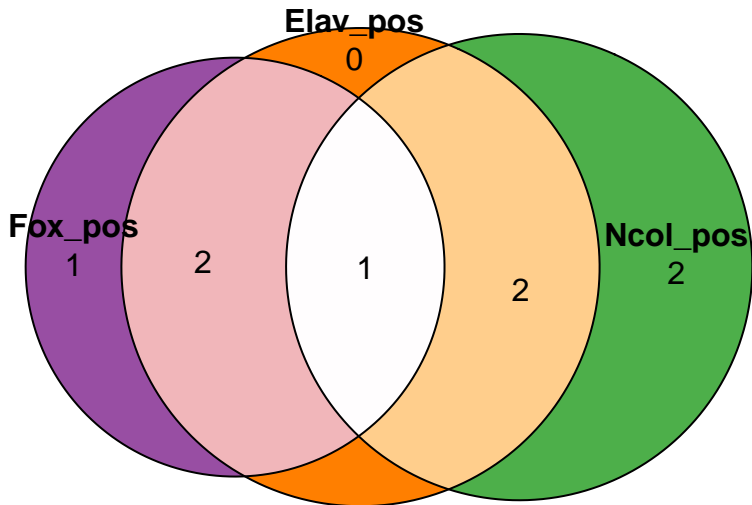


Nvec_vc1.1_XM_032366217.2

Pkinase:CSNK1A1/CSNK1A1L; Pkinase.HG31.4:CSNK1A1/CS

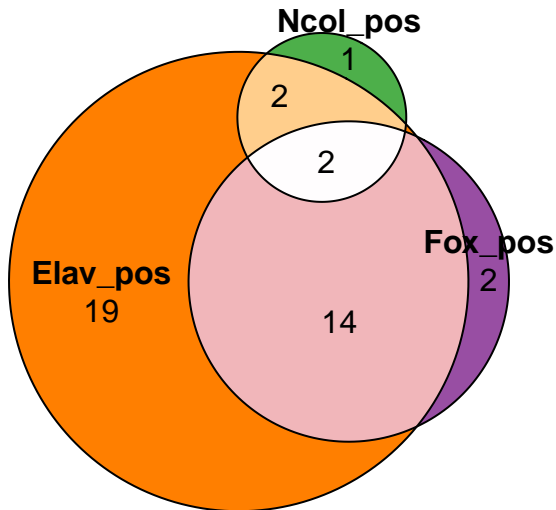


Nvec_vc1.1_XM_032365977.2
zf-C2H2.HG1.33:like:AC008;

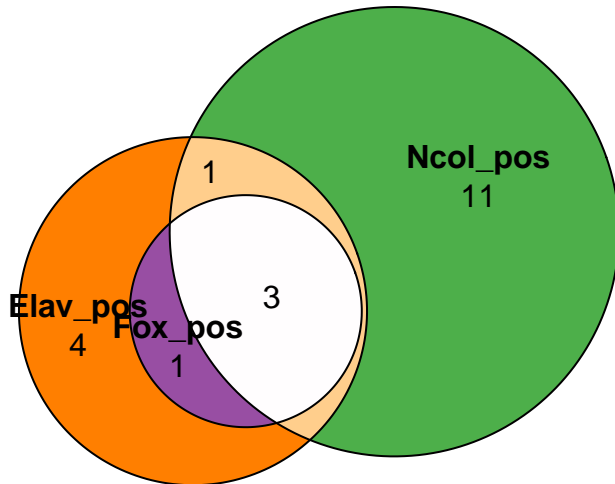


Nvec_vc1.1_XM_048729407.1

Dual specificity protein ; Rhodanese/DSPc/

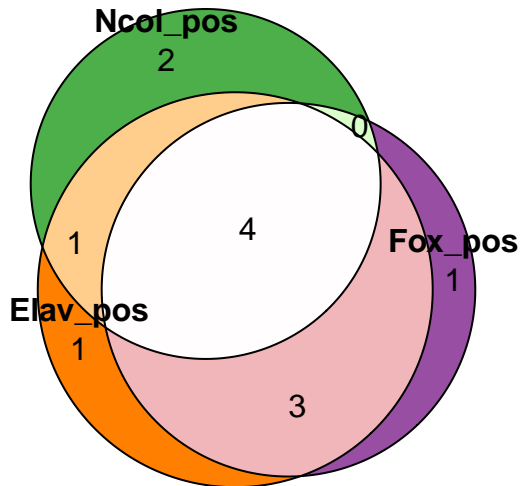


Nvec_vc1.1_XM_032366205.2
epithelial–Na:like:ASIC1/; ASC/



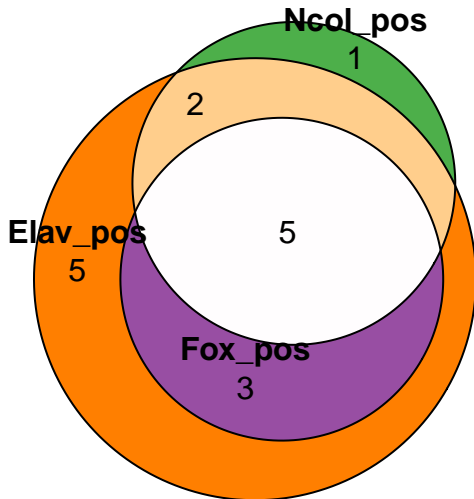
Nvec_vc1.1_XM_048722001.1

Mannosyl-oligosaccharide ; Glyco_hydro_47/

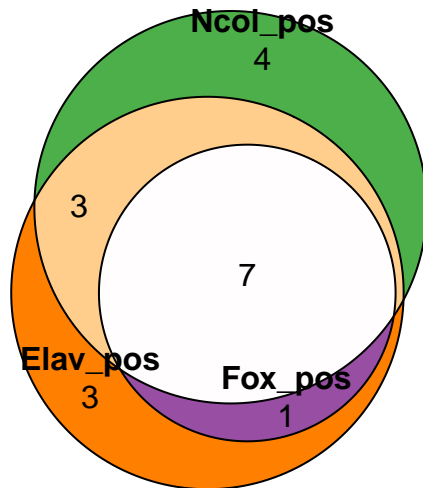


Nvec_vc1.1_XM_032377365.2

BTB:like:RHOBTB1/RHOBTB2/; Ras/BTB/BTB/

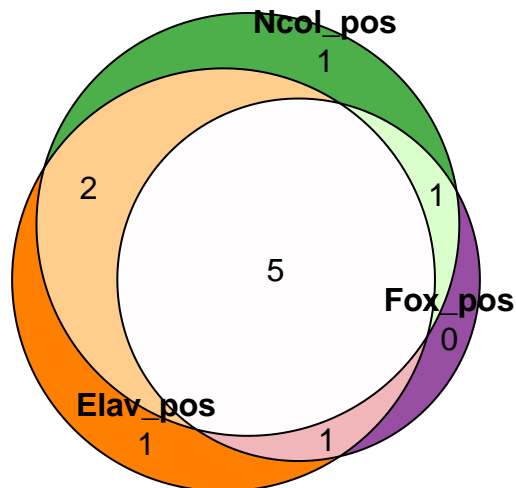


Nvec_vc1.1_XM_032363418.2
; Myb.HG9.0:NA

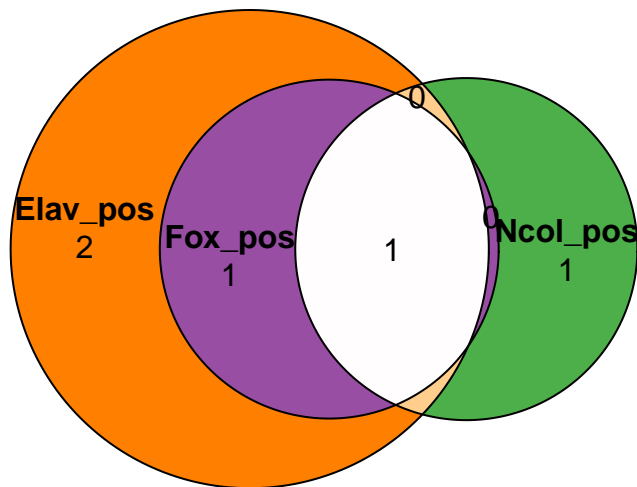


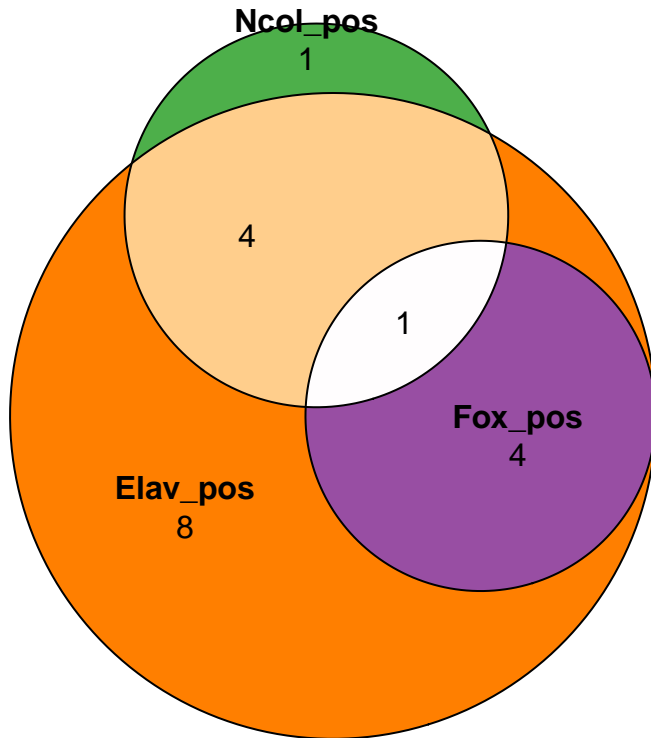
Nvec_vc1.1_XM_048722789.1

BTB:like:CCIN/GAN/KLHL11:: BTB/BACK/Kelch_1/Kelch_1/



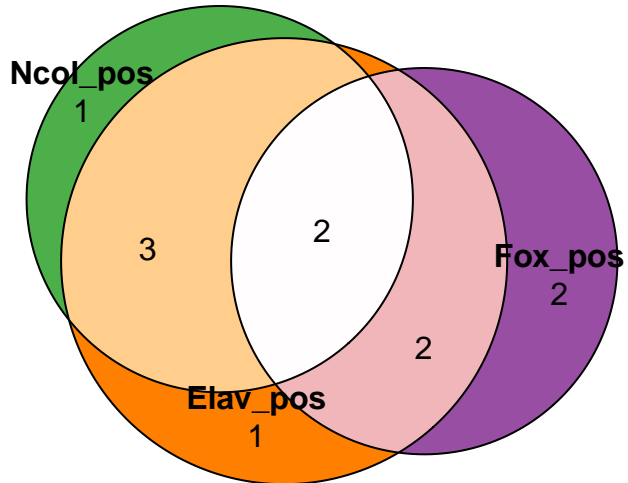
Nvec_vc1.1_XM_032373933.2
Nuclear distribution prot; NUDE_C/





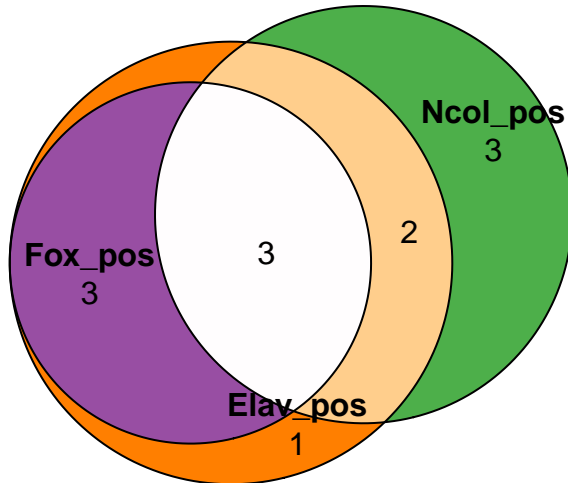
Nvec_vc1.1_XM_032363165.2

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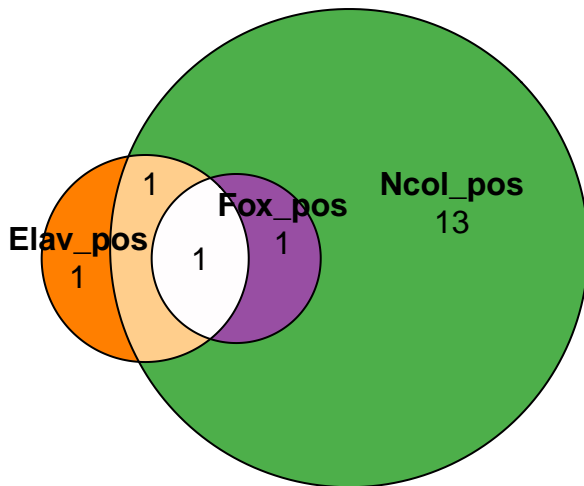
Nvec_vc1.1_XM_001638802.3

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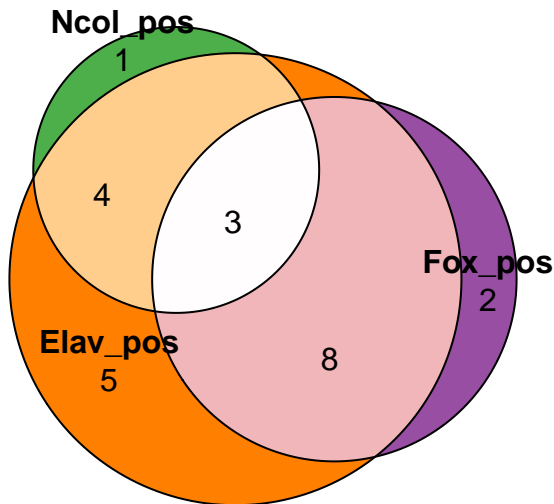
Nvec_vc1.1_XM_032374127.2

Lysine-specific histone d; Amino_oxidase/



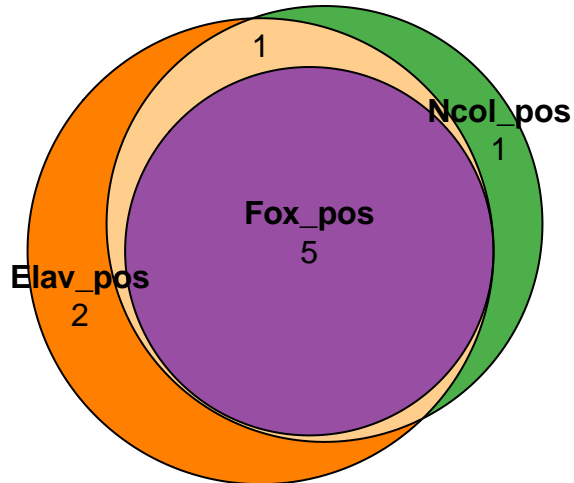
Nvec_vc1.1_XM_001639658.3

Disrupted in renal carcin; MFS_1/



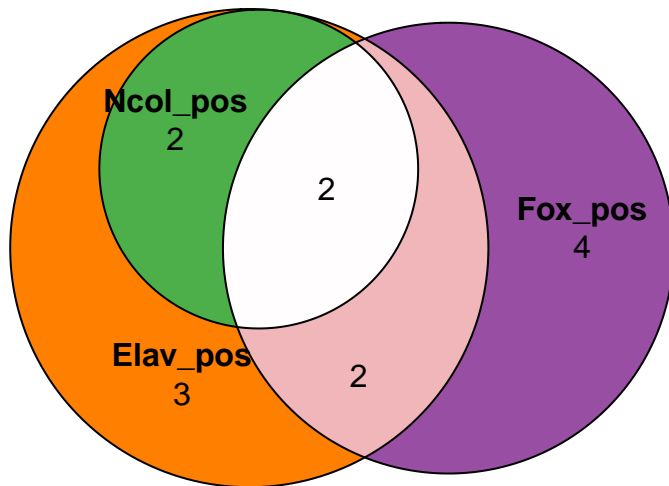
Nvec_vc1.1_XM_048731444.1

;



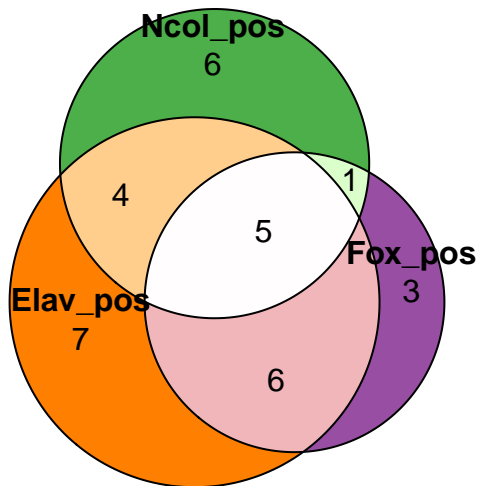
Nvec_vc1.1_XM_048732230.1

Putative uncharacterized ; Carboxyl_trans/



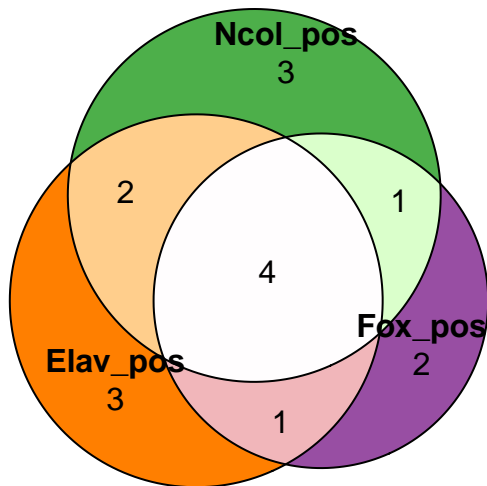
Nvec_vc1.1_XM_032366211.2

Homeodomains.HG1.34:HMX1/; HMX1/HMX2/HMX3



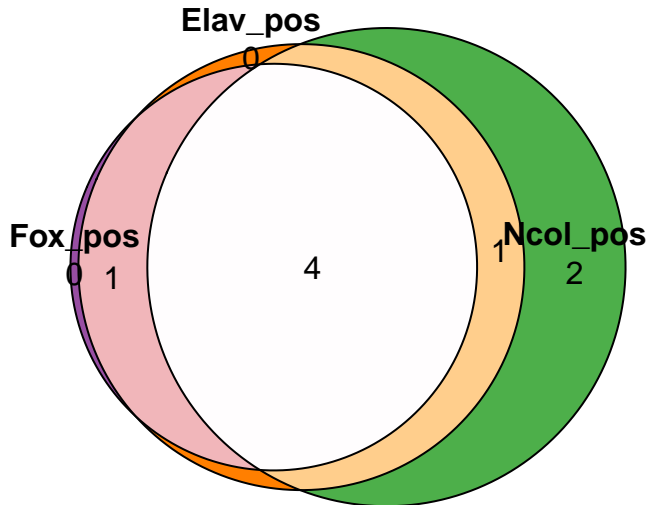
Nvec_v1g238274

;



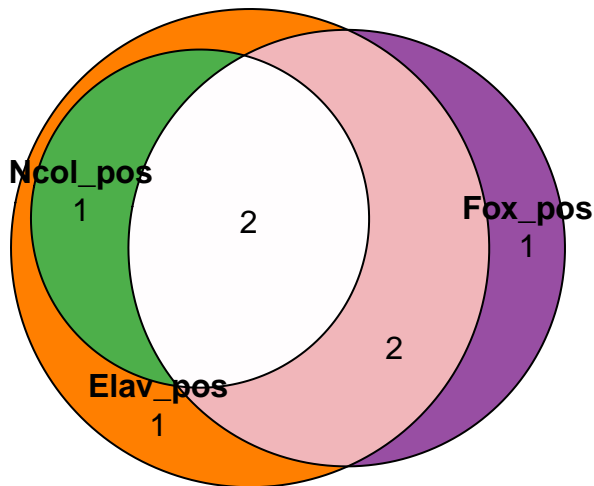
Nvec_vc1.1_XM_001641053.3

Ribonuclease kappa;



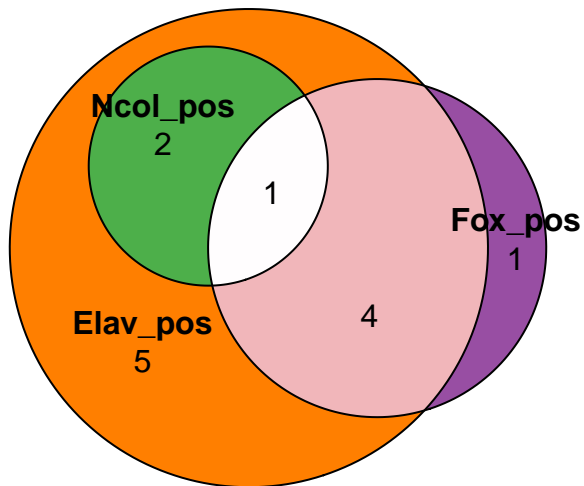
Nvec_vc1.1_XM_032366206.2

Disrupted in renal carcin; MFS_1/



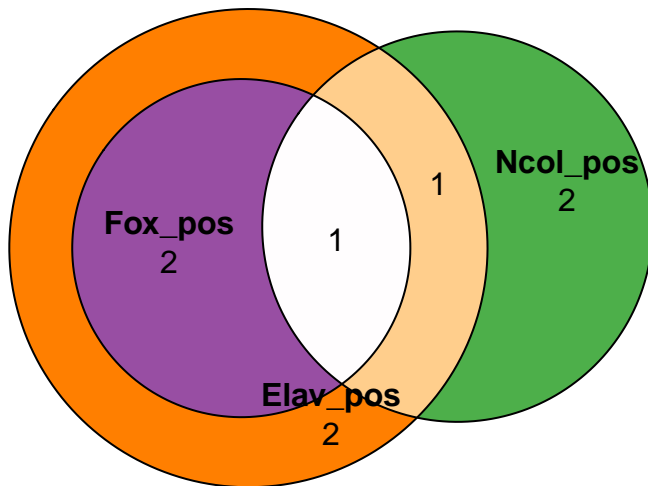
Nvec_vc1.1_XM_032362465.2

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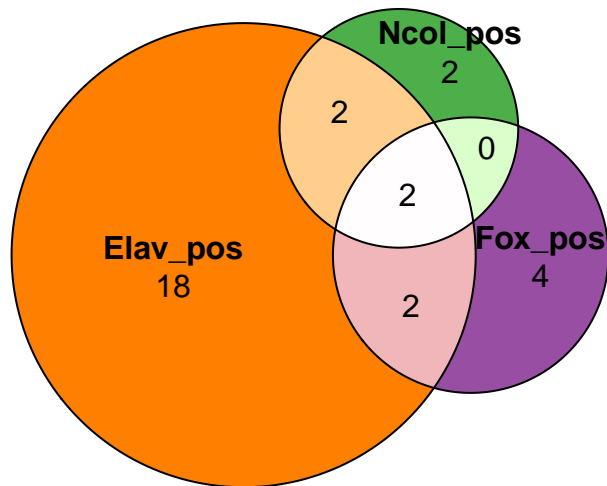
Nvec_vc1.1_XM_032372025.2

