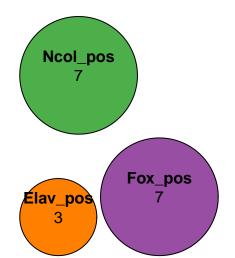
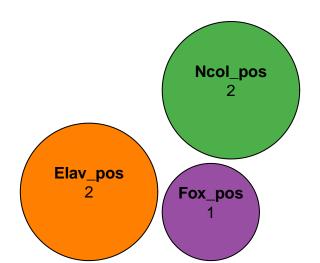
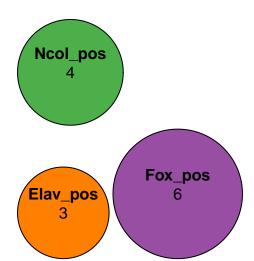
Nvec_vc1.1_XM_032385685.2 GPCR-rhodopsin:like:GPR17; 7tm_1/



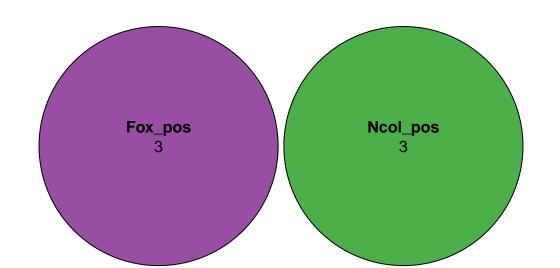
Nvec_vc1.1_XM_032367127.2 bZIP.HG2.10:DBP/HLF/TEF; DBP/HLF/TEF



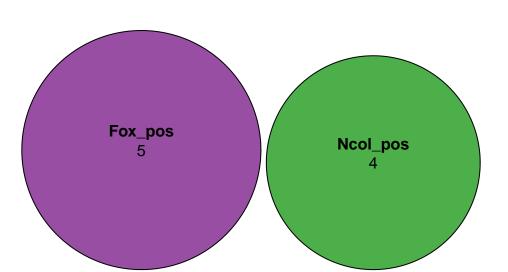
Nvec_vc1.1_XM_048720228.1 Sushi, nidogen and EGF-li; NIDO/



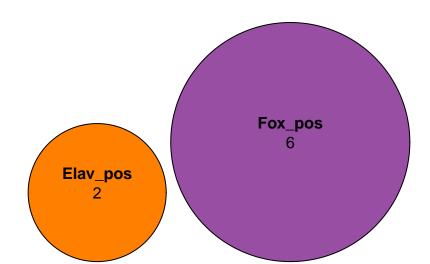
Nvec_vc1.1_XM_048734432.1 Folliculin–interacting pr; FNIP_N/FNIP_M/FNIP_C/



Nvec_vc1.1_XM_001627750.3



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

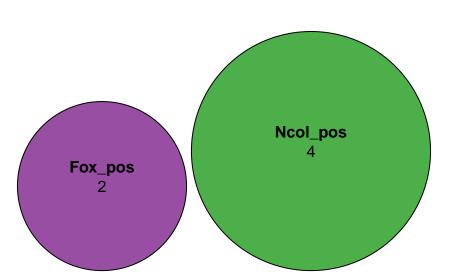


Nvec_vc1.1_XM_001627964.3

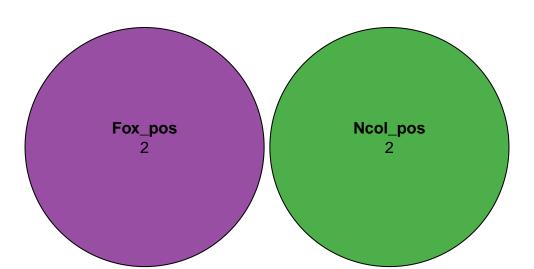
n-NaKCa:KCNA1/KCNA10/KC; ion-NaKCa.HG1.6:KCNA1/KC

Fox_pos Elav pos 13

Nvec_vc1.1_XM_048729250.1

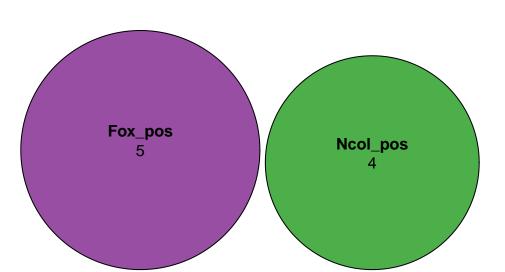


Nvec_vc1.1_XM_001641405.3



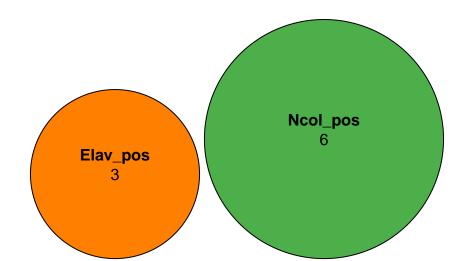
Nvec_vc1.1_XM_032375896.2

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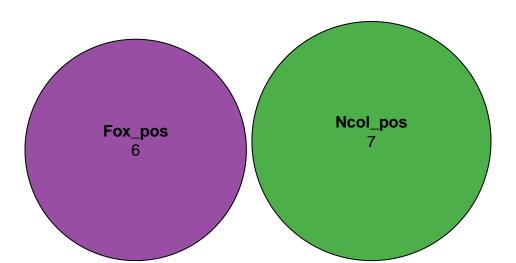