HLH.HG1.0:AC019117.4/AHR/; ARNTL/ARNTL2

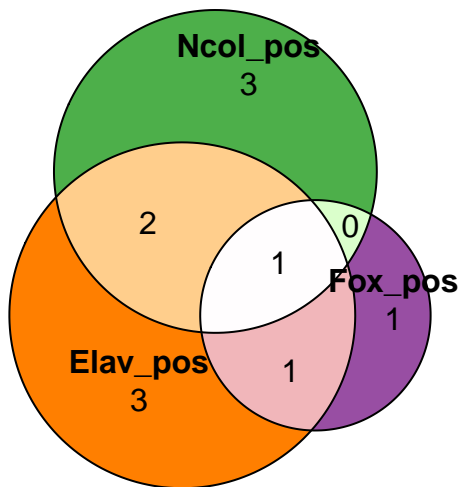


Nvec_vc1.1_XM_001635978.3

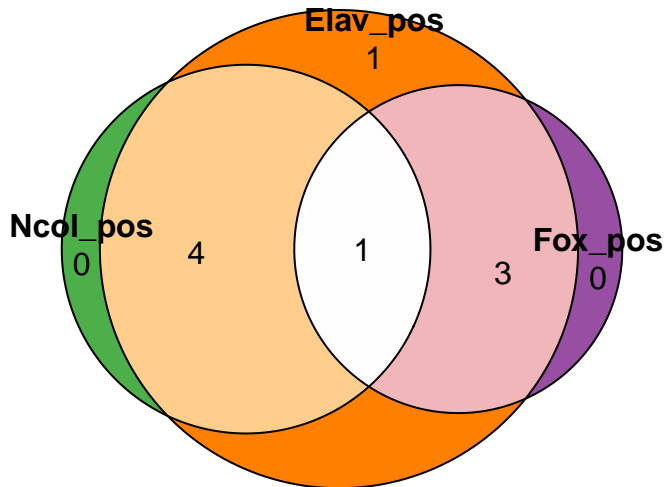
GPCR-glutamate:CASR/GPRC6; GPCR-glutamate.HG1.0:CAS



Nvec_vc1.1_XM_001633032.3
cDNA FLJ54459, highly sim; PARG_cat/

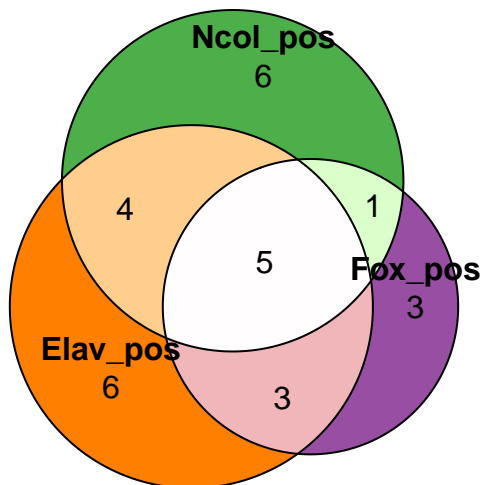


Nvec_vc1.1_XM_048728351.1
WSB1 protein; WD40/SOCS_box/



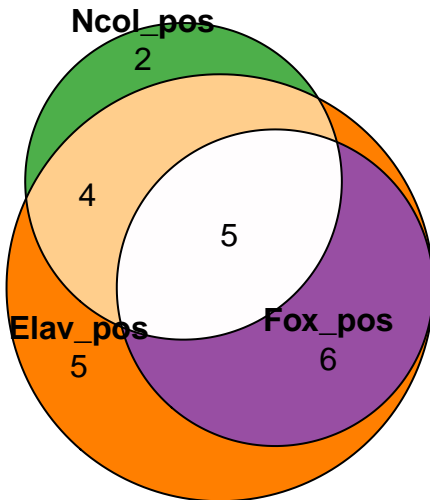
Nvec_vc1.1_XM_048722339.1

HLH.HG1.0:AC019117.4/AHR/; AC019117.4/AHR/AHRR



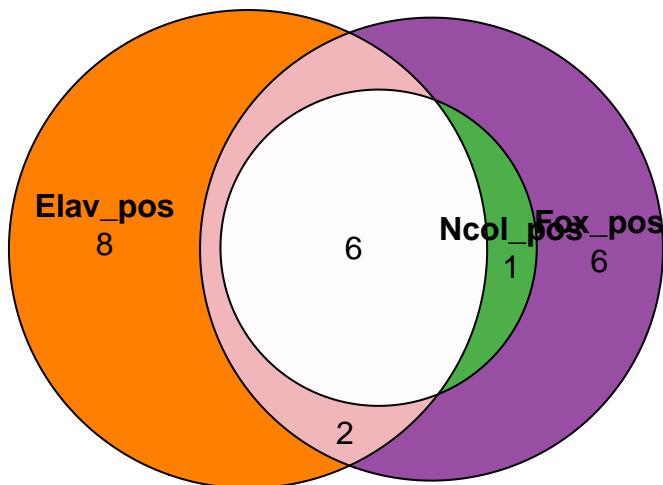
Nvec_vc1.1_XM_048727185.1

GR AF-1 specific histone ; zf-UBP/ADP_ribosyl_GH/



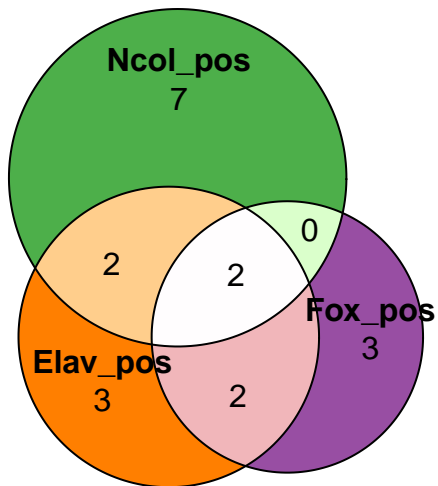
Nvec_vc1.1_XM_032384105.2

DZ:SHANK1/SHANK2/SHANK3; PDZ.HG2.12:SHANK1/SHANK



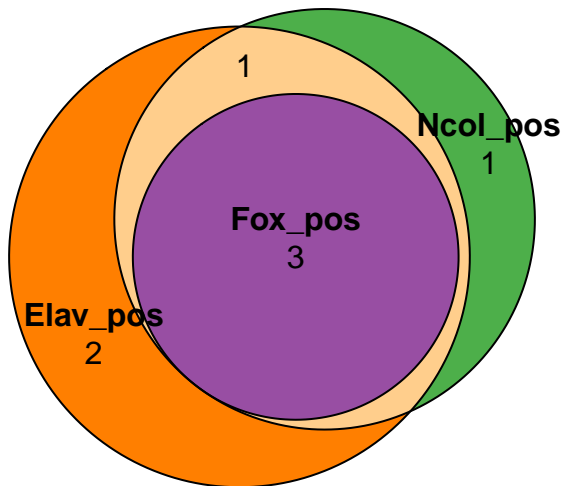
Nvec_vc1.1_XM_032384841.2

UHRF1BP1 protein;



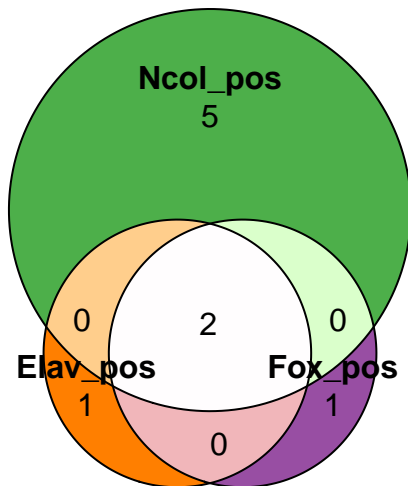
Nvec_vc1.1_XM_048721983.1

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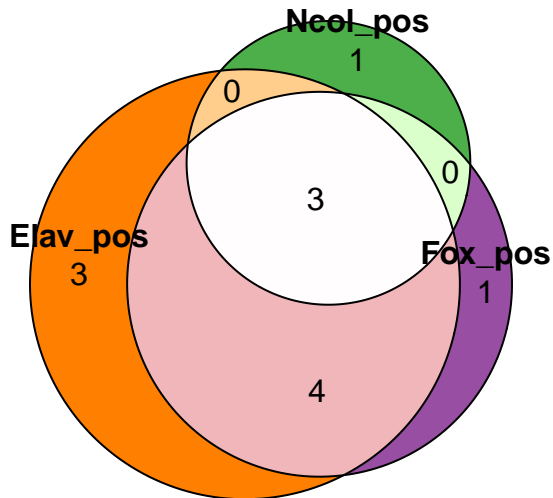


Nvec_vc1.1_XM_001640949.3

Transmembrane protein 106; DUF1356/

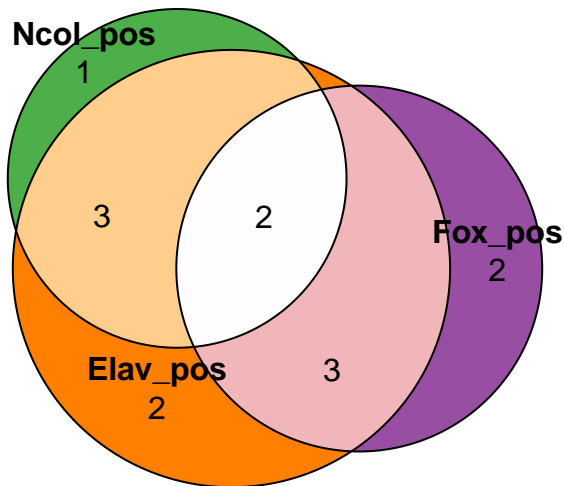


Nvec_vc1.1_XM_048723524.1
NFX1_ZF.HG1.1:NFX1; NFX1



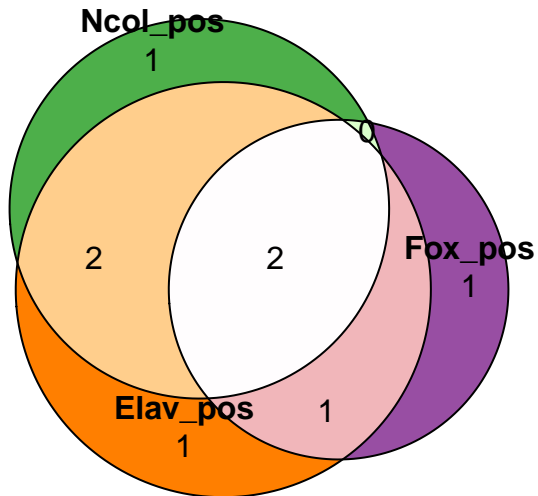
Nvec_vc1.1_XM_032386734.2

Ubiquitin conjugation fac; Ufd2P_core/Ufd2P_core/U-b



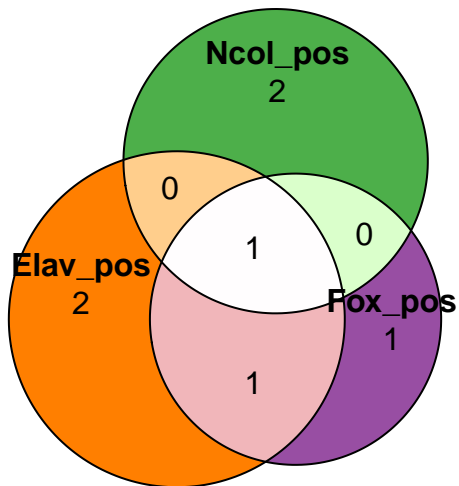
Nvec_vc1.1_XM_001633494.3

Tigger transposable eleme; DDE_1/



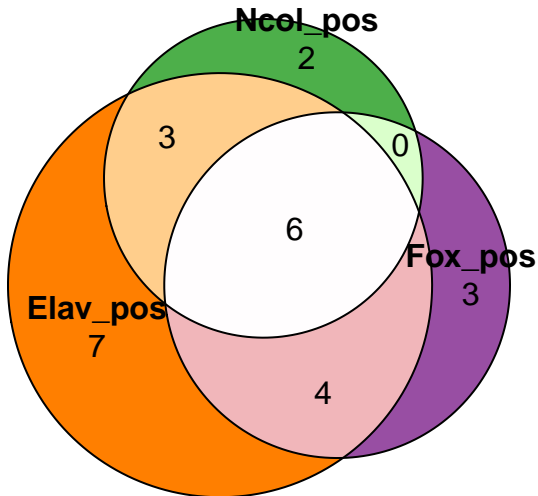
Nvec_vc1.1_XM_048731996.1

Phosphatidylinositol 3,4,; Exo_endo_phos/

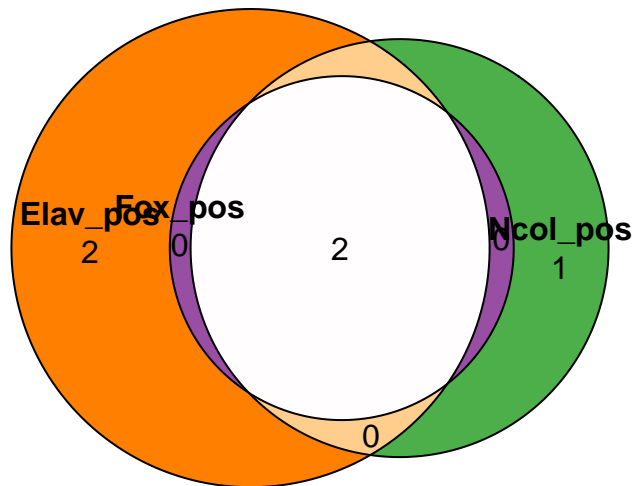


Nvec_vc1.1_XM_001636428.3

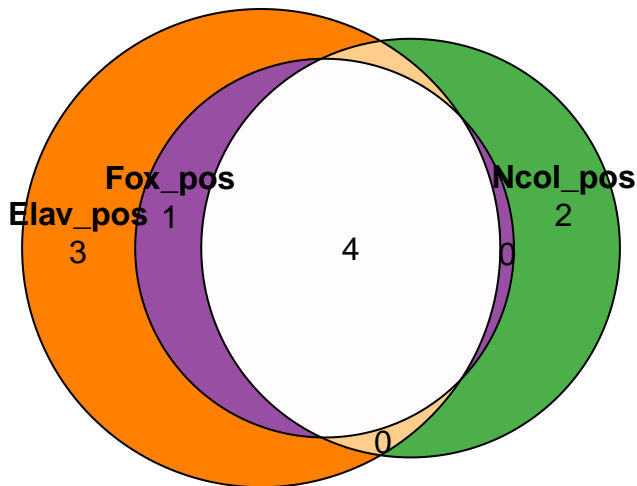
Chromosome 14 open readin; DUF766/



Nvec_vc1.1_XM_001628341.3
Progesterin and adipoQ rece; HlyIII/

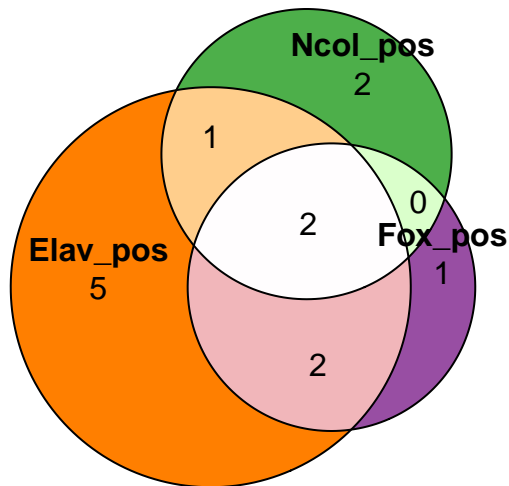


Nvec_vc1.1_XM_048720046.1
Tubulin-specific chaperon; TFCD_C/



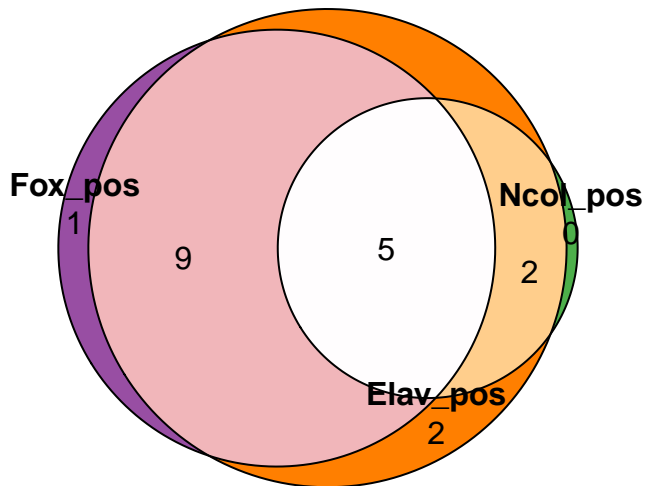
Nvec_vc1.1_XM_001625226.3

Solute carrier family 15 ; PTR2/PTR2/



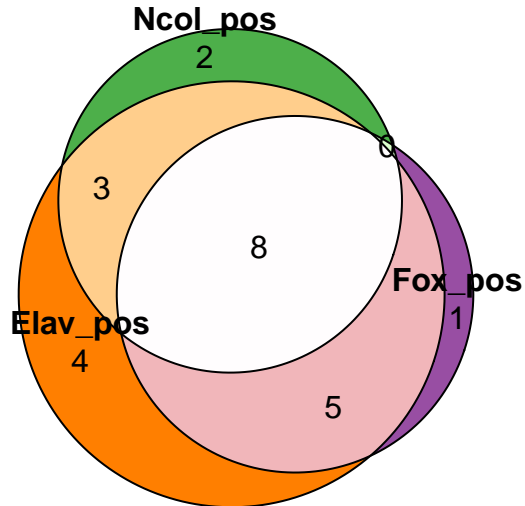
Nvec_vc1.1_XM_001632898.3

Cationic amino acid trans; AA_permease_2/

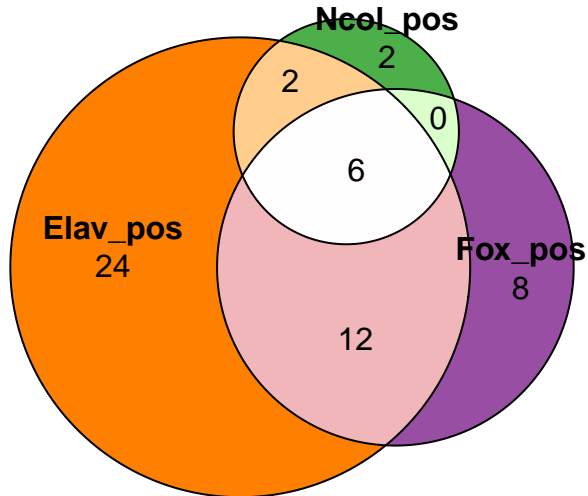


Nvec_vc1.1_XM_001620810.3

E3 ubiquitin-protein liga;

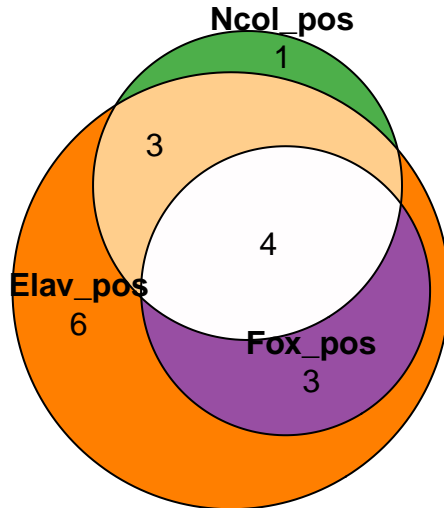


Nvec_vc1.1_XM_032378223.2
cDNA FLJ35331 fis, clone ;



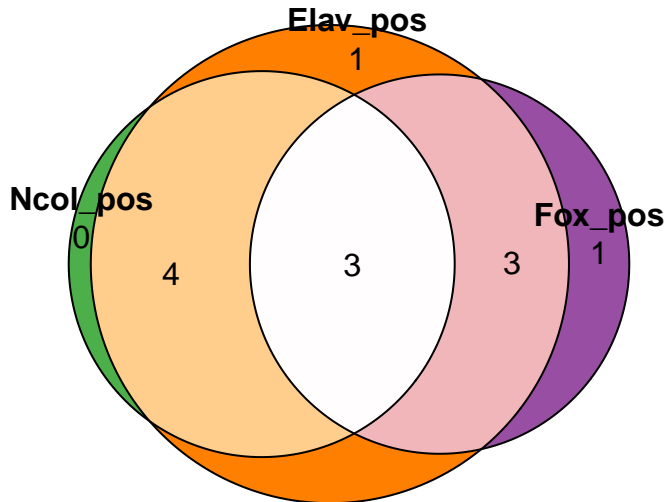
Nvec_vc1.1_XM_001626170.3

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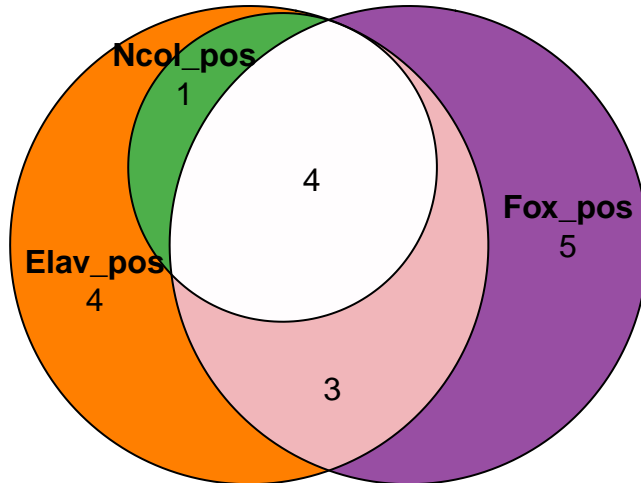
Nvec_vc1.1_XM_032376793.2