Nvec_vc1.1_XM_032376323.2

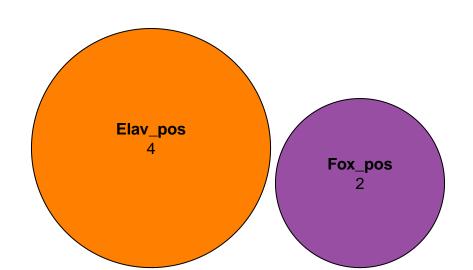
Pkinase:like:GUCY2C/GUCY2; Pkinase_Tyr/HNOBA/Guanyla



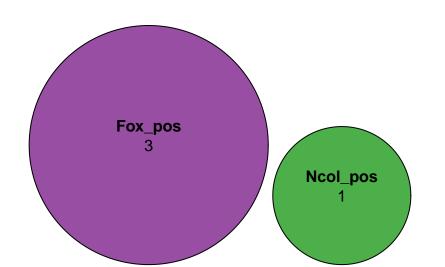
Nvec_vc1.1_XM_032381641.2 b(0,+)-type amino acid tr; AA_permease_2/



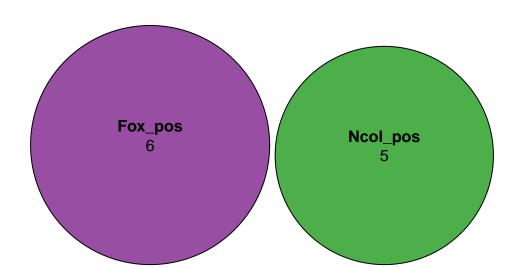
Nvec_vc1.1_XM_001632745.3 Protein NipSnap homolog 3; NIPSNAP/NIPSNAP/



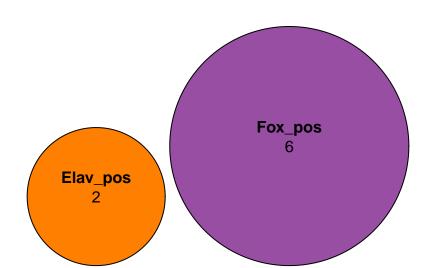
Nvec_vc1.1_XM_048724152.1 GDP-fucose protein O-fuco; O-FucT/



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

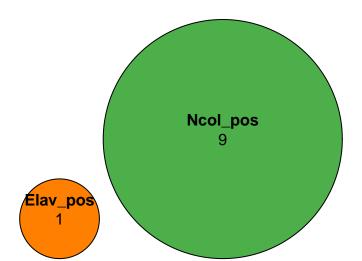


Nvec_vc1.1_XM_048731890.1 GPCR-rhodopsin:like:GPR17; 7tm_1/



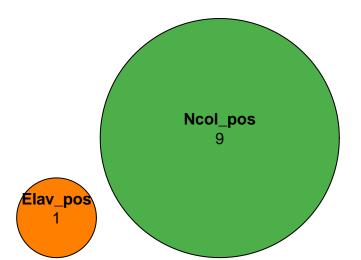
Nvec_NVE13556

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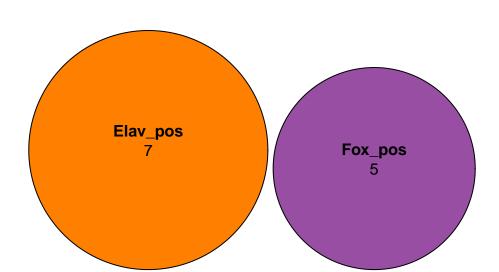


Nvec_vc1.1_XM_032387399.2

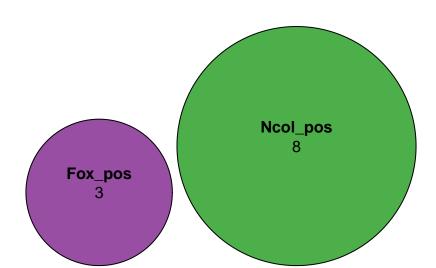
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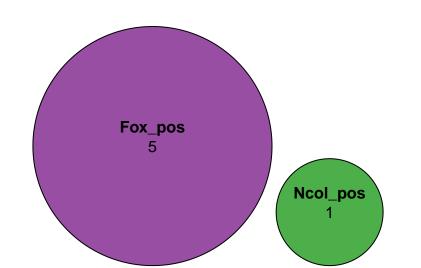
Nvec_vc1.1_XM_048729747.1 Kinesin-like protein KIF1; Kinesin/FHA/