Ubiquitin-conjugating enz; UQ_con/



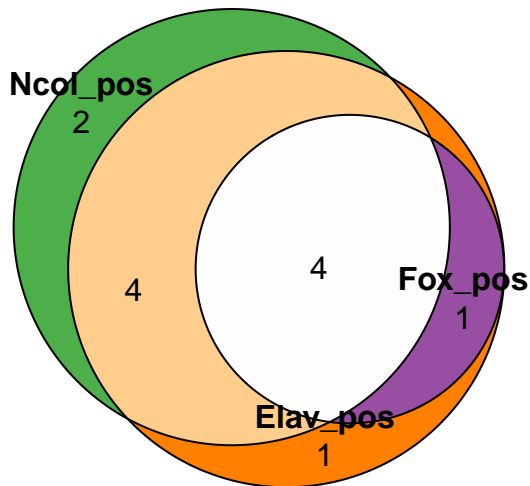
Nvec_vc1.1_XM_048728336.1

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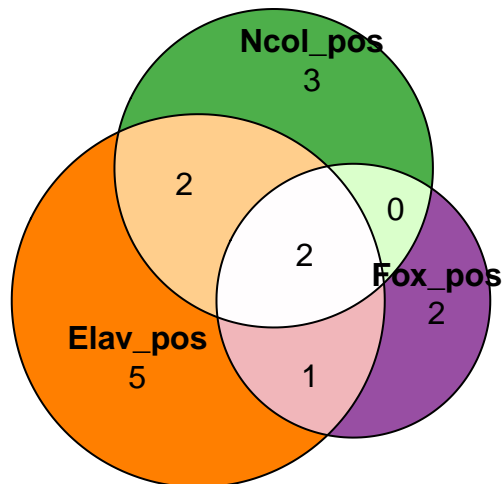
Nvec_vc1.1_XM_032382276.2

Macoilin; Macoilin/Macoilin/



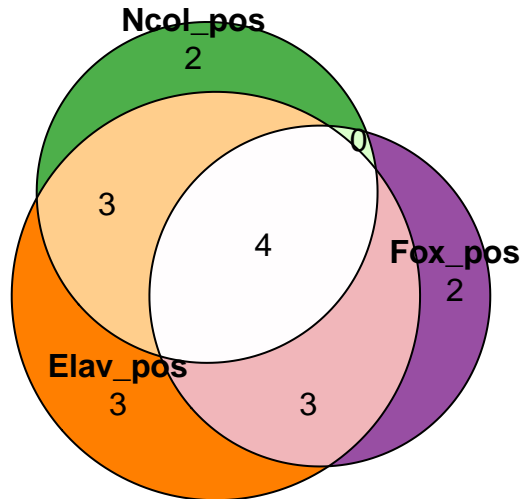
Nvec_vc1.1_XM_001631176.3

Pkinase:RPS6KA4/RPS6KA5; Pkinase.HG120.0:RPS6KA4/R



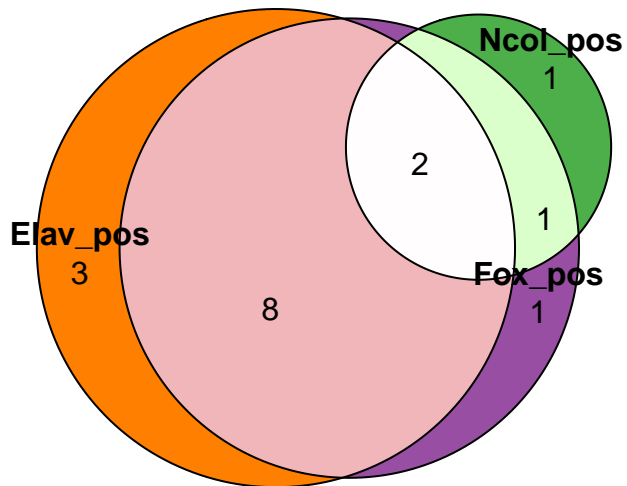
Nvec_vc1.1_XM_032385492.2

Phosphoinositide phosphol; EF-hand_like/PI-PLC-X/PI-

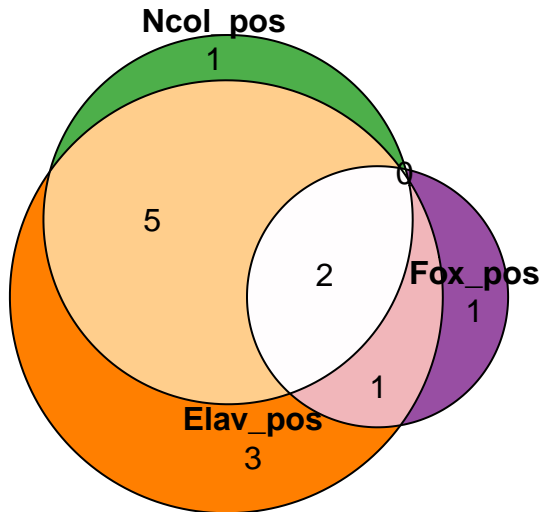


Nvec_vc1.1_XM_048734633.1

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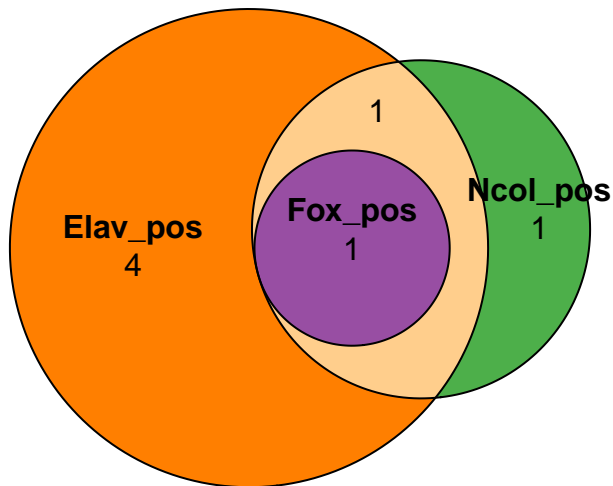


Nvec_vc1.1_XM_032367447.2
Sodium-dependent neutral ; SNF/



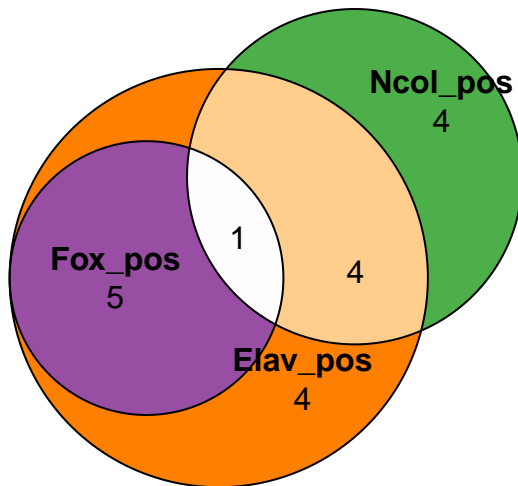
Nvec_vc1.1_XM_032381065.2

ion-NaKCa:TRPA1; ion-NaKCa.HG4.1:TRPA1

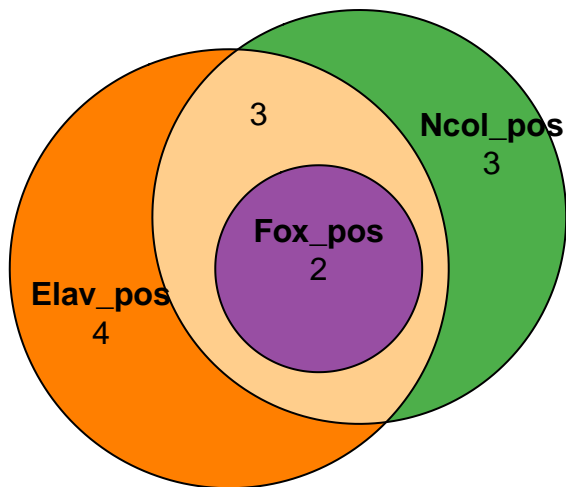


Nvec_vc1.1_XM_048726563.1

BSD domain-containing pro; BSD/

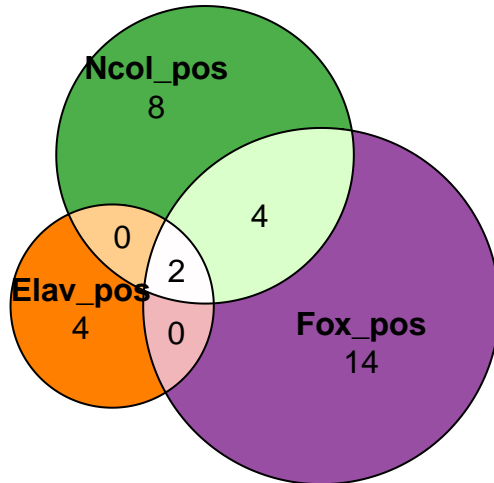


Nvec_vc1.1_XM_001624118.3
Glucose-6-phosphatase 2; PAP2/



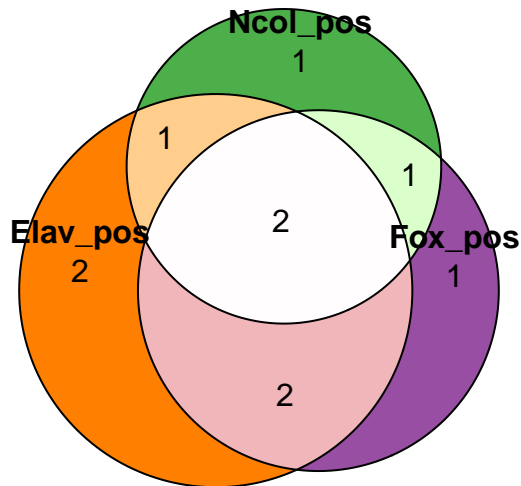
Nvec_vc1.1_XM_048730697.1

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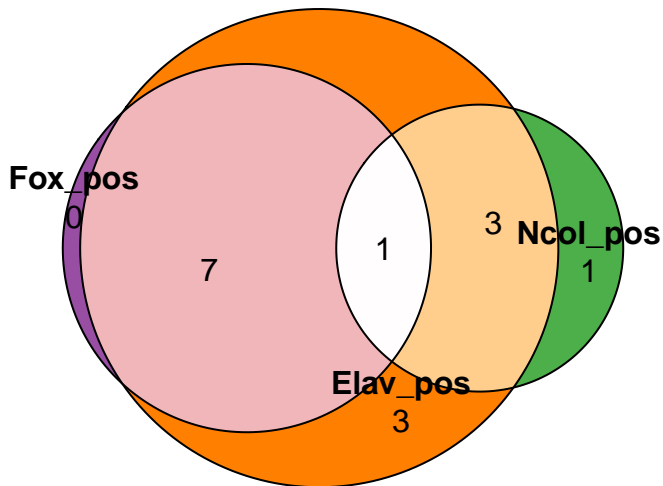
Nvec_vc1.1_XM_048726080.1

Oxidative stress-responsi;



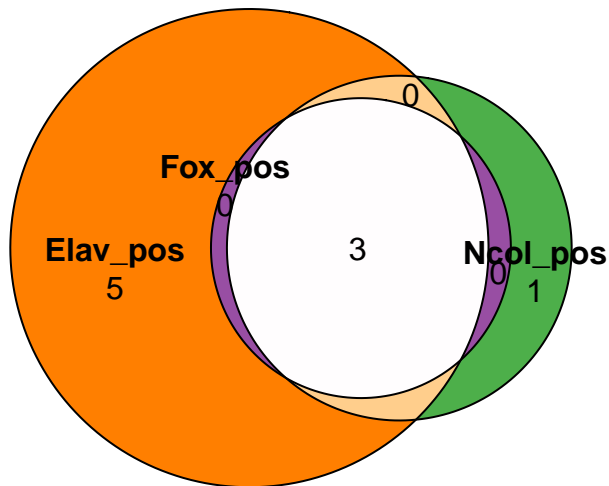
Nvec_vc1.1_XM_048732328.1

fibronectin:like:FNDC3A/F; fn3/fn3/fn3/fn3/fn3/fn3/f



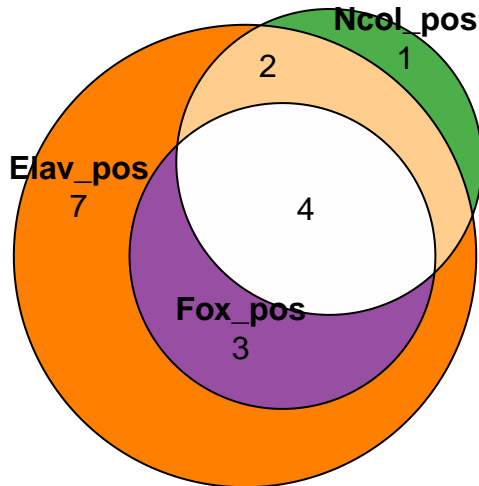
Nvec_vc1.1_XM_048725313.1

N6-adenosine-methyltransf; MT-A70/

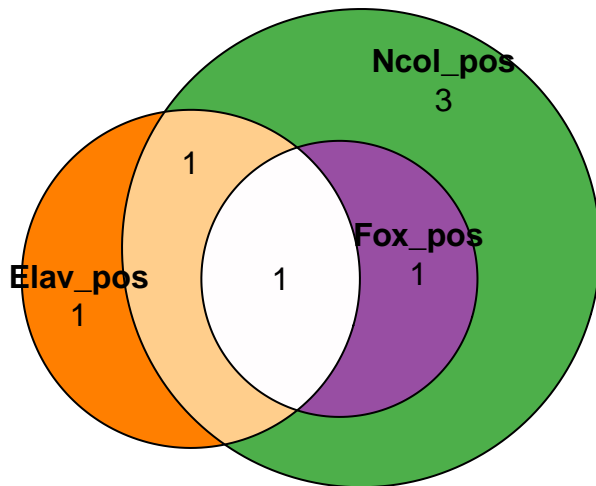


Nvec_vc1.1_XM_048732109.1

P2X purinoceptor 7;

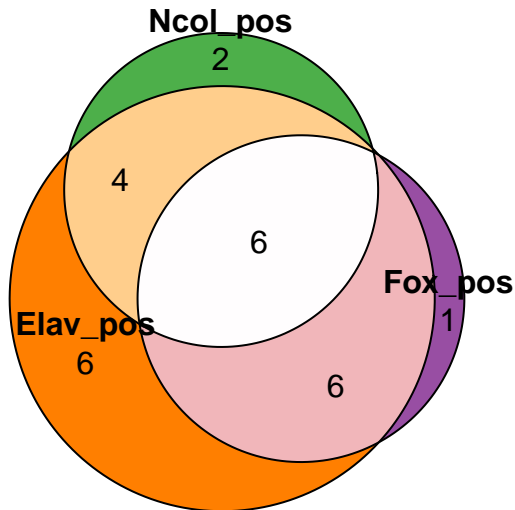


Nvec_vc1.1_XM_032374236.2
HCG1811773; zf-RING_2/CUE/

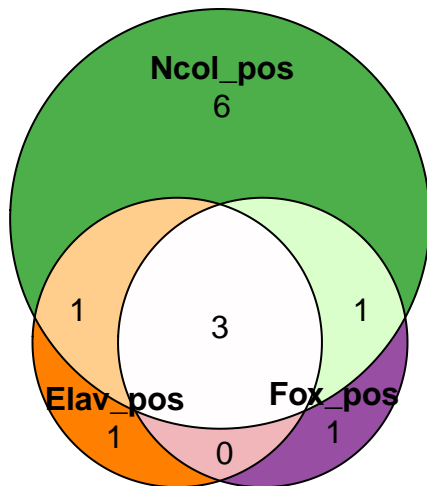


Nvec_vc1.1_XM_001627225.3

Lipoma HMGIC fusion partn; L_HGMIC_fpl/

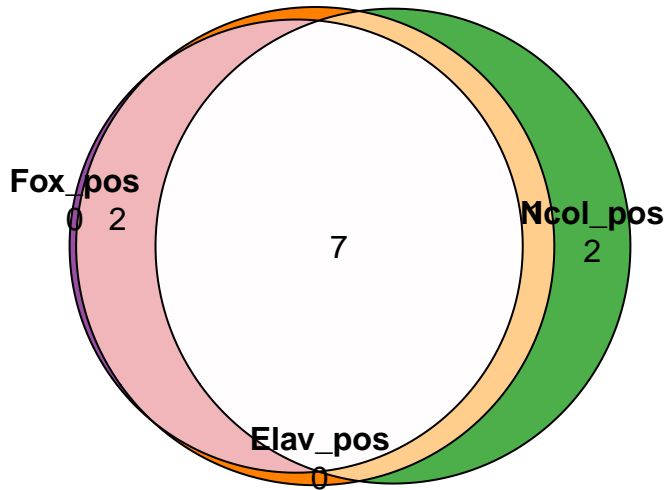


Nvec_vc1.1_XM_001640821.3
Lysoplasmalogenase-like p; YhhN/



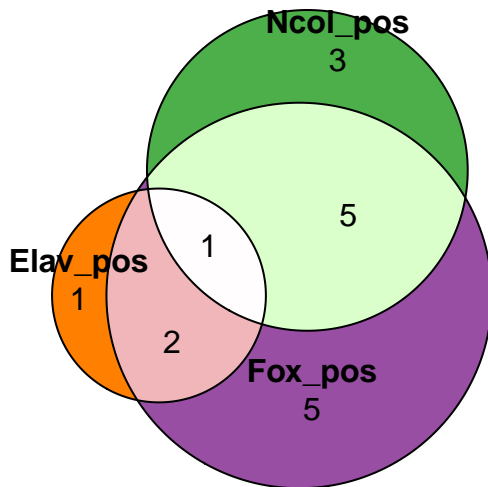
Nvec_vc1.1_XM_048734662.1

Golgi-specific brefeldin ; Sec7_N/Sec7/



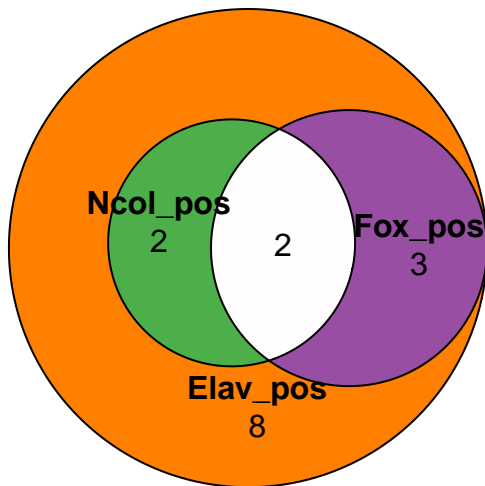
Nvec_vc1.1_XM_001636754.3

TB:BTBD1/BTBD2/BTBD3/BTB; BTB.HG3.0:BTBD1/BTBD2/BT

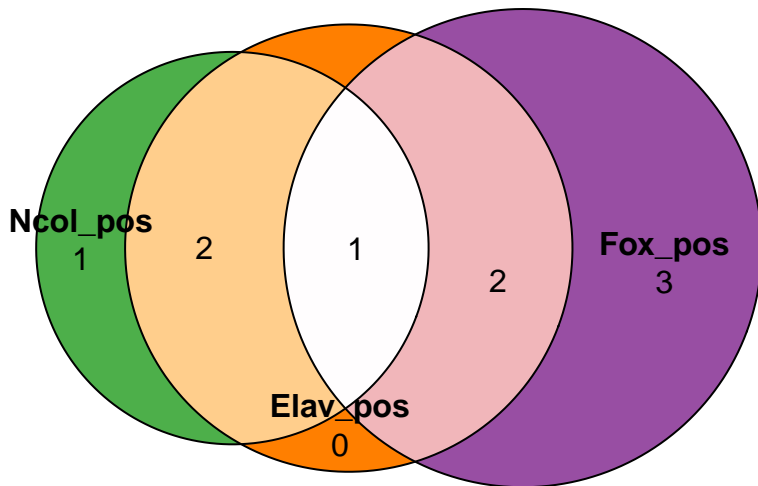


Nvec_vc1.1_XM_032364941.2

Pkinase:TNK1/TNK2; Pkinase.HG78.0:TNK1/TNK2

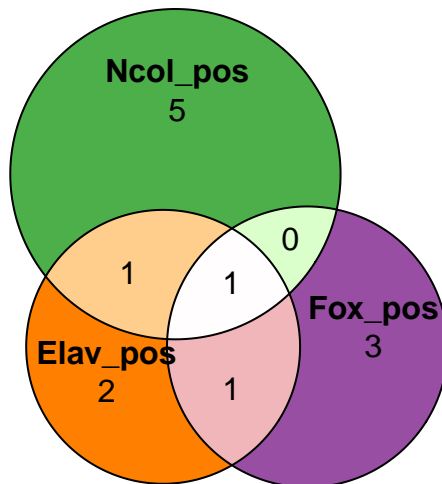


Nvec_vc1.1_XM_048726140.1
; Herpes_alk_exo/



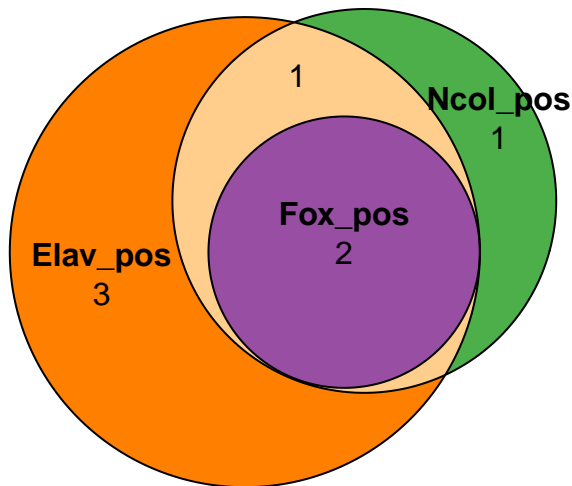
Nvec_vc1.1_XM_032362594.2

MARVEL:like:CMTM3/CMTM4/C; MARVEL/



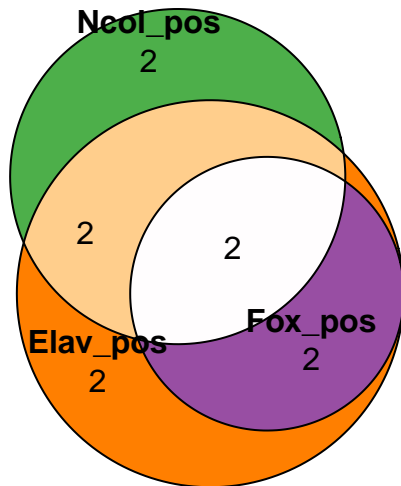
Nvec_vc1.1_XM_032381104.2

Structural maintenance of; Trehalose_recP/HATPase_c_



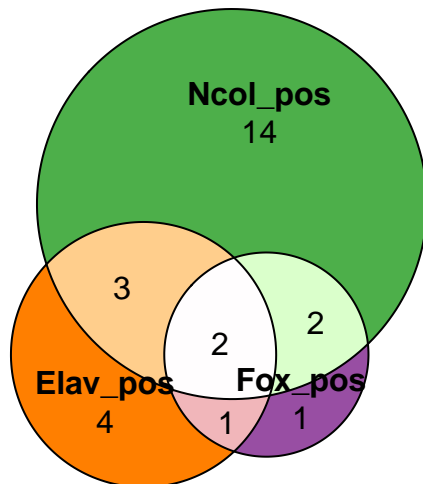
Nvec_vc1.1_XM_048728032.1

Snurportin-1; Snurportin1/



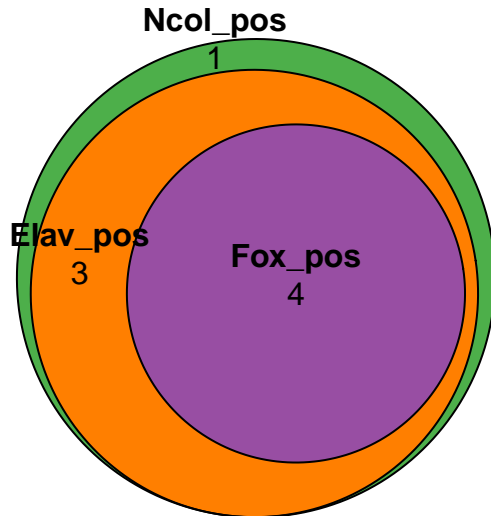
Nvec_vc1.1_XM_001638049.3

Platelet-activating facto; Lipase_GDSL_2/



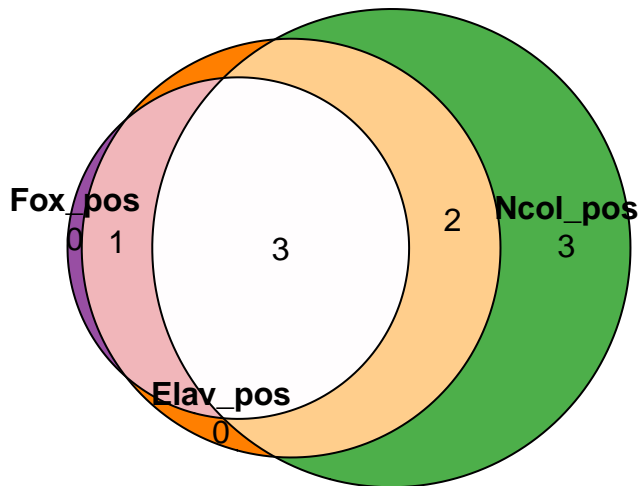
Nvec_vc1.1_XM_032375382.2

Transmembrane protein 104; Aa_trans/



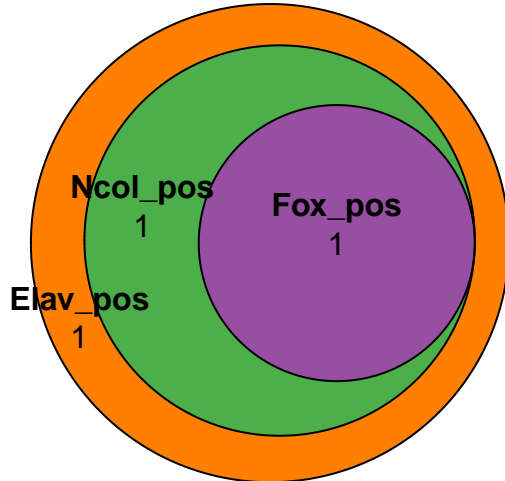
Nvec_vc1.1_XM_001636301.3

Poly (ADP-ribose) polymer;



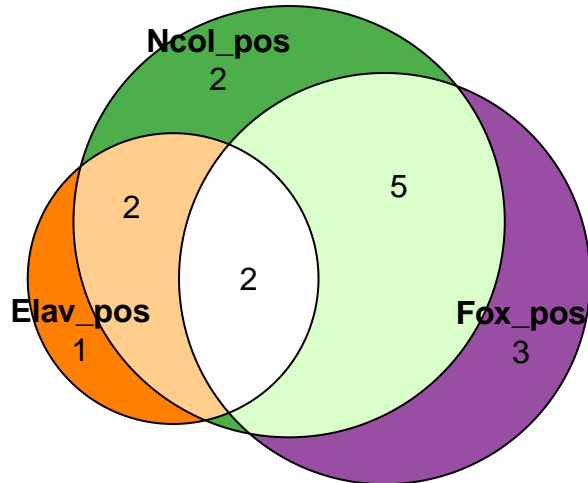
Nvec_vc1.1_XM_001630350.3

Importin subunit alpha; IBB/Arm/Arm/Arm/Arm/Arm/A



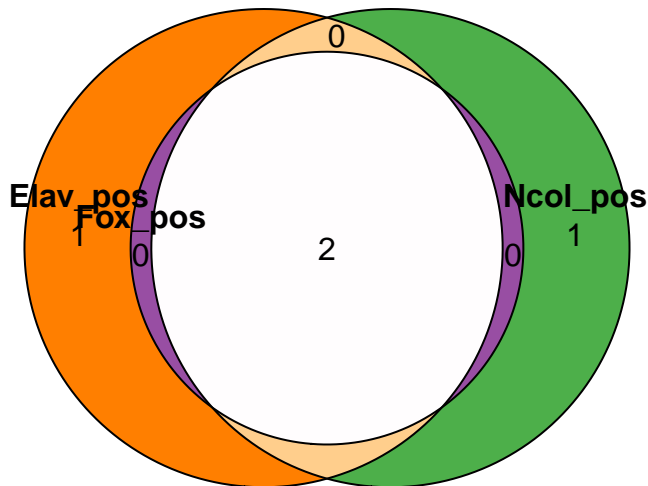
Nvec_vc1.1_XM_048727498.1

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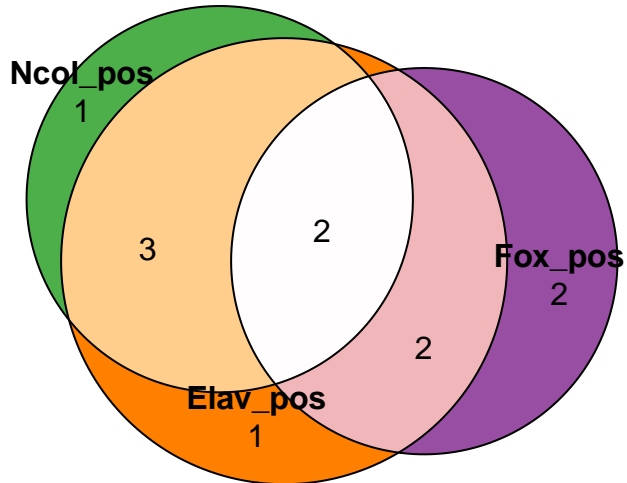
Nvec_vc1.1_XM_048726451.1

Pkinase:RPS6KB1/RPS6KB2; Pkinase.HG122.0:RPS6KB1/R

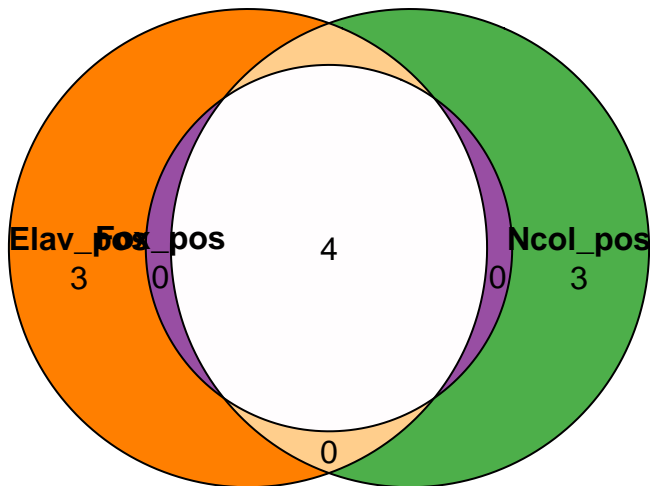


Nvec_vc1.1_XM_048724381.1

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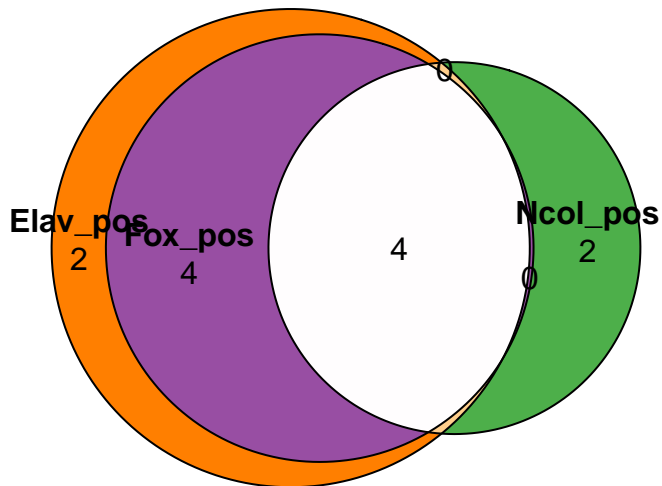


Nvec_vc1.1_XM_032386760.2
cDNA FLJ34132 fis, clone ;

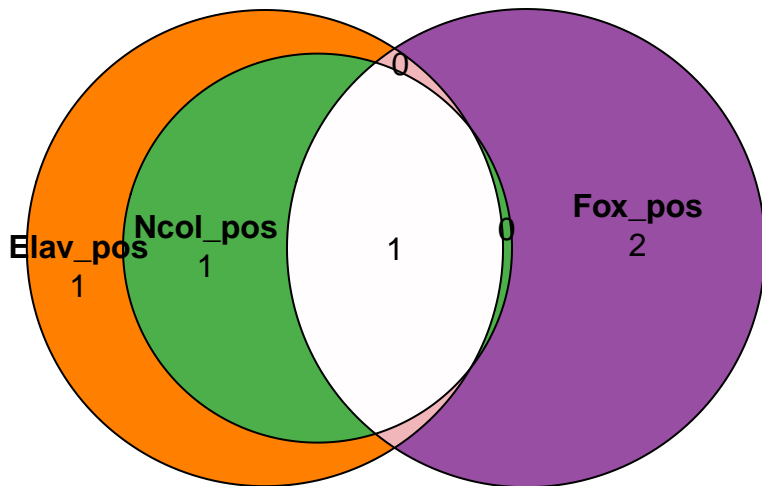


Nvec_vc1.1_XM_032365565.2

RRM:like:AP002373/PABPC1/; RRM_1/

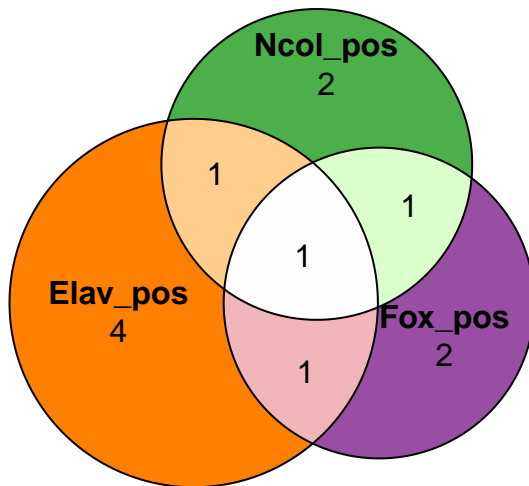


Nvec_vc1.1_XM_001637171.3
Solute carrier family 35,; EamA/TPT/

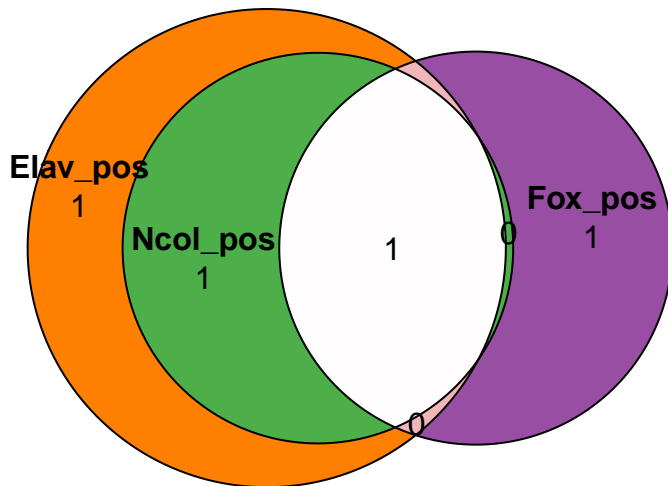


Nvec_vc1.1_XM_032384009.2

Afadin– and alpha–actinin; Tctex–1/ADIP/ADIP/

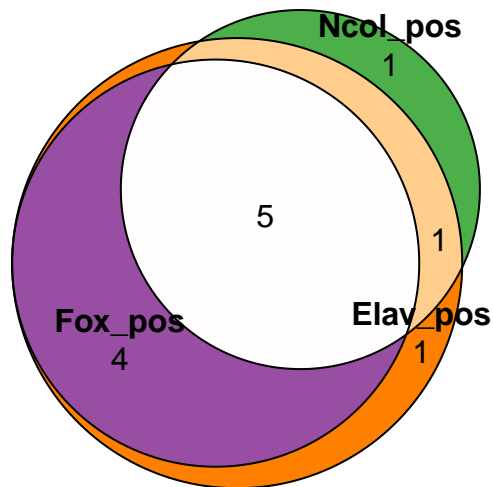


Nvec_vc1.1_XM_048722690.1
SET:SETD3; SET.HG7.3:SETD3



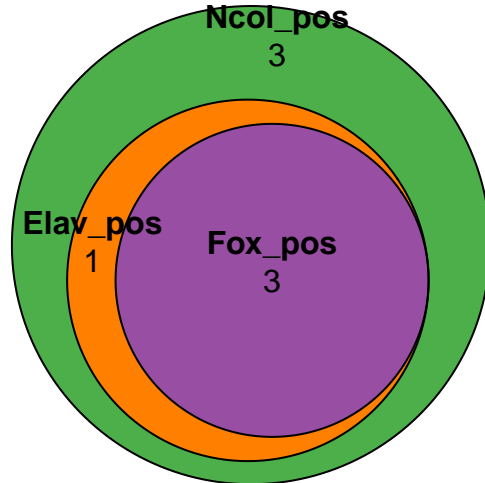
Nvec_vc1.1_XM_001626746.3

Protein THEM6; 4HBT_2/



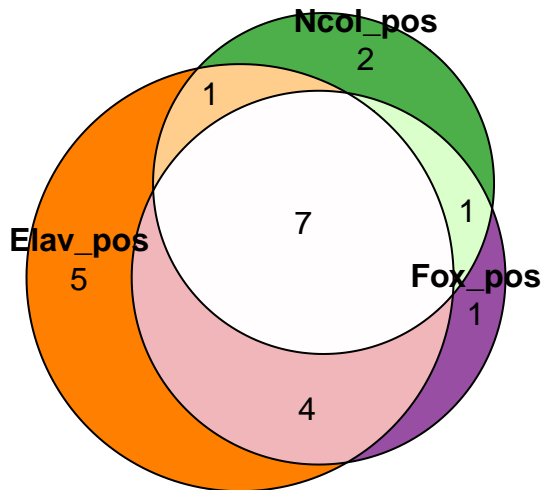
Nvec_vc1.1_XM_048732495.1

Prolow-density lipoprotei; Ldl_recept_a/Ldl_recept_a



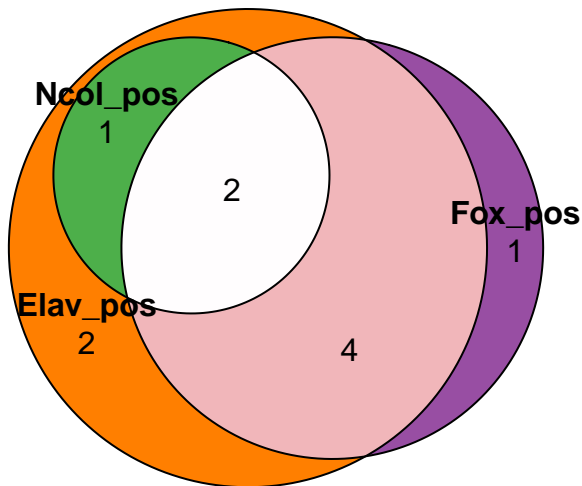
Nvec_vc1.1_XM_048723913.1

FKTN protein; LicD/

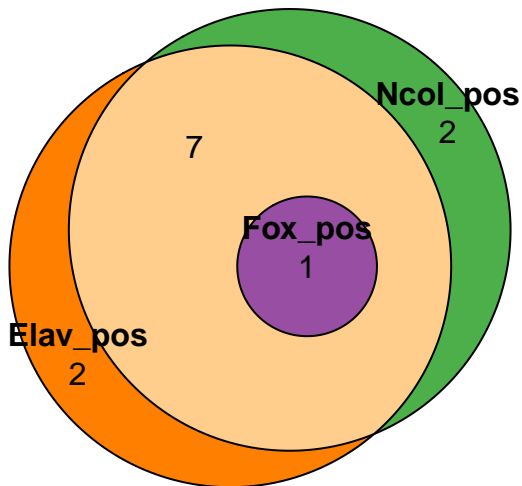


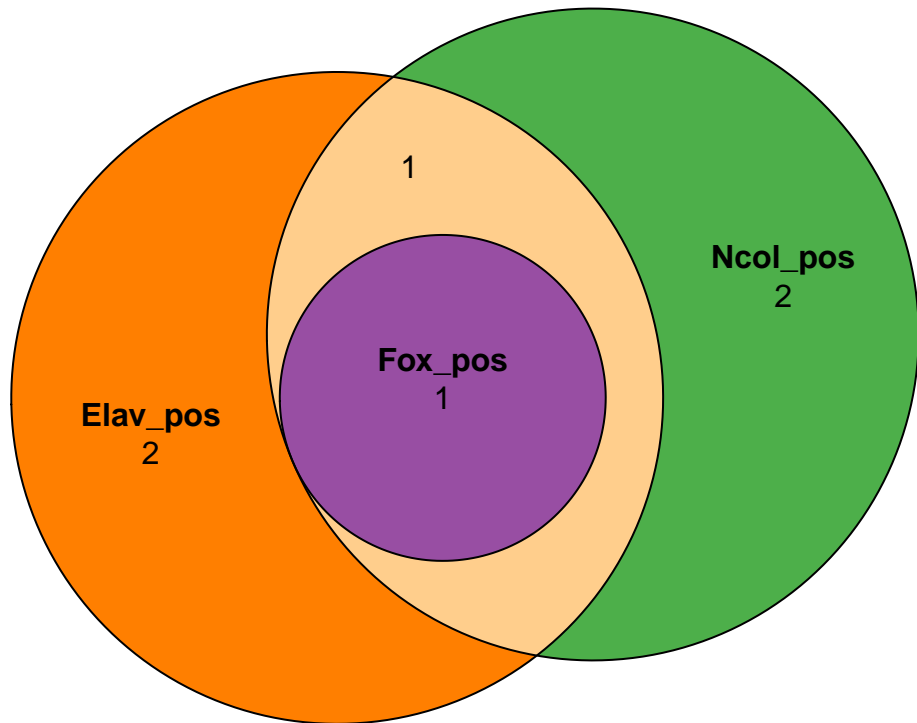
Nvec_vc1.1_XM_048727213.1

Beta-glucuronidase; Glyco_hydro_2_N/Glyco_hyd

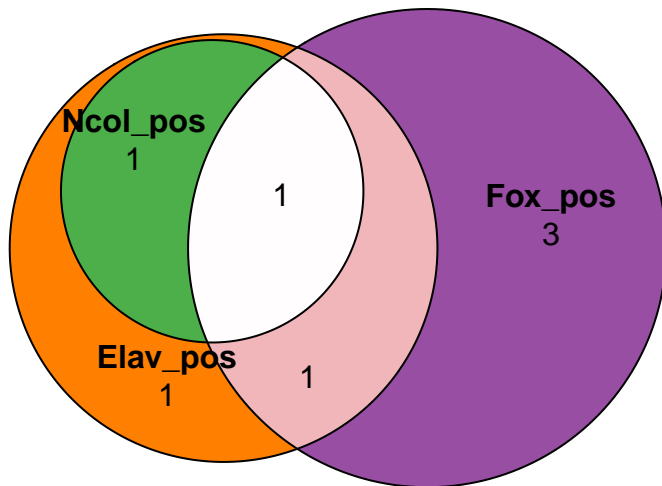


Nvec_vc1.1_XM_032380282.2
XK-related protein; XK-related/

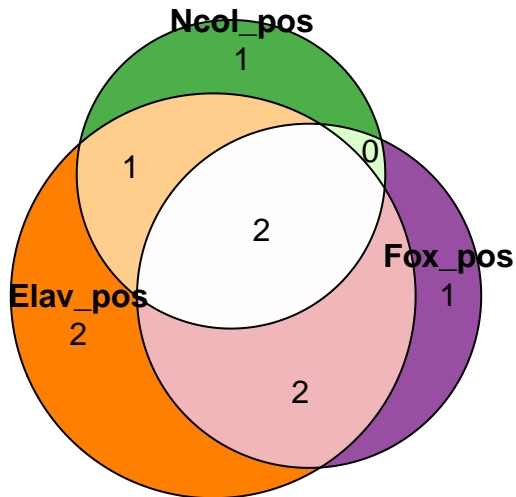




Nvec_vc1.1_XM_001630495.3
Signal peptidase complex ; SPC12/

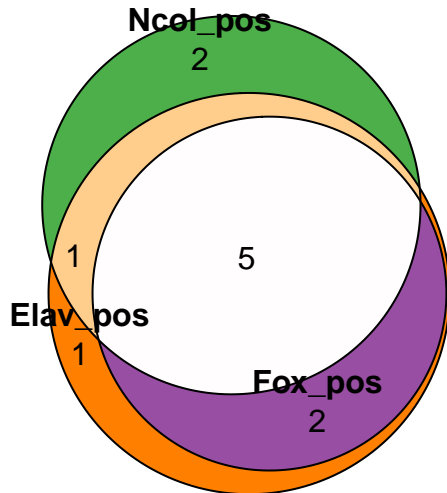


Nvec_vc1.1_XM_032365707.2
Epididymis secretory prot; PBP/



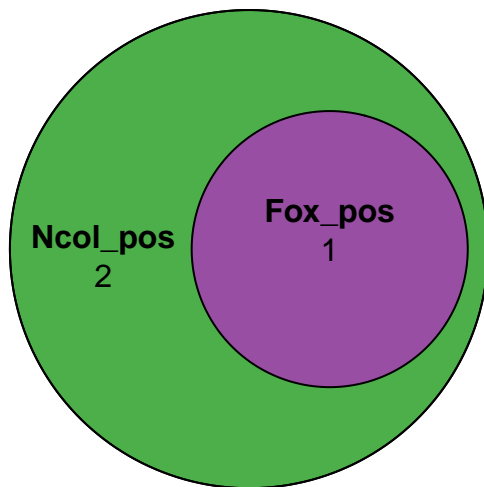
Nvec_vc1.1_XM_048719807.1

;