Nvec_vc1.1_XM_032382956.2 Galectin; VHL/

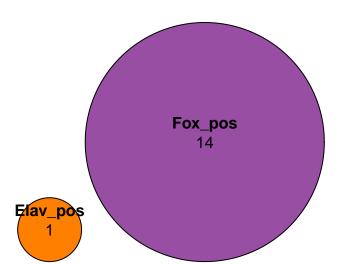


Nvec_vc1.1_XM_001627245.3 cDNA FLJ56852, highly sim; I_LWEQ/



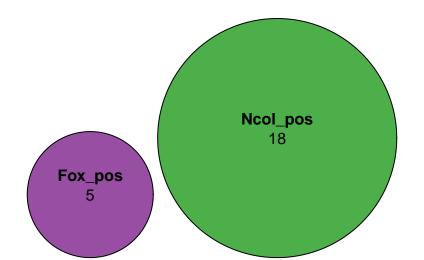
Nvec_vc1.1_XM_001640356.3

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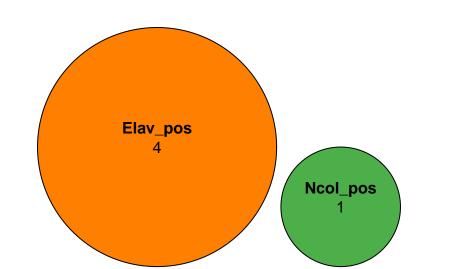


Nvec_vc1.1_XM_032377937.2

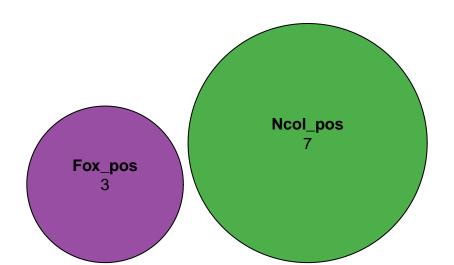
ion-neuro-gated:like:GABR; Neur_chan_LBD/Neur_chan_m



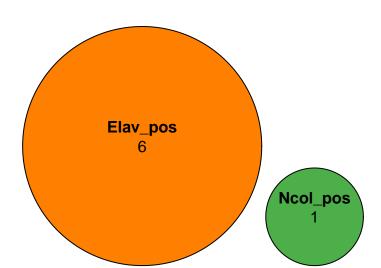
Nvec_vc1.1_XM_001636964.3 cDNA FLJ77769, highly sim; NAD_kinase/



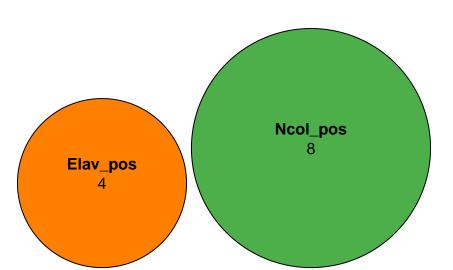
Nvec_vc1.1_XM_001623818.3 ion=NaKCa:like:KCNA1/KCNA; BTB_2/lon_trans/



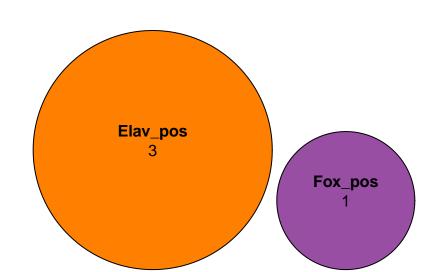
Nvec_vc1.1_XM_001637022.3 GPCR-rhodopsin:NA; 7tm_1/



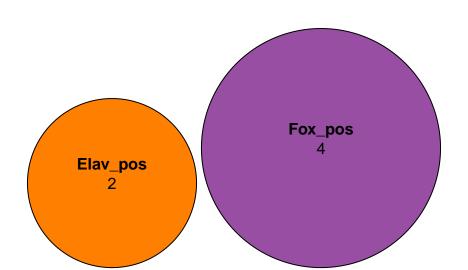
Nvec_vc1.1_XM_048731921.1



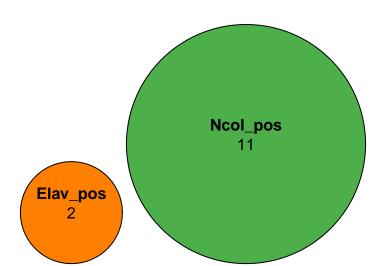
Nvec_vc1.1_XM_001631508.3 YTH:YTHDC1; YTH.HG1.0:YTHDC1



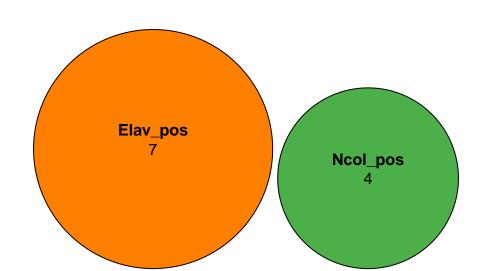
Nvec_vc1.1_XM_032370981.2 Synaptotagmin:SYT12; Synaptotagmin.HG1.38:SYT1



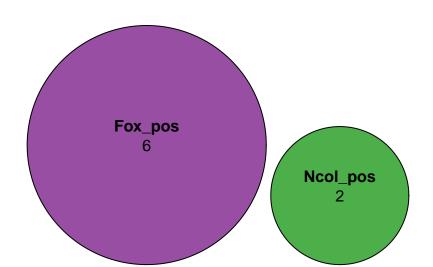
Nvec_vc1.1_XM_032384420.2 Sp6-9; SP6/SP7/SP8/SP9



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

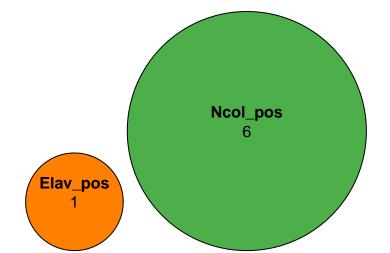


Nvec_vc1.1_XM_001637149.3 RAP1A, member of RAS onco; Ras/

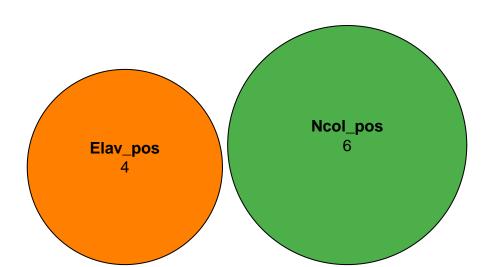


Nvec_vc1.1_XM_001628711.3

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

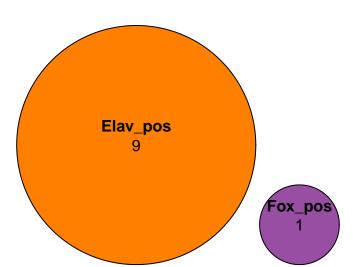


Nvec_vc1.1_XM_001630845.3 ion-glutamate:NA; Ion_trans_2/

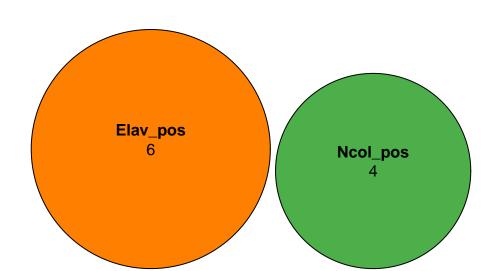


Nvec_vc1.1_XM_032379049.2

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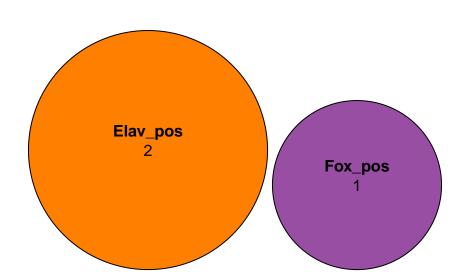


Nvec_vc1.1_XM_032376506.2 Zinc finger protein 862; Dimer_Tnp_hAT/

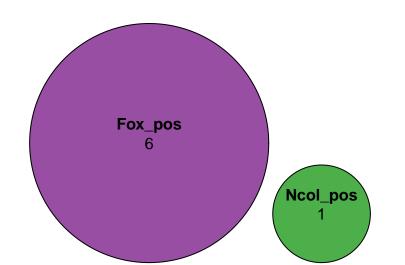


Nvec_vc1.1_XM_032370048.2

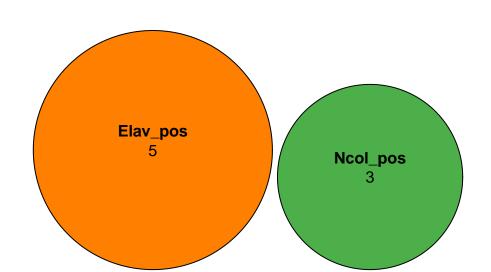
CCCH:MKRN1/MKRN2/MKRN3; zf-CCCH.HG1.17:MKRN1/MK



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

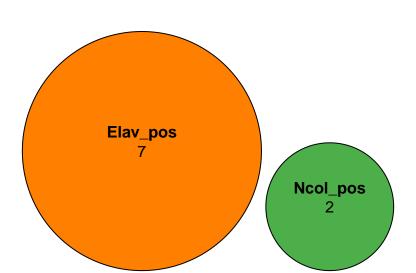


Nvec_vc1.1_XM_048731252.1 OTU domain-containing pro; OTU/

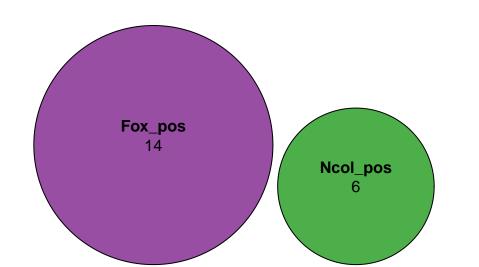


Nvec_vc1.1_XM_001633821.3

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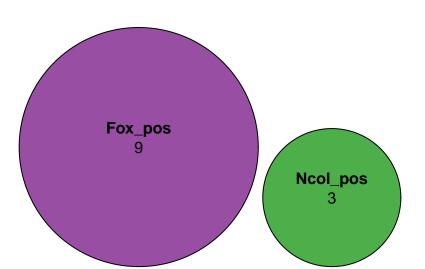