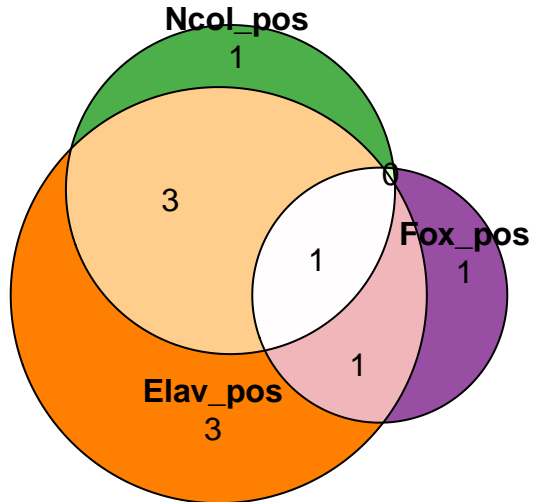
Nvec_vc1.1_XM_048723796.1

Ciliogenesis-associated T;



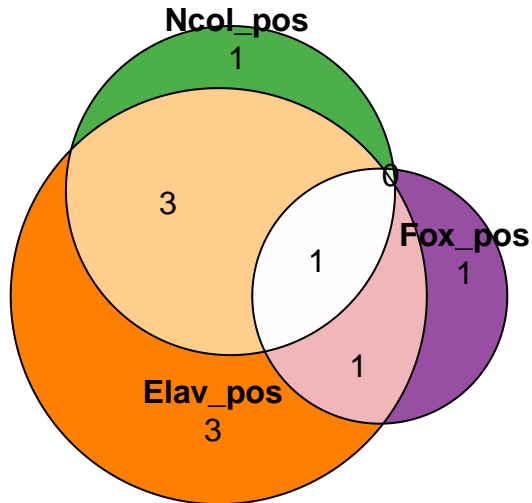
Nvec_vc1.1_XM_048722975.1

;



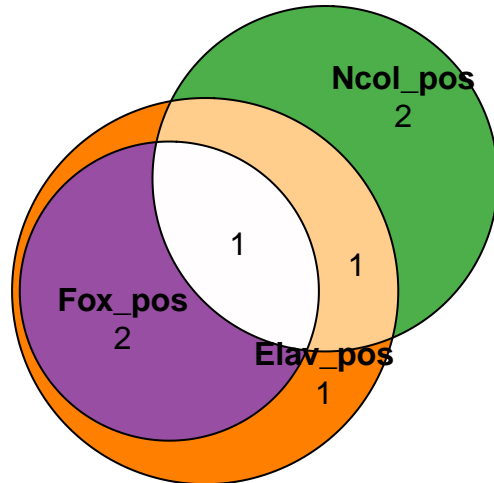
Nvec_vc1.1_XM_048722976.1

Two pore segment channel ;



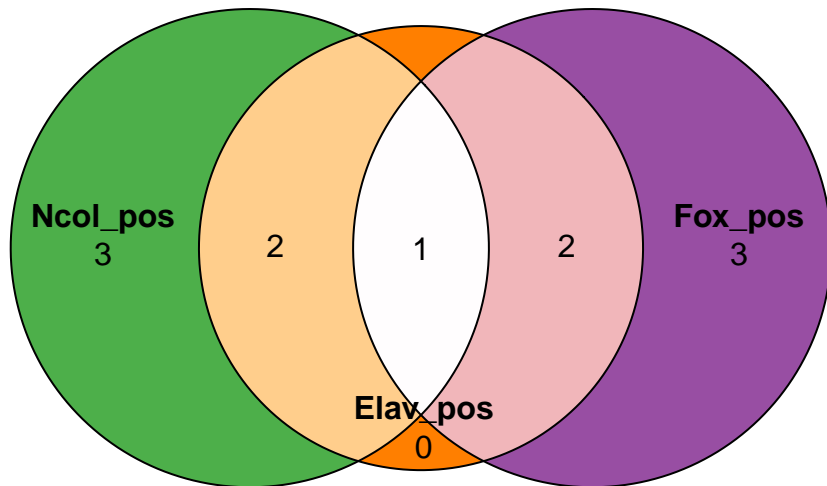
Nvec_vc1.1_XM_032377267.2

;



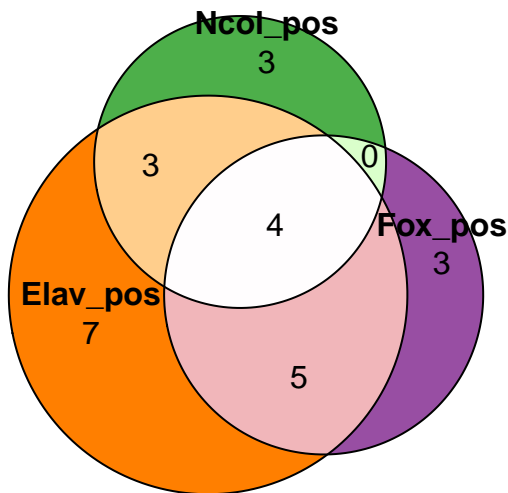
Nvec_vc1.1_XM_032371194.2

Uncharacterized protein D;



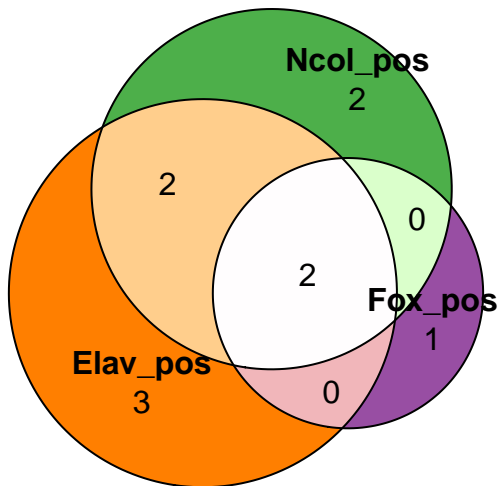
Nvec_vc1.1_XM_032379786.2

F-BAR and double SH3 doma; FCH/SH3_1/SH3_9/



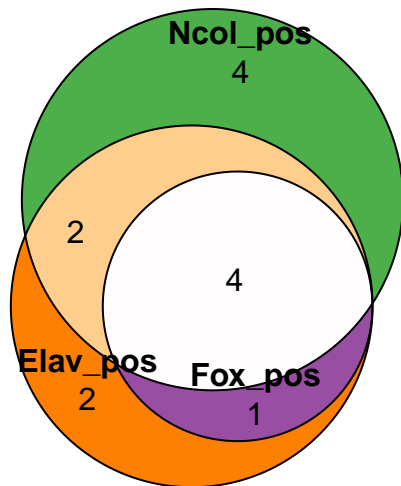
Nvec_vc1.1_XM_032373331.2

;



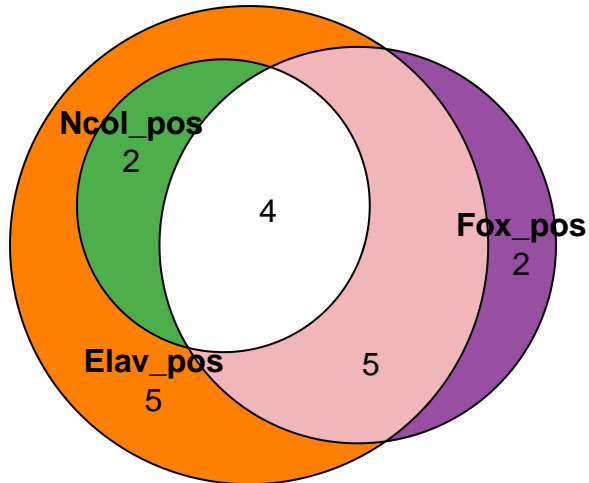
Nvec_vc1.1_XM_048723910.1

FAS-associated factor 1; UBX/

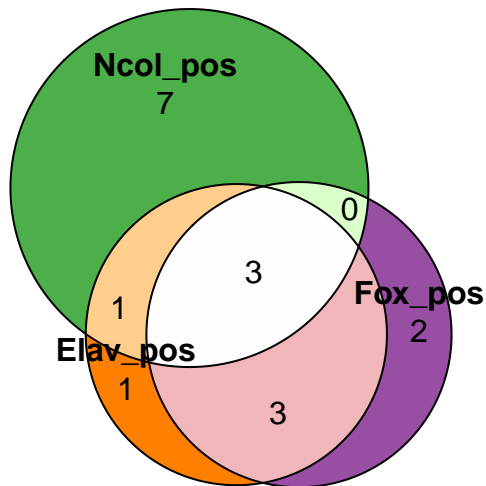


Nvec_vc1.1_XM_032387016.2

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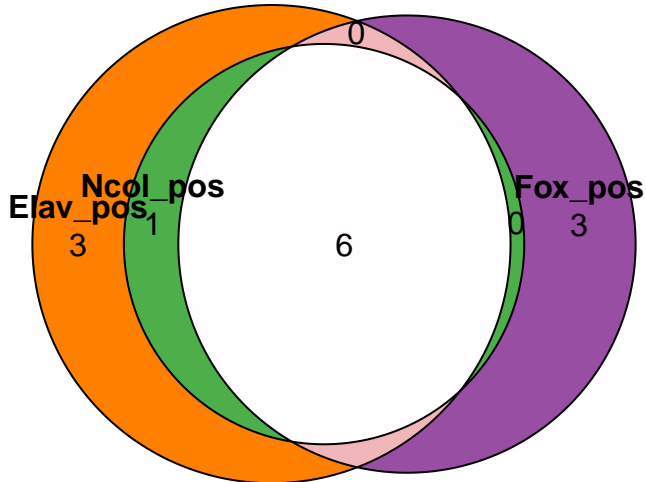


Nvec_vc1.1_XM_032387198.2
NIPA-like protein 2; Mg_trans_NIPA/



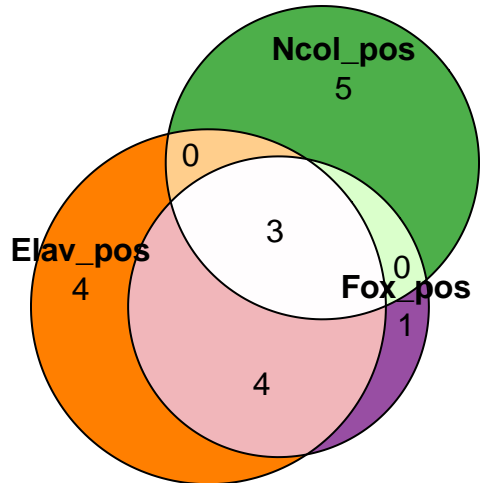
Nvec_vc1.1_XM_001632239.3

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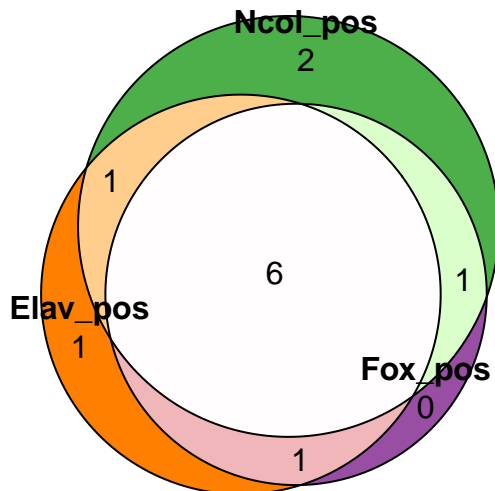
Nvec_vc1.1_XM_048728888.1

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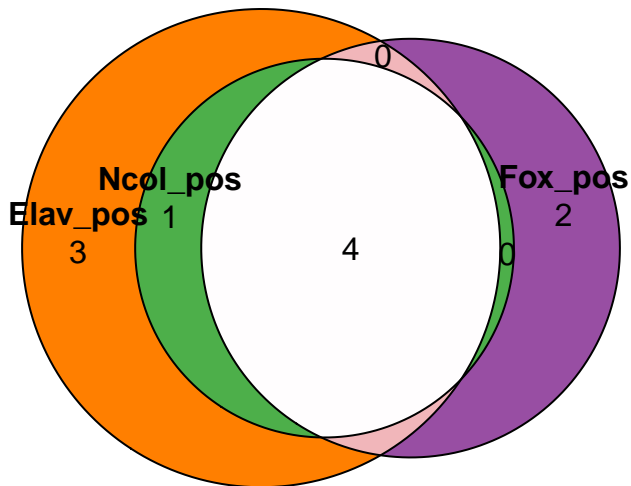
Nvec_vc1.1_XM_032377344.2

SAM:DDHD2/SEC23IP; SAM.HG7.1:DDHD2/SEC23IP



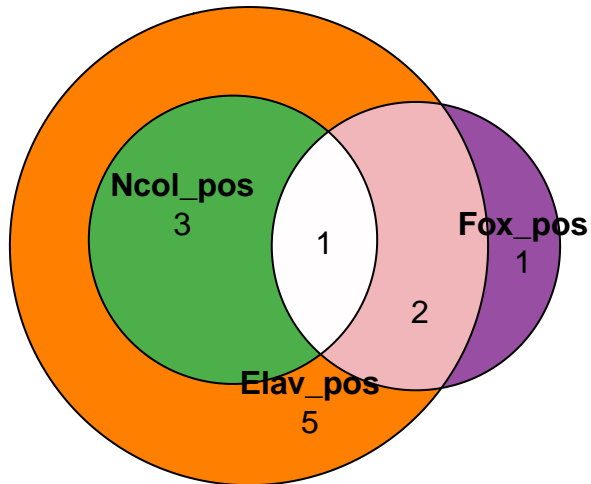
Nvec_vc1.1_XM_032370882.2

Non-lysosomal glucosylcer; GBA2_N/DUF608/



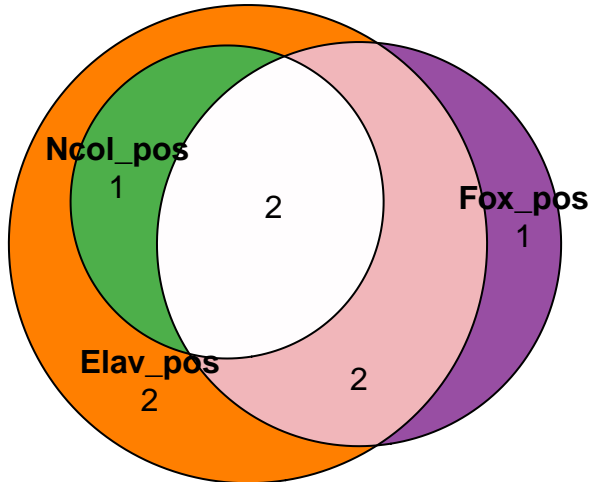
Nvec_vc1.1_XM_032383039.2

Potassium voltage-gated c;



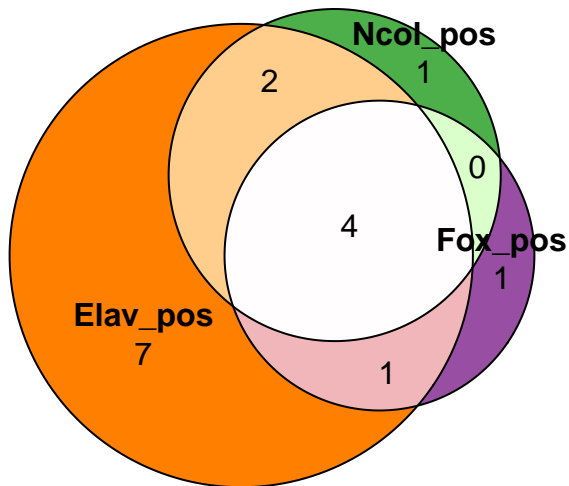
Nvec_vc1.1_XM_048724132.1

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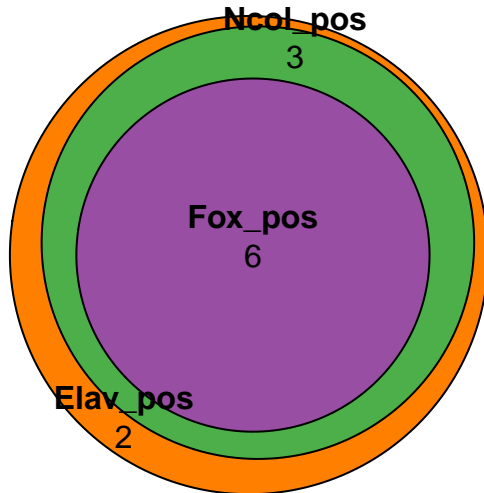
Nvec_vc1.1_XM_001641722.3

N(4)–(beta–N–acetylglucos; Asparaginase_2/



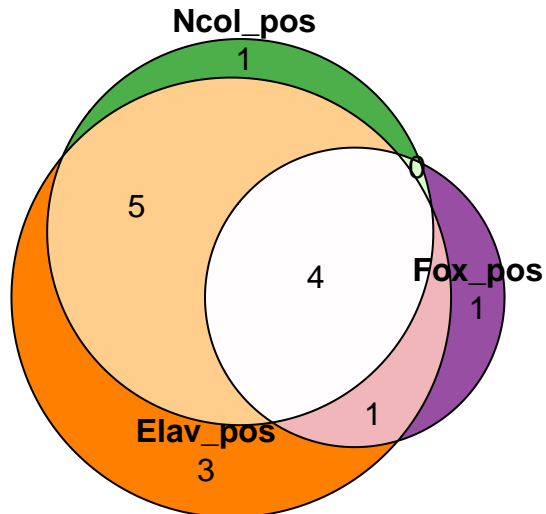
Nvec_vc1.1_XM_032380857.2

Receptor-type tyrosine-pr; Y_phosphatase/



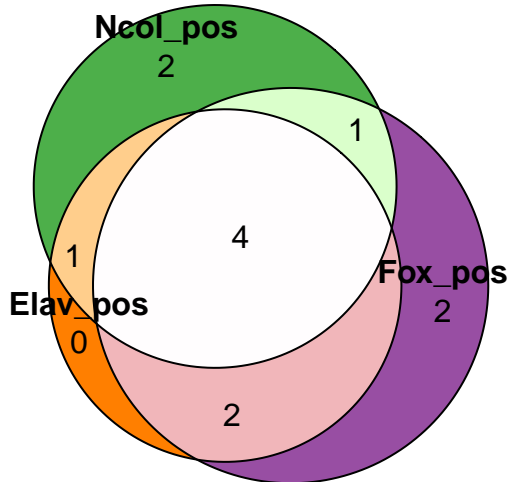
Nvec_vc1.1_XM_032376533.2

E3 ubiquitin-protein liga; PA/zf-RING_2/



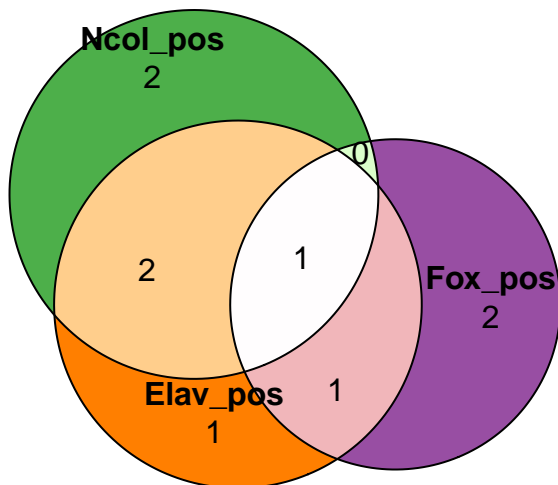
Nvec_vc1.1_XM_032366290.2

Islet cell autoantigen 1,; Arfaptin/ICA69/



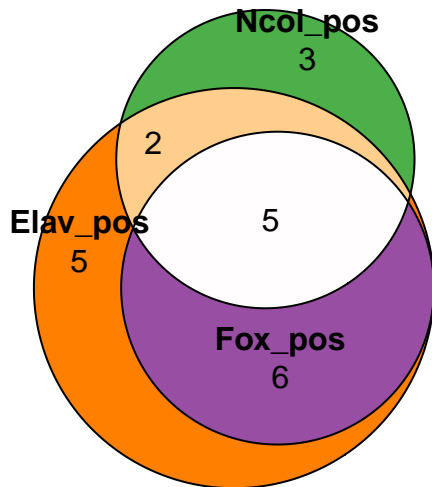
Nvec_vc1.1_XM_001639626.3

Protein SPT2 homolog; SPT2/



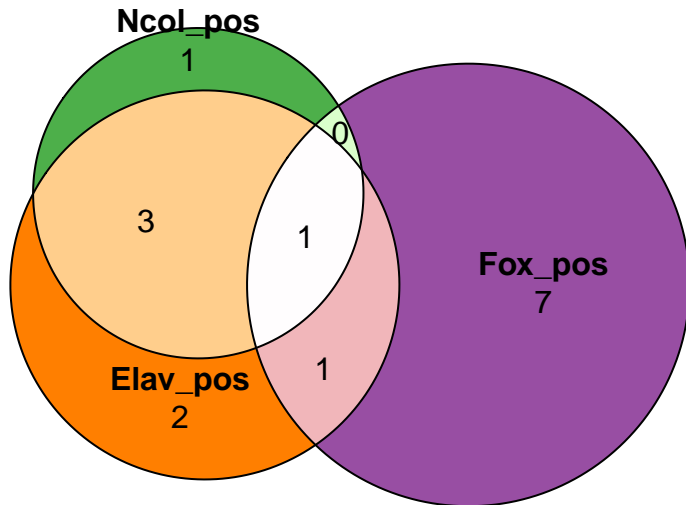
Nvec_vc1.1_XM_032366824.2

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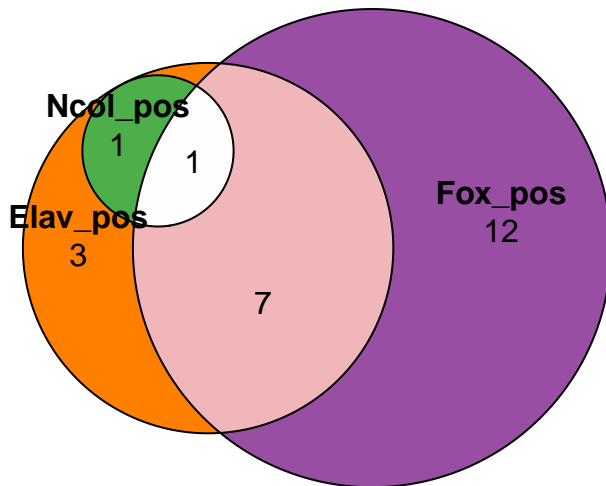
Nvec_vc1.1_XM_001631500.3