Nvec_vc1.1_XM_032379911.2 Sparc/osteonectin, cwcv a; Thyroglobulin_1/

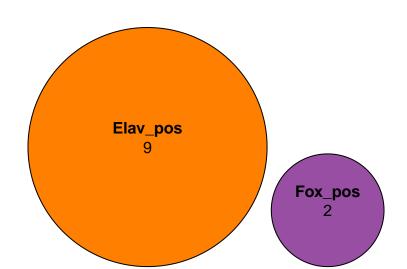


Nvec_vc1.1_XM_032370084.2

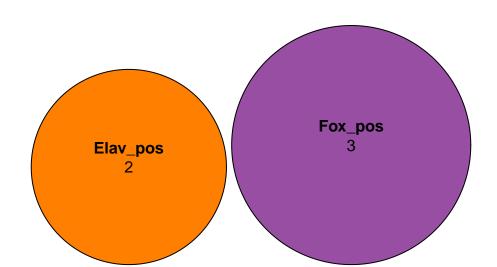
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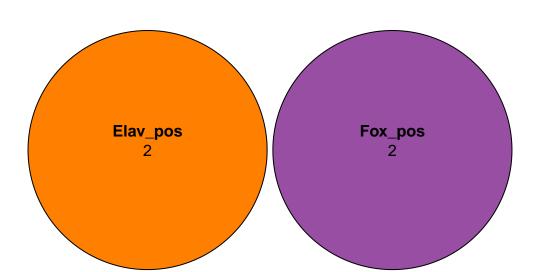
Nvec_vc1.1_XM_048723783.1 Cilia- and flagella-assoc;

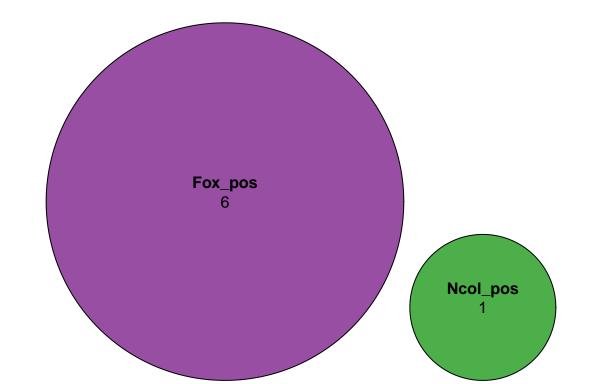


Nvec_vc1.1_XM_048725946.1 WD repeat–containing prot;

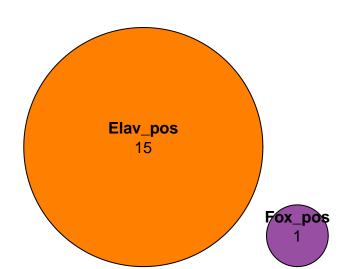


Nvec_vc1.1_XM_001636610.3

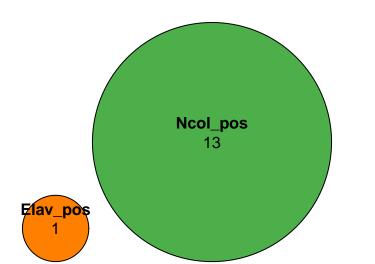




Nvec_vc1.1_XM_001628279.3 HLH.HG1.76:ATOH8; ATOH8

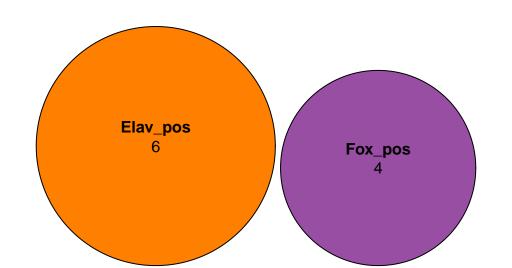


Nvec_vc1.1_XM_032377736.2 Protein FAM173B; Methyltransf_23/

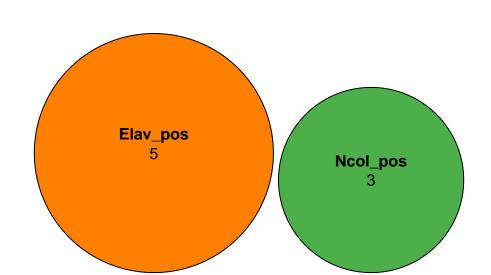


Nvec_vc1.1_XM_032361849.2

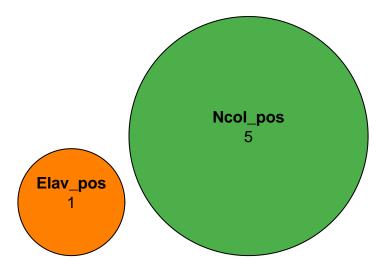
Ets.HG1.2:ELK1/ELK3/ELK4; ELK1/ELK3/ELK4



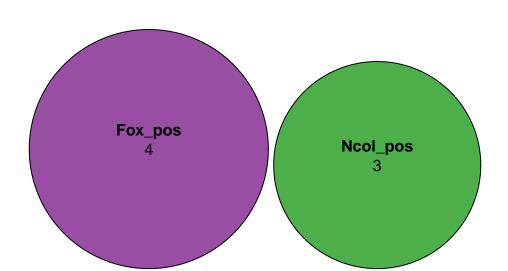
Nvec_vc1.1_XM_048731003.1 ion=NaKCa:NA; Ion_trans_2/



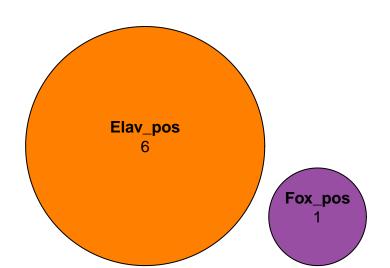
Nvec_vc1.1_XM_001639611.1 ; EF-hand_7/



Nvec_vc1.1_XM_032362682.2 CEA protein; V-set/lg_2/

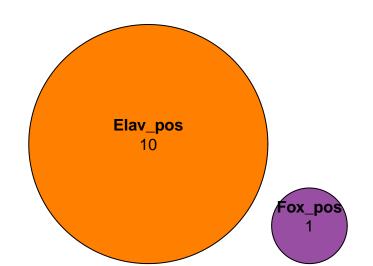


Nvec_vc1.1_XM_032380524.2 Pkinase:PIK3R4; Pkinase.HG180.0:PIK3R4

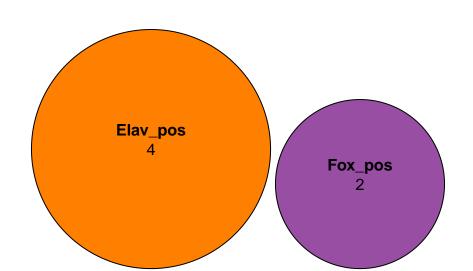


Nvec_vc1.1_XM_032378372.2

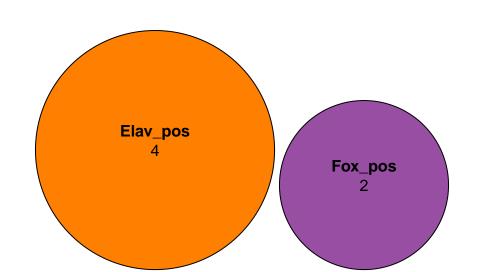
GPCR-rhodopsin:like:ADORA; 7tm_1/



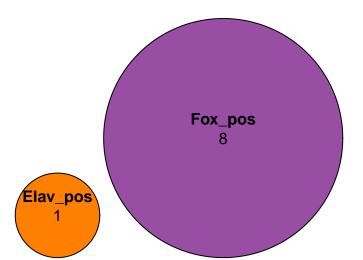
Nvec_vc1.1_XM_032386263.2 Rhombotin-1; LIM/LIM/



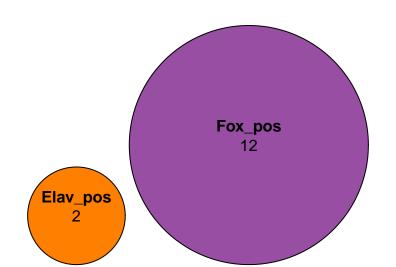
Nvec_vc1.1_XM_032378457.2 GPCR-rhodopsin:NA; 7TM_GPCR_Srw/



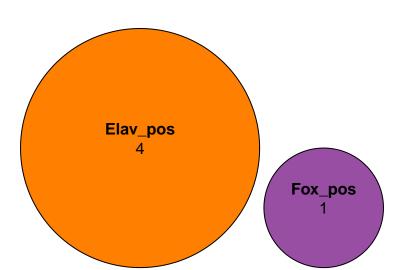
Nvec_vc1.1_XM_032379543.2 ; DUF885/



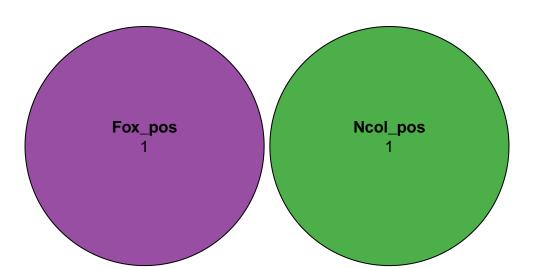
Nvec_vc1.1_XM_001633143.3 Replication factor C subu; AAA/Rep_fac_C/