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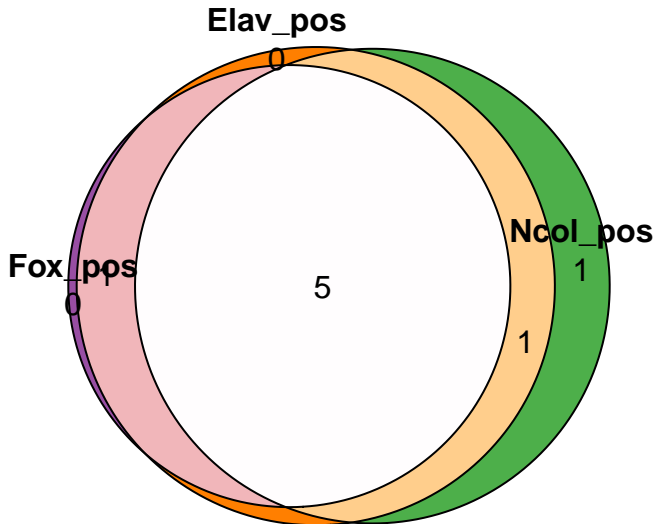


Nvec_vc1.1_XM_032383213.2

Alanyl (Membrane) aminope; Peptidase_M1/ERAP1_C/

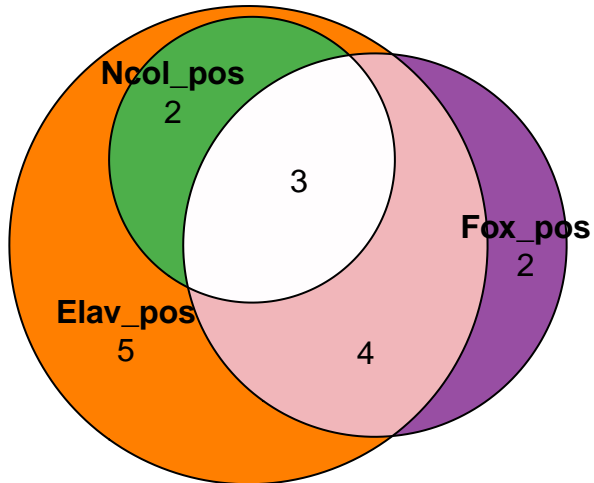


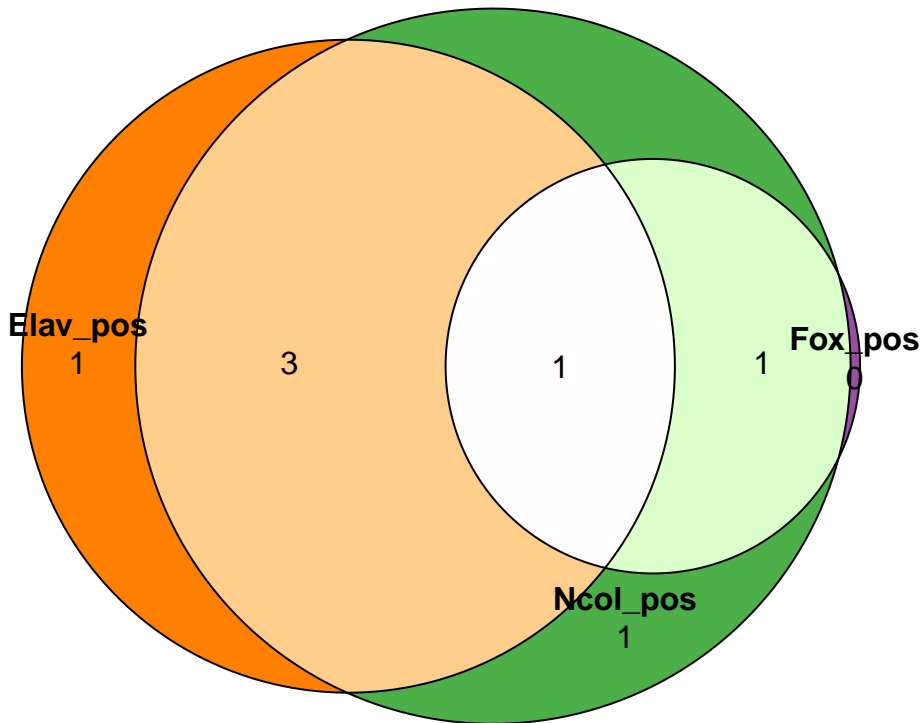
Nvec_vc1.1_XM_001639895.3
Hyaluronidase; Glyco_hydro_56/



Nvec_vc1.1_XM_048722735.1

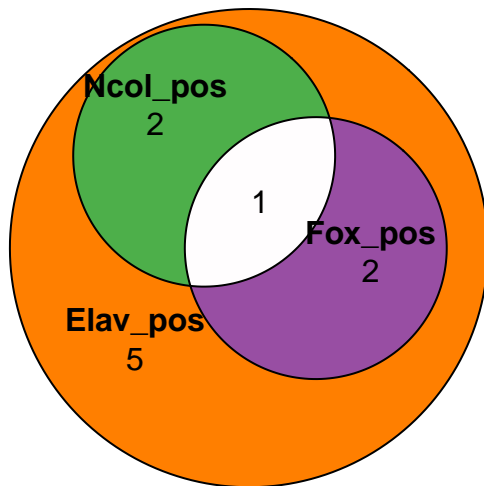
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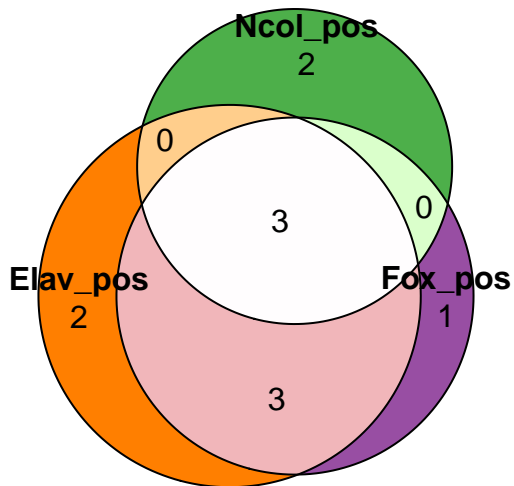
Nvec_vc1.1_XM_001632034.3

Zinc transporter ZIP11; Zip/



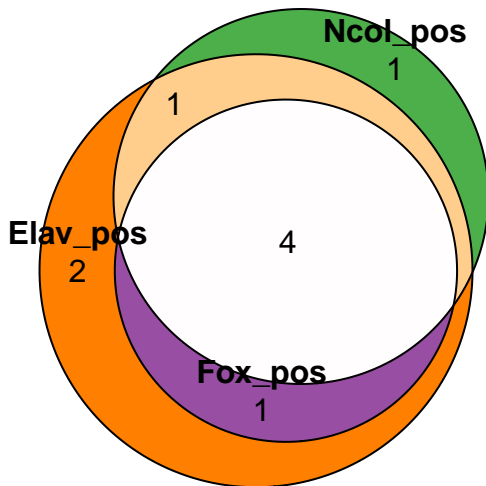
Nvec_vc1.1_XM_048732867.1

Selenoprotein T; Rdx/



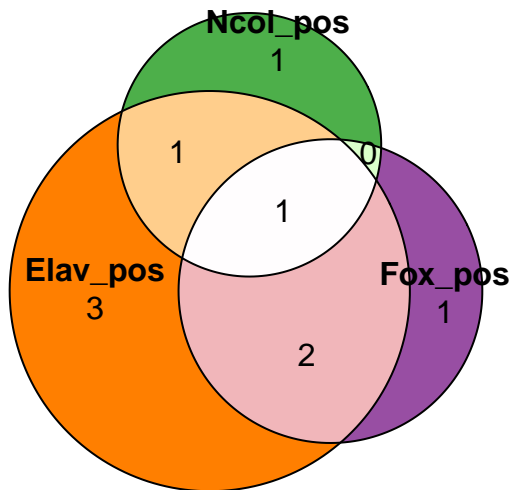
Nvec_vc1.1_XM_048720590.1

Synaptotagmin:PIK3C2A/PIK; Synaptotagmin.HG2.3:PIK3C



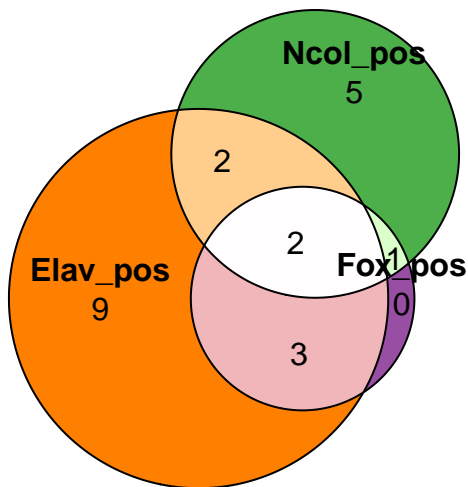
Nvec_vc1.1_XM_048726128.1

Transmembrane protein 14C; Tmemb_14/

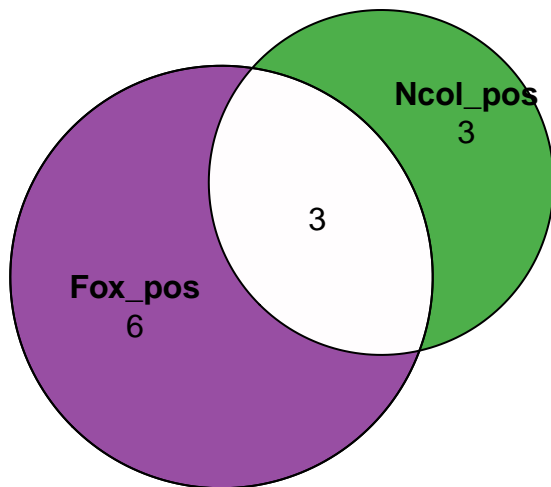


Nvec_vc1.1_XM_032384401.2

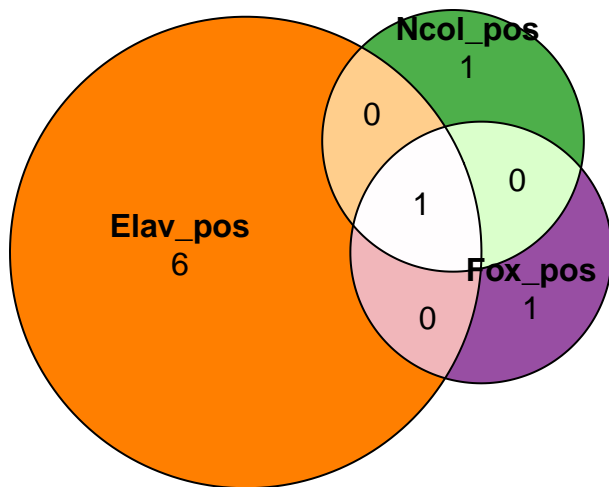
GKAP:like:DLGAP1/DLGAP2/D; GKAP/



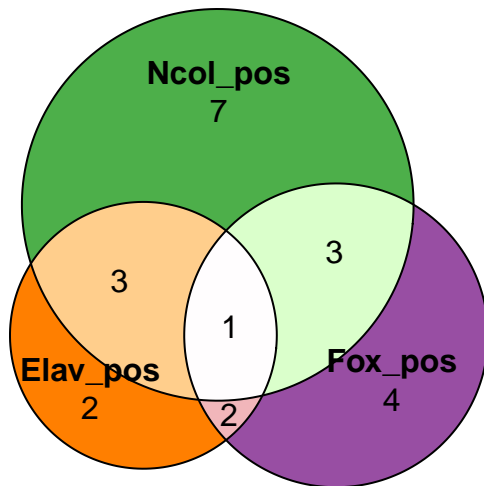
Nvec_vc1.1_XM_001626886.3
Selenoprotein S; Selenoprotein_S/



Nvec_vc1.1_XM_032373753.2
cDNA FLJ51906, highly sim; HSP20/

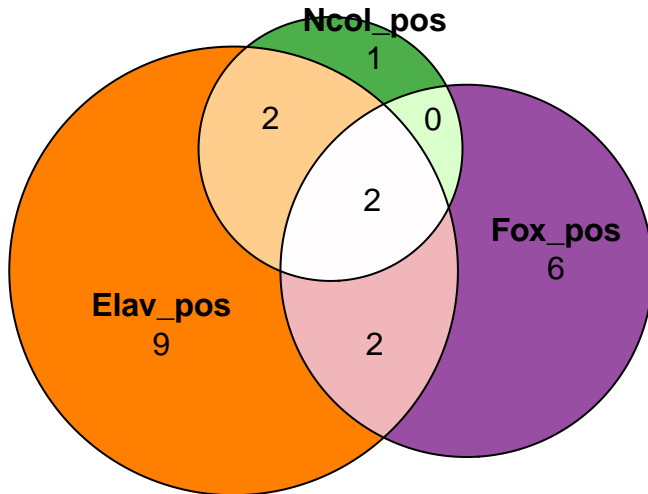


Nvec_vc1.1_XM_032373261.2
cDNA, FLJ94988, highly si; EF-hand_7/



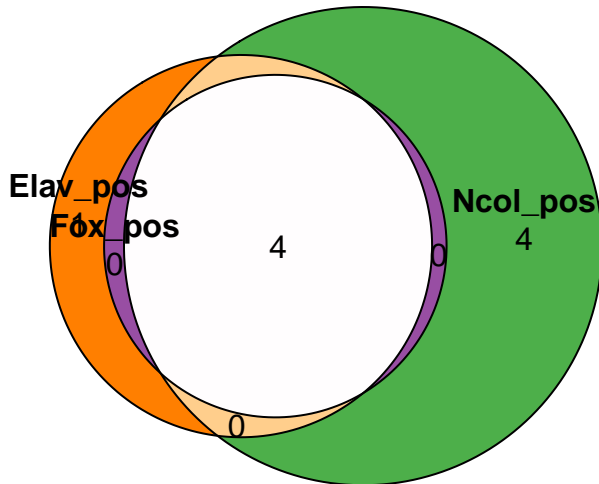
Nvec_vc1.1_XM_048729836.1

E3 ubiquitin-protein liga;



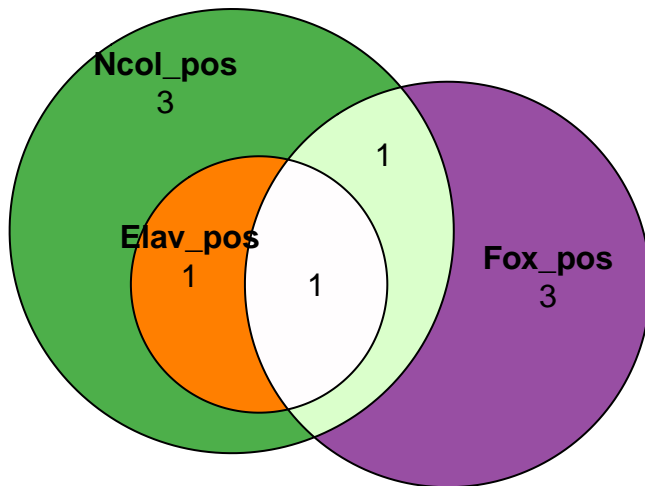
Nvec_vc1.1_XM_048719979.1

Leukocyte receptor cluste; SAC3_GANP/



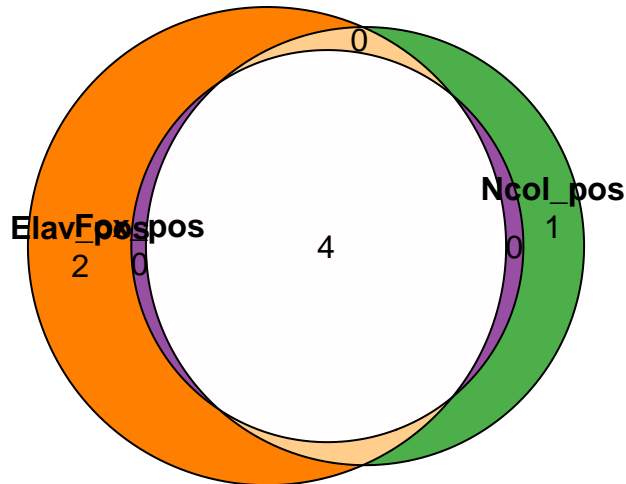
Nvec_vc1.1_XM_048722888.1

Transmembrane protein 79; MAPEG/



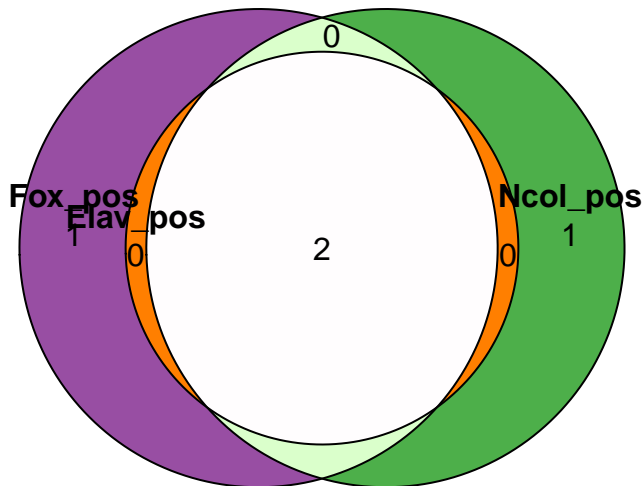
Nvec_vc1.1_XM_048733377.1

WD repeat domain 19; WD40/



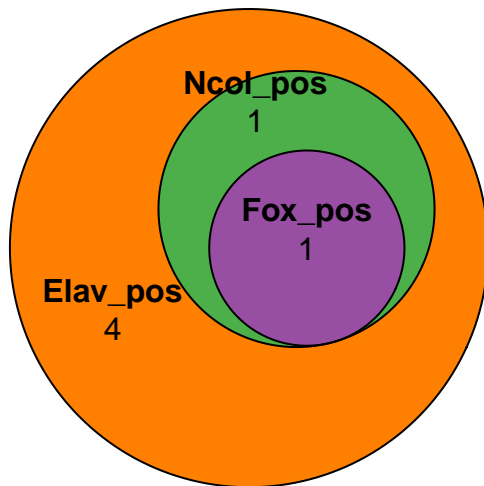
Nvec_vc1.1_XM_001627406.3

Ufm1-specific protease 2; Peptidase_C78/



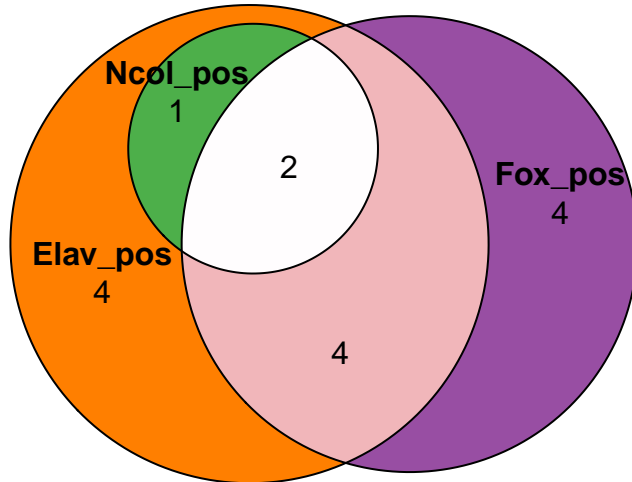
Nvec_vc1.1_XM_048734604.1

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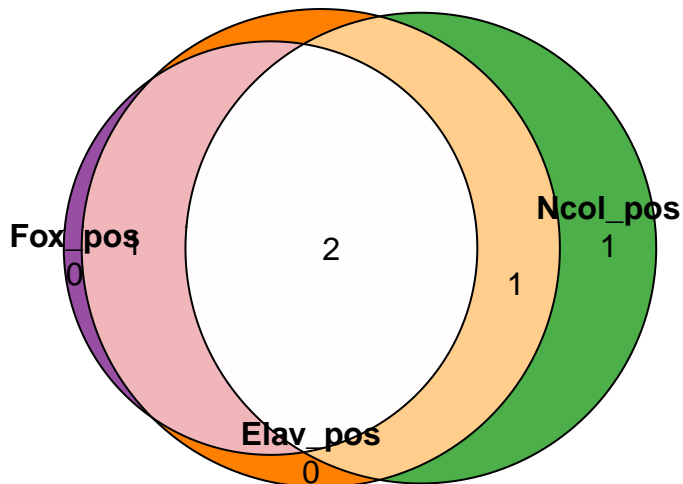
Nvec_vc1.1_XM_001632720.3

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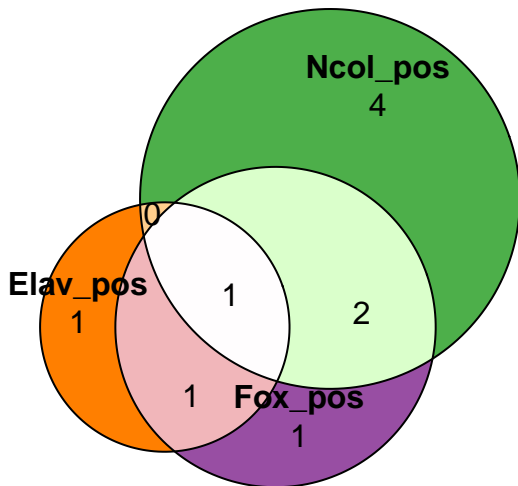
Nvec_vc1.1_XM_048734120.1

DNA annealing helicase an;

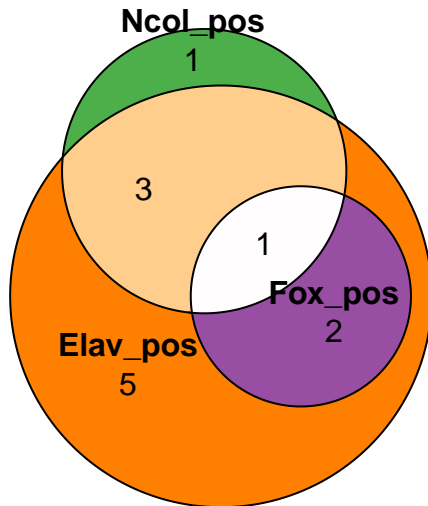


Nvec_vc1.1_XM_032383407.2

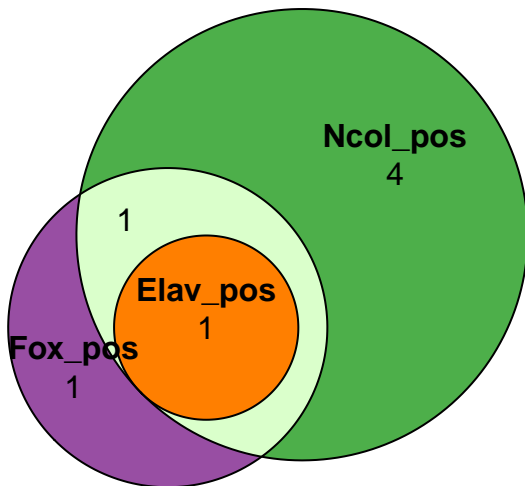
cDNA FLJ51019, highly sim; Ammonium_transp/