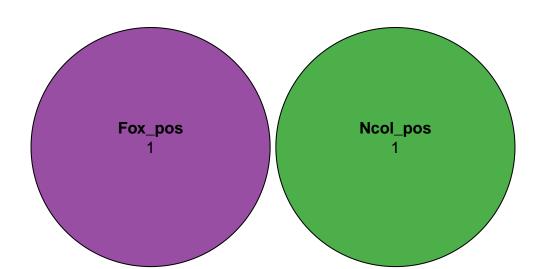
Nvec_vc1.1_XM_048720909.1



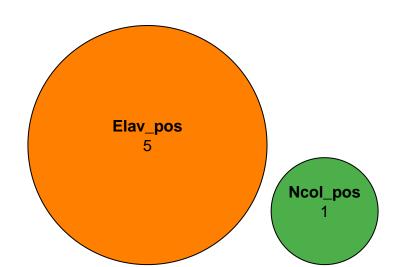
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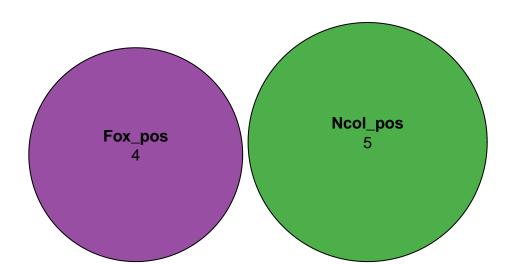
Nvec_vc1.1_XM_001623762.3 Vesicular glutamate trans; MFS_1/



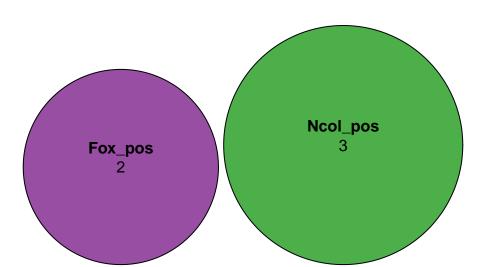
Nvec_vc1.1_XM_001627181.3 Myeloid leukemia factor 1; Mlf1IP/



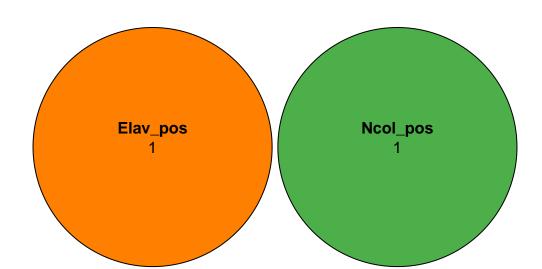
Nvec_vc1.1_XM_032372769.2 fibronectin:like:AXL/MERT; I-set/lg_2/fn3/fn3/



Nvec_vc1.1_XM_048734459.1 E3 ubiquitin-protein liga;

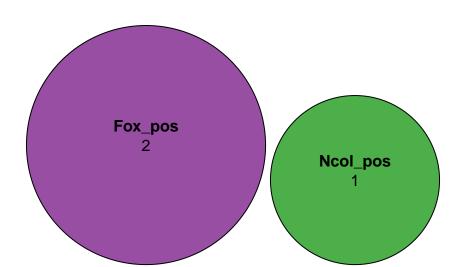


Nvec_vc1.1_XM_001636661.3 Serine/threonine-protein; Metallophos/

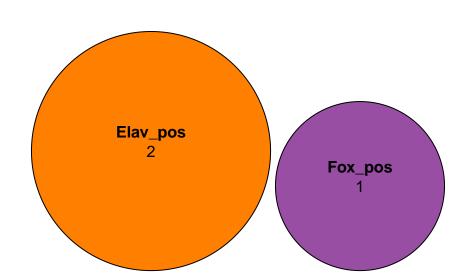


Nvec_vc1.1_XM_001639688.3

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

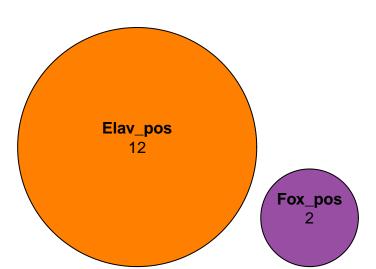


Nvec_vc1.1_XM_032374307.2 Major facilitator superfa; MFS_2/

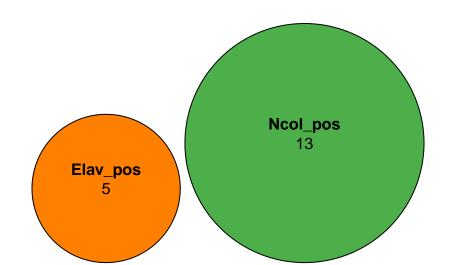


Nvec_vc1.1_XM_032365655.2

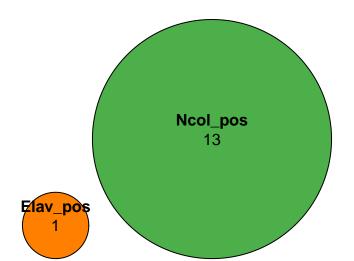
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Nvec_vc1.1_XM_048720751.1 cDNA FLJ33589 fis, clone ; Asparaginase_2/