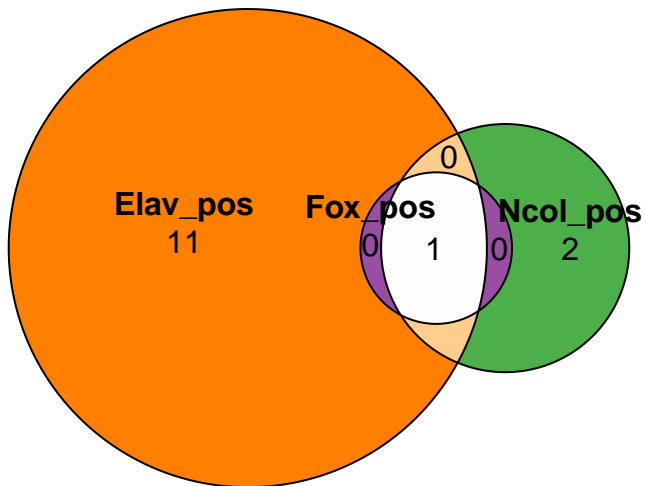
Nvec_vc1.1_XM_032373921.2
; BNIP3/BNIP3/



Nvec_vc1.1_XM_032384411.2
cDNA FLJ90515 fis, clone ; Zip/

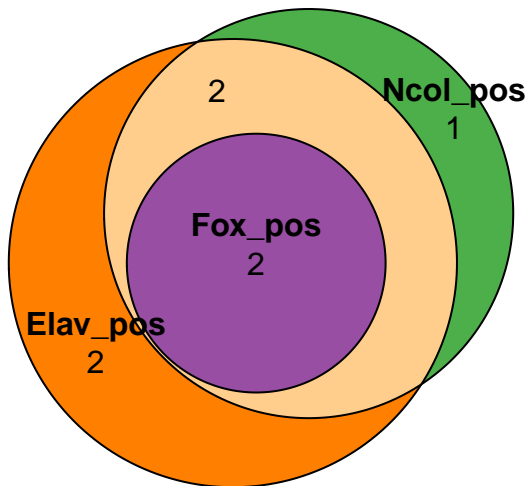


Nvec_vc1.1_XM_048722812.1
Tub.HG1.2:TUB/TULP3; TULP3



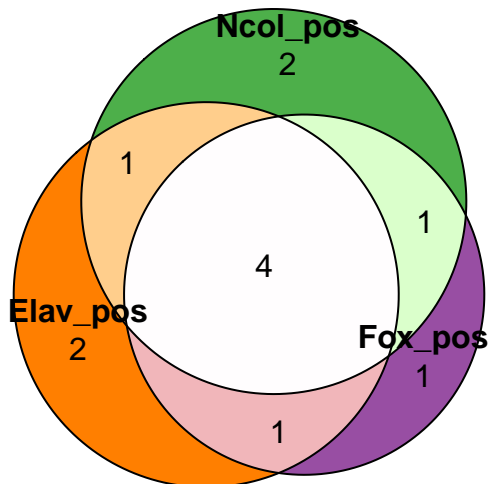
Nvec_vc1.1_XM_048732912.1

TTF1; Myb.HG6.1:TTF1



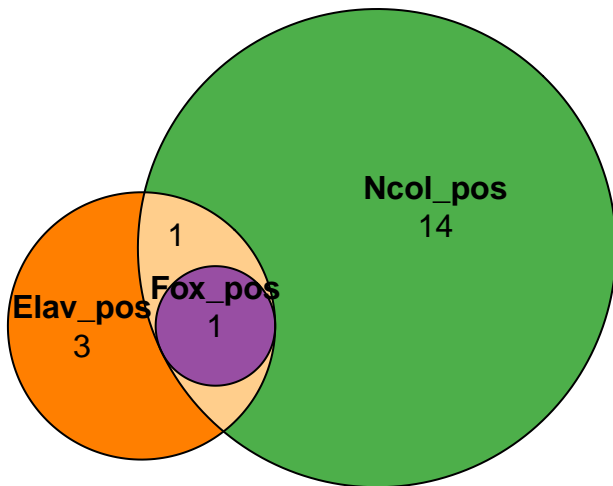
Nvec_vc1.1_XM_001635459.3

RasGAP:RASA2/RASA3; RasGAP.HG1.1:RASA2/RASA3

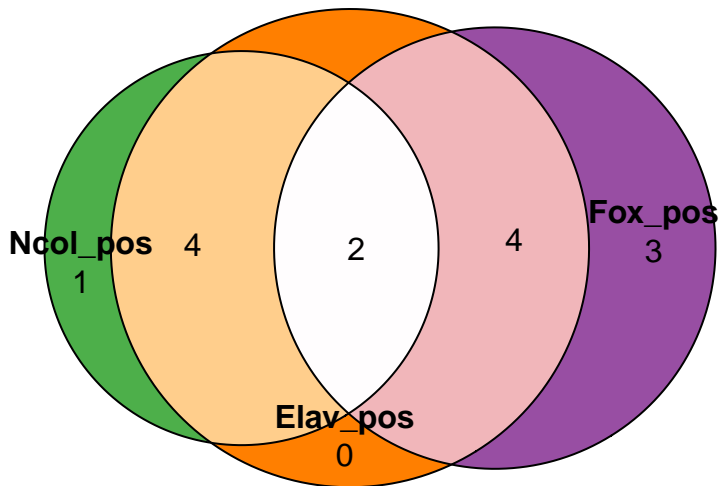


Nvec_vc1.1_XM_032371040.2

ion-NaKCa:like:KCND1/KCND; BTB_2/Ion_trans/

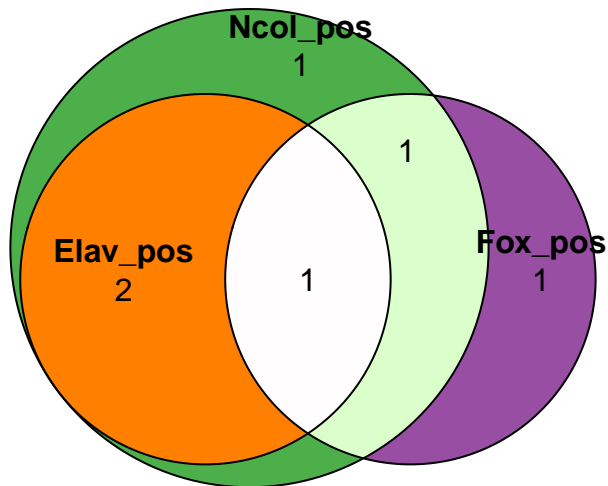


Nvec_vc1.1_XM_032372495.2
Tripartite motif-containi; zf-RING_5/



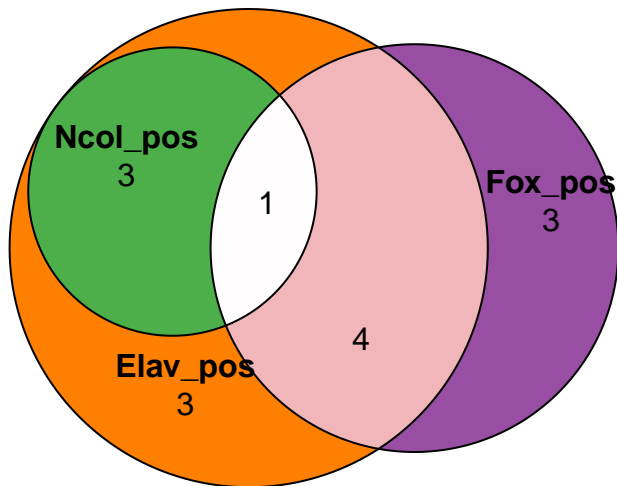
Nvec_vc1.1_XM_048719781.1

WW domain-binding protein; GRAM/WWbp/



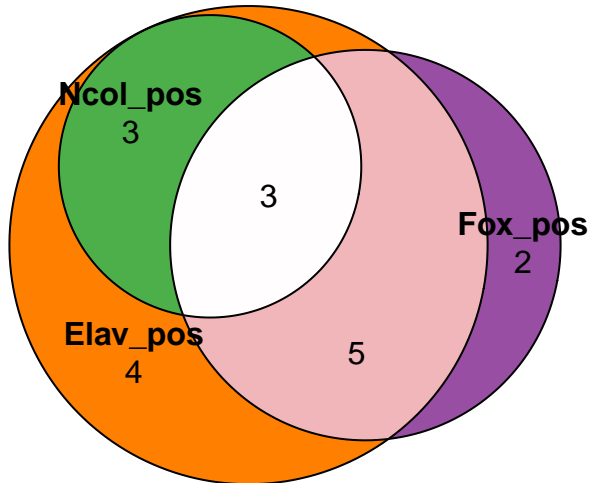
Nvec_v1g232650

NADH dehydrogenase (Ubiqu; NADH_B2/

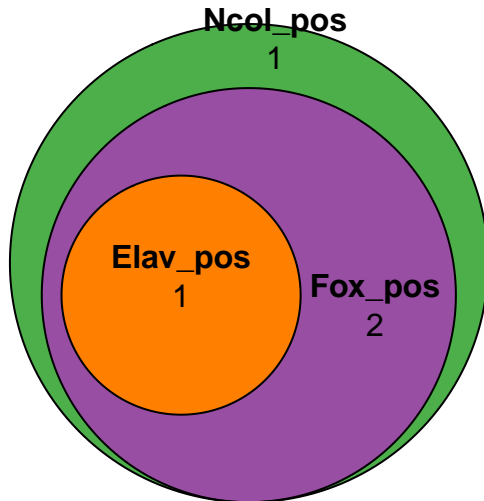


Nvec_vc1.1_XM_048722989.1

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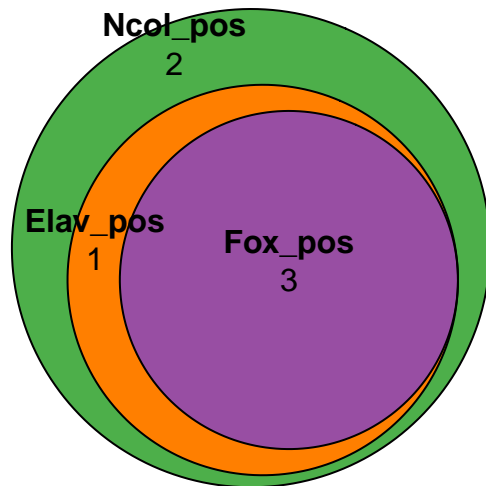


Nvec_vc1.1_XM_001623641.3
Gem-associated protein 6; Gemin6/

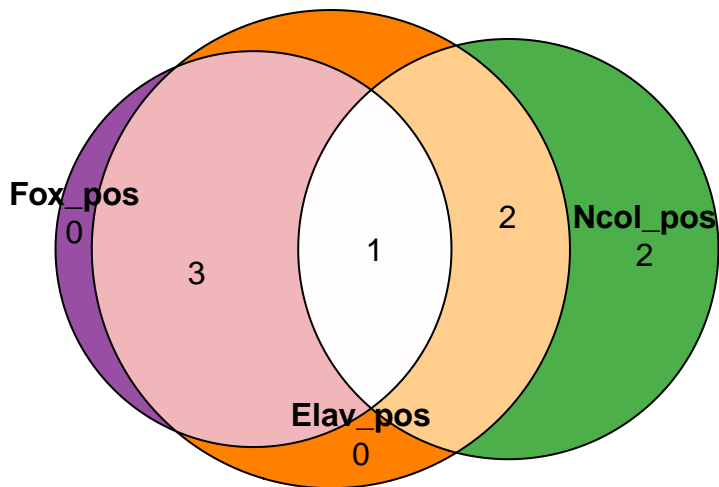


Nvec_vc1.1_XM_032387376.2

Cysteine protease ATG4B; Peptidase_C54/

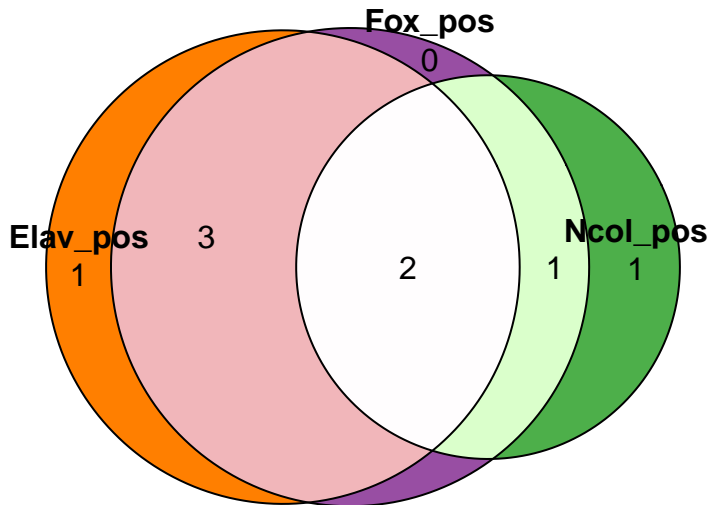


Nvec_vc1.1_XM_032377206.2
DCN1-like protein 3; Cullin_binding/



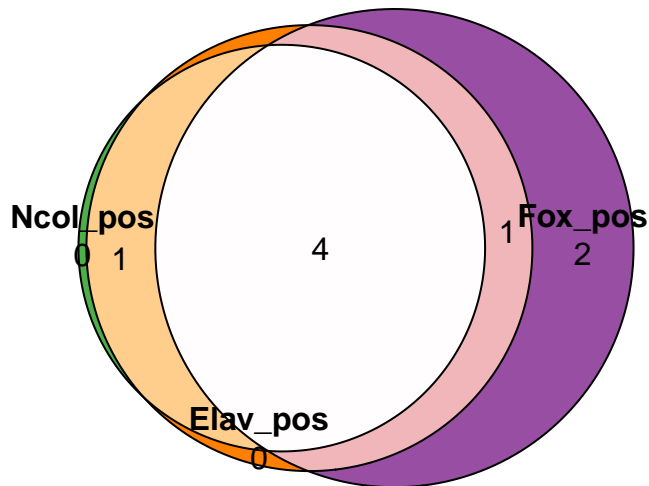
Nvec_vc1.1_XM_048730893.1

cDNA FLJ31473 fis, clone ; EST1/EST1_DNA_bind/



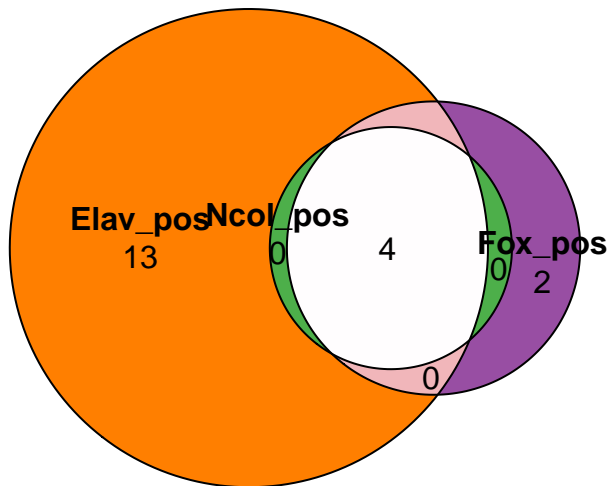
Nvec_vc1.1_XM_001639496.3

CDW9/WDR51B; WD40/WD40/WD40/WD40/WD40/



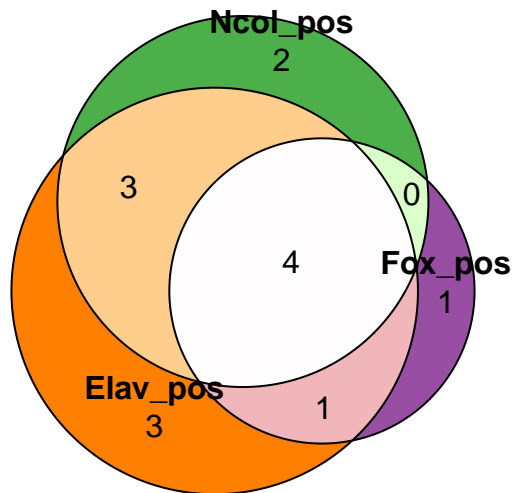
Nvec_vc1.1_XM_032372281.2

TNF receptor-associated f; zf-C3HC4/zf-TRAF/zf-TRAF/



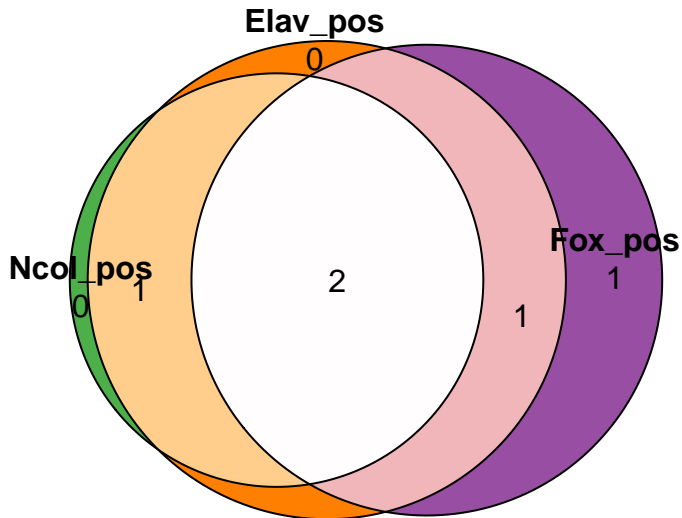
Nvec_vc1.1_XM_001635718.3

WD repeat domain 45; WD40/

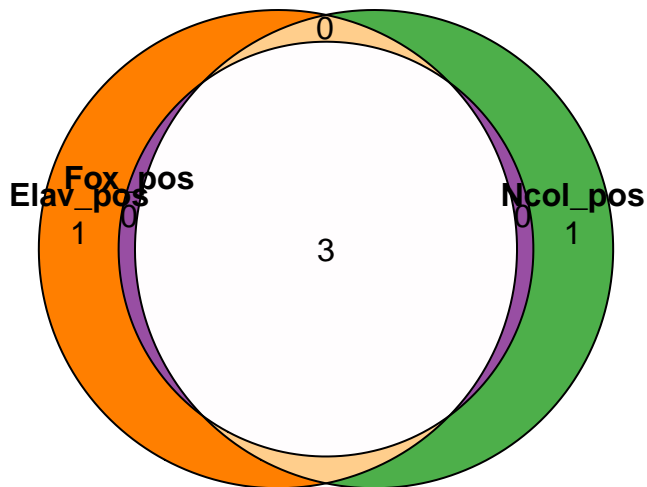


Nvec_vc1.1_XM_032373393.2

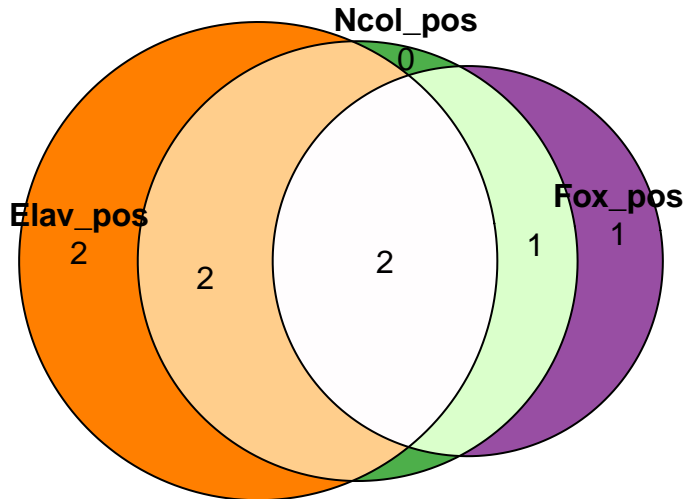
Dynactin subunit 4; Dynactin_p62/Dynactin_p62



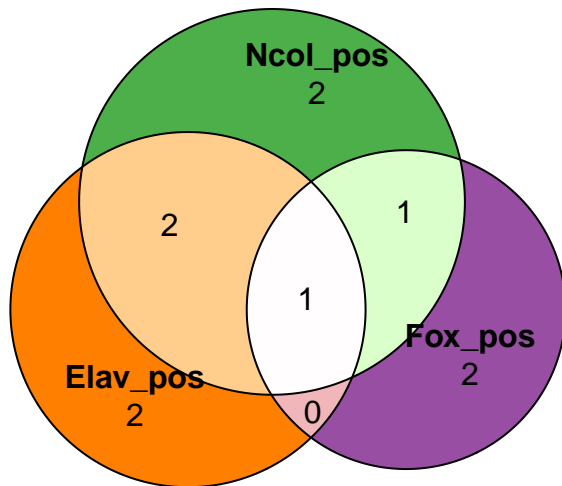
Nvec_vc1.1_XM_032381835.2
; zf-RING_2/



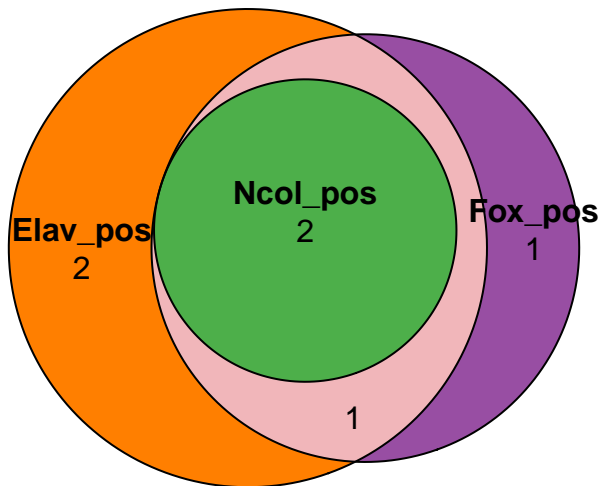
Nvec_vc1.1_XM_001634495.3
; PB1/



Nvec_vc1.1_XM_048724740.1
Translocation protein SEC; Sec62/

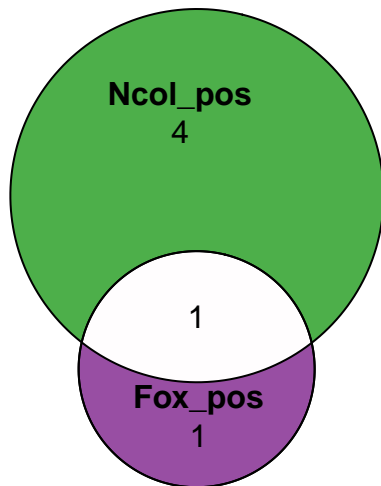


Nvec_vc1.1_XM_032373709.2
cDNA FLJ59118; BNIP2/CRAL_TRIO_2/



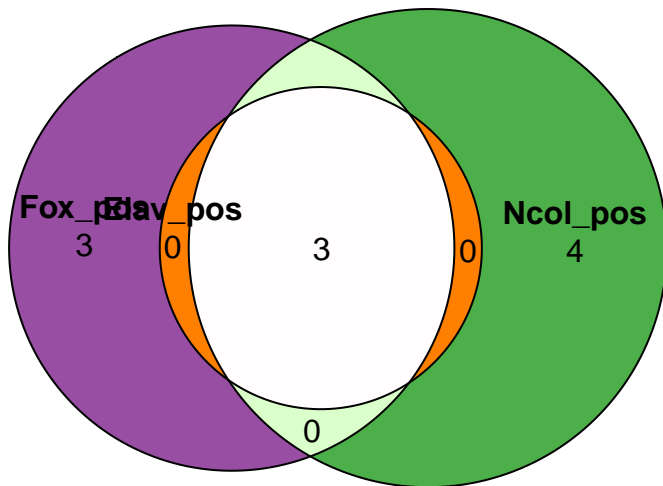
Nvec_vc1.1_XM_032378615.2

;

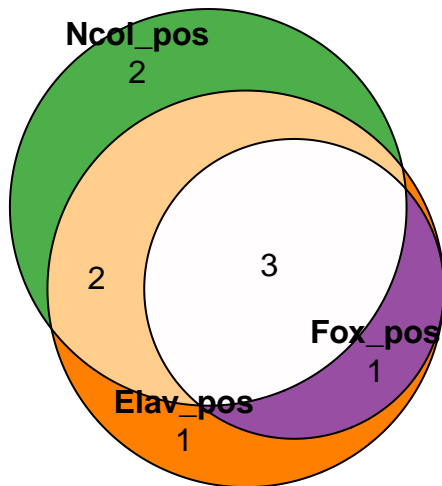


Nvec_vc1.1_XM_048725768.1

;

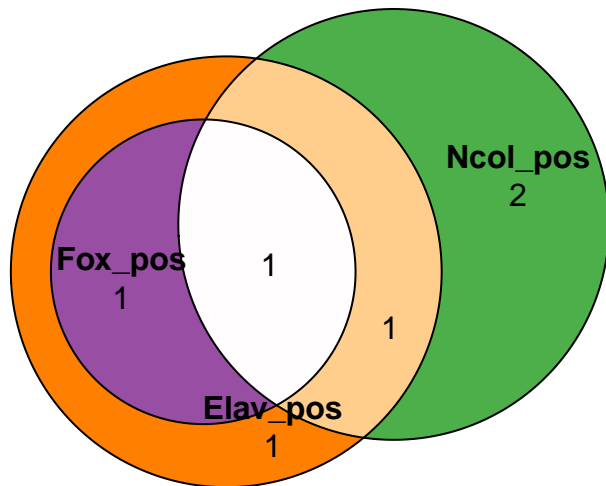


Nvec_vc1.1_XM_048726725.1
Rab GTPase-binding effect; FYVE/



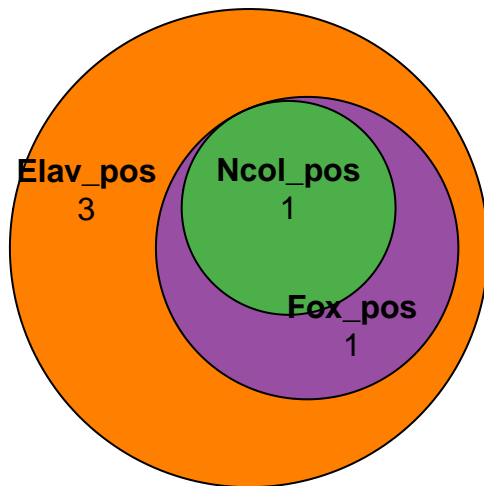
Nvec_vc1.1_XM_048728573.1

Putative uncharacterized ; PDEase_I/



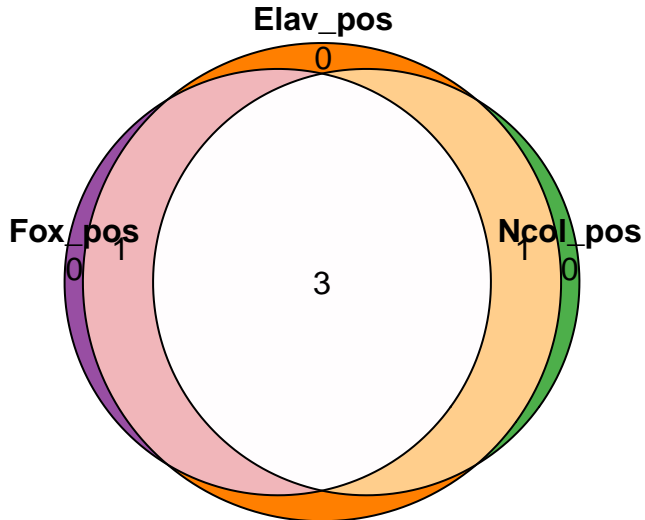
Nvec_vc1.1_XM_032379526.2

Protein FAM35A;

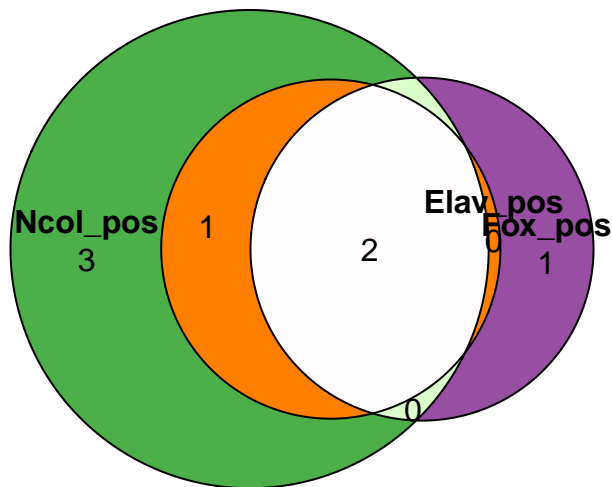


Nvec_NVE4126

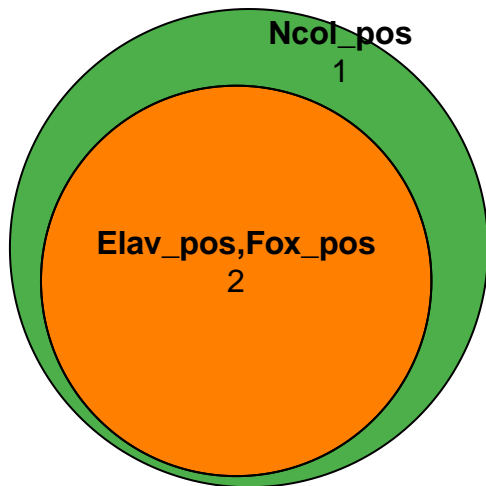
;



Nvec_vc1.1_XM_001621797.3
SIR2:SIRT7; SIR2.HG5.0:SIRT7

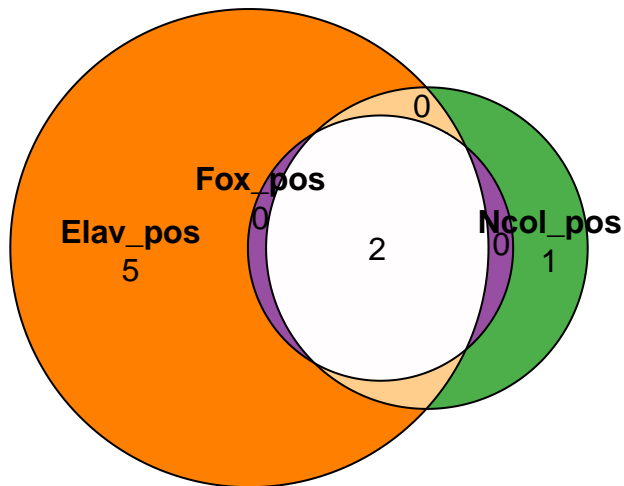


Nvec_vc1.1_XM_001635893.3
cDNA FLJ54041, highly sim; LETM1/



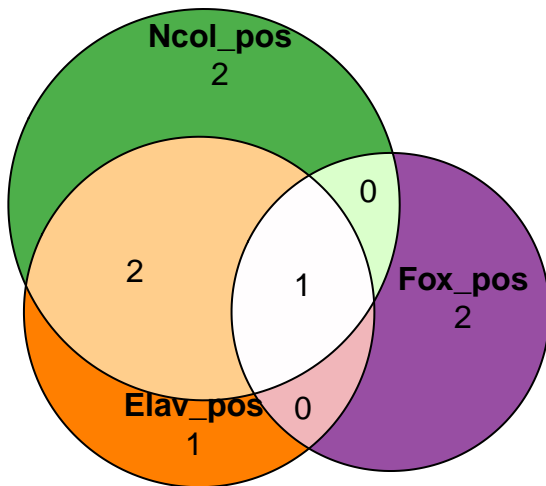
Nvec_vc1.1_XM_048725801.1

Ectopic P granules protei;



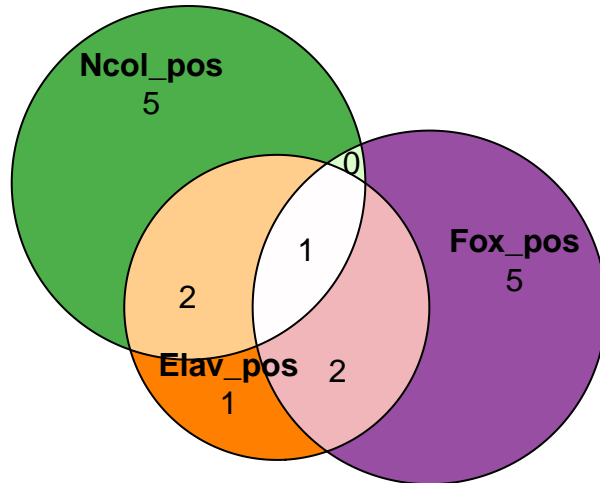
Nvec_vc1.1_XM_032378974.2

E3 ubiquitin-protein liga; TPR_11/TPR_1/zf-RING_2/



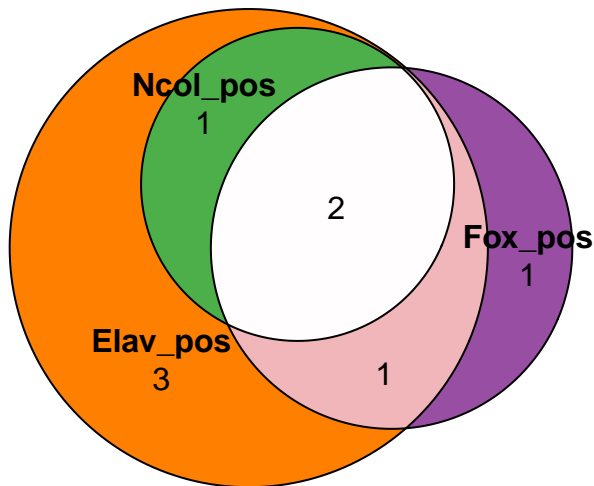
Nvec_vc1.1_XM_001635153.3

;



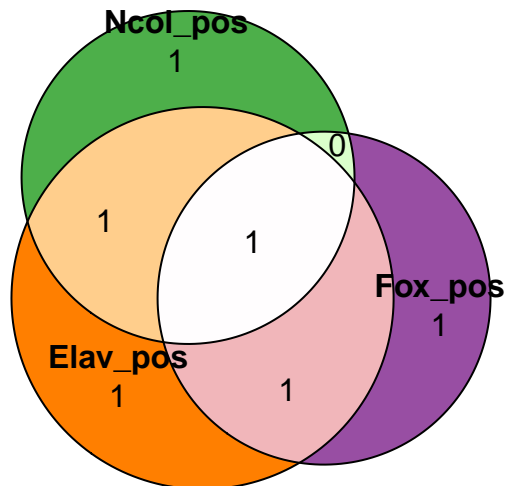
Nvec_vc1.1_XM_001638519.3

CYP:CYP17A1/CYP21A2; CYP.HG1.16:CYP17A1/CYP21A

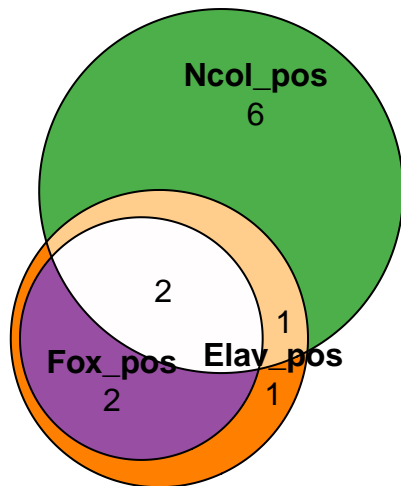


Nvec_vc1.1_XM_001628946.3

bZIP.HG1.19:ATF6/ATF6B; ATF6/ATF6B

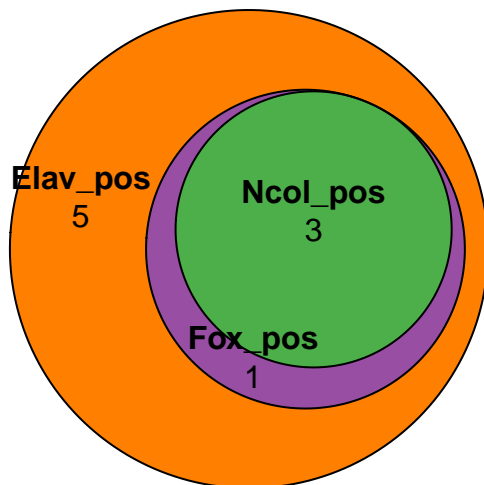


Nvec_vc1.1_XM_048721680.1
Chromatin modification-re; NuA4/

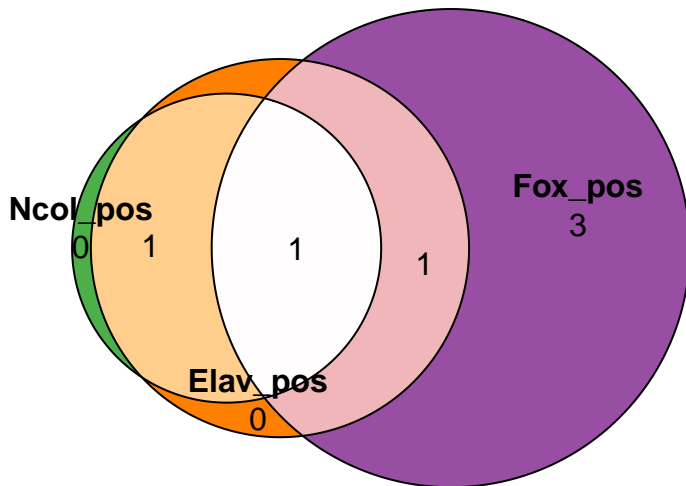


Nvec_vc1.1_XM_032379449.2

RRM:HNRNPR/SYNCRIP; RRM.HG11.1:HNRNPR/SYNCRIP

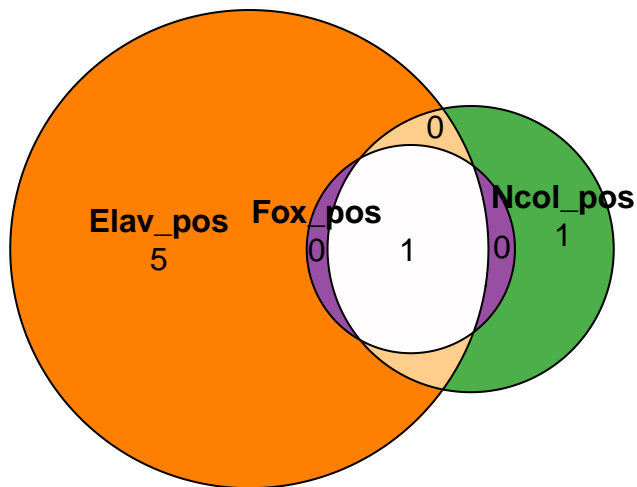


Nvec_vc1.1_XM_001641313.3
RING finger protein 24; zf-RING_2/



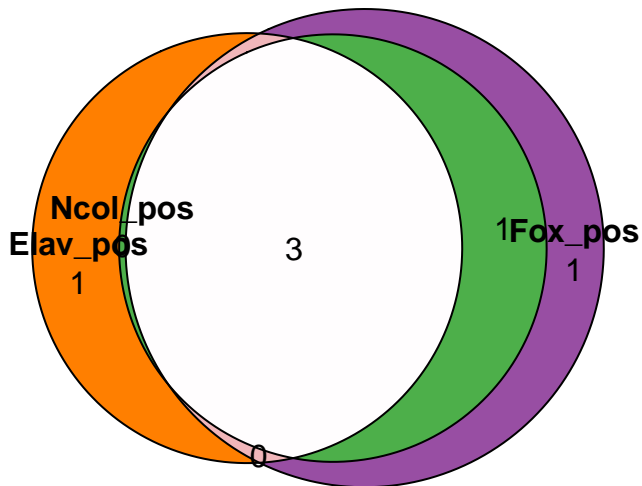
Nvec_vc1.1_XM_032384609.2

Nucleobindin 2; EF-hand_7/

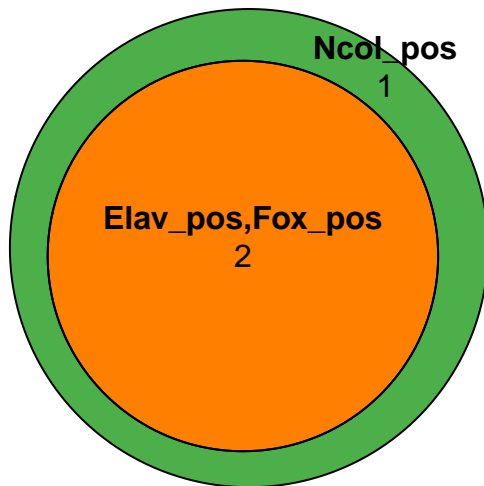


Nvec_vc1.1_XM_001638822.3

HIG1 domain family, membe; HIG_1_N/

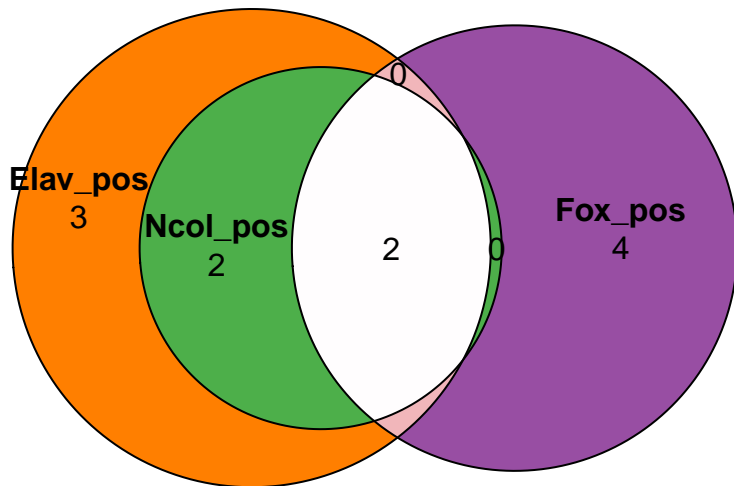


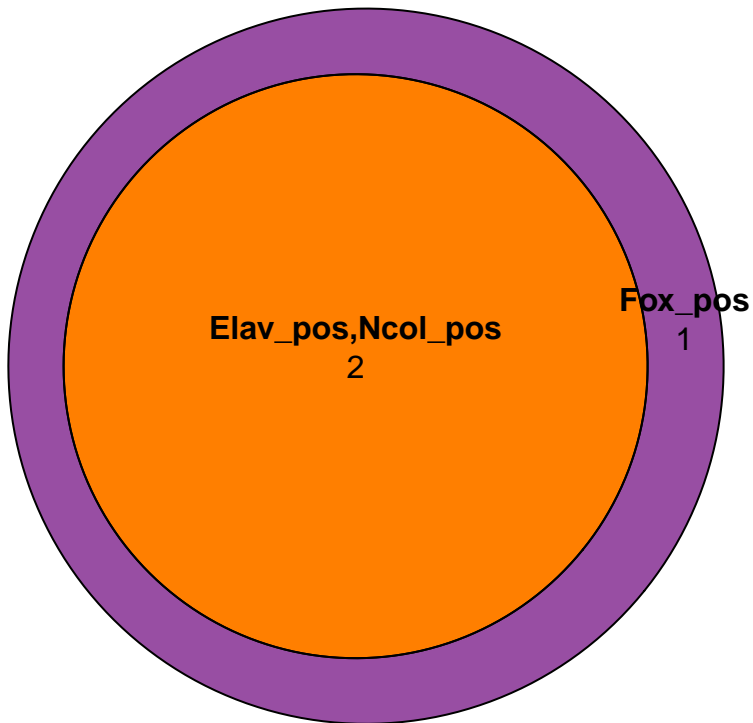
Nvec_vc1.1_XM_048730080.1
Thiopurine S-methyltransf; TPMT/



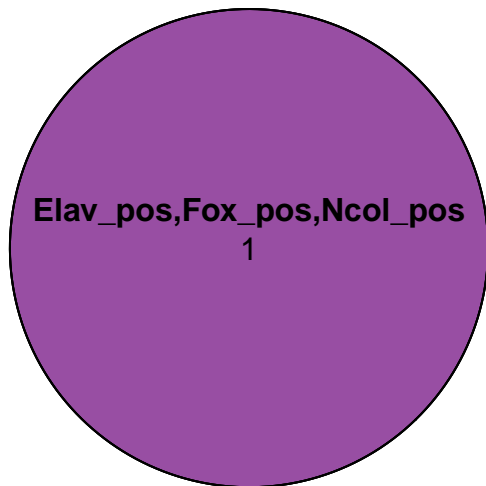
Nvec_vc1.1_XM_032373576.2

Lin-10 homolog (C. elegan; UPF0183/





Nvec_vc1.1_XM_032363015.2
Vesicular, overexpressed ; WBP-1/



Nvec_vc1.1_XM_048729139.1
Epoxide hydrolase 4; Abhydrolase_6/