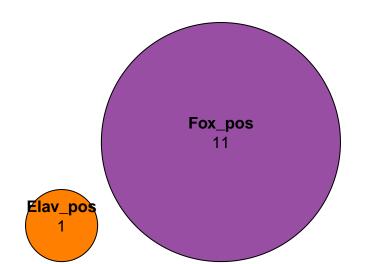


Nvec_vc1.1_XM_048720757.1 cDNA FLJ33589 fis, clone ; Asparaginase_2/Asparagina



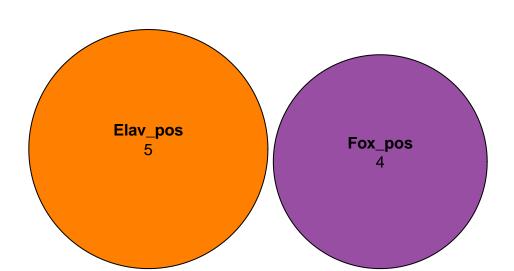
Nvec_vc1.1_XM_032362262.2

GPCR-rhodopsin:like:CCKAR; 7tm_1/

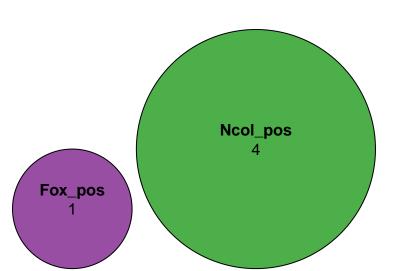


Nvec_vc1.1_XM_048734599.1

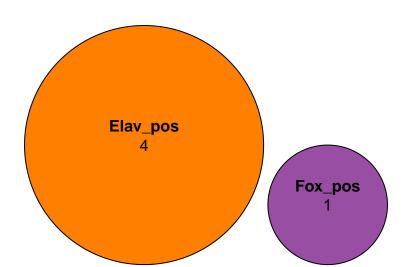
GPCR-rhodopsin:like:ADORA; 7tm_1/



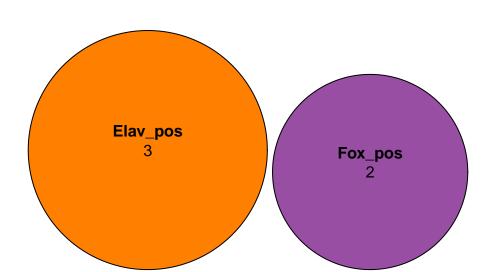
Nvec_vc1.1_XM_048723891.1



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

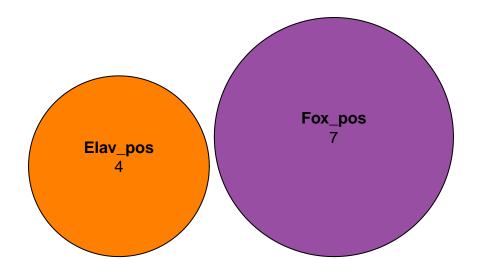


Nvec_vc1.1_XM_032363157.2

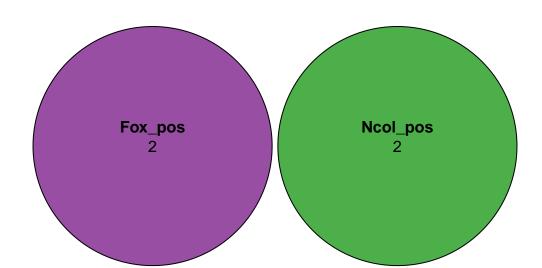


Nvec_vc1.1_XM_001633244.3

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

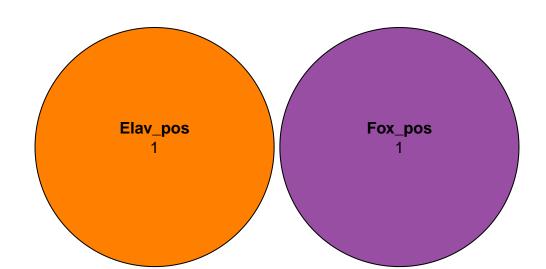


Nvec_vc1.1_XM_001639954.3 GPCR-rhodopsin:like:GPR27; 7tm_1/

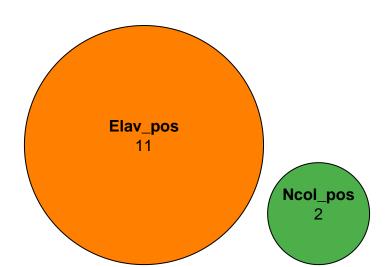


Nvec_vc1.1_XM_032372557.2

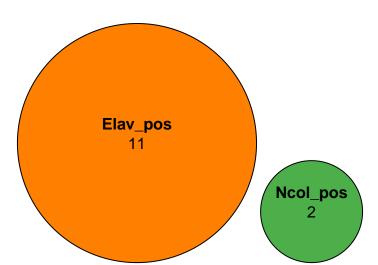
Synaptotagmin:like:FO6814; C2/C2/



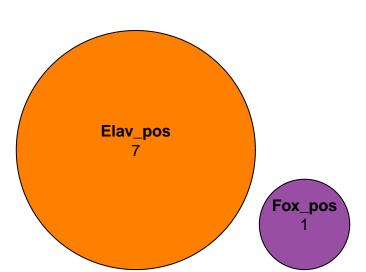
Nvec_vc1.1_XM_001636882.3 ADP-ribosylation factor-I; Arf/



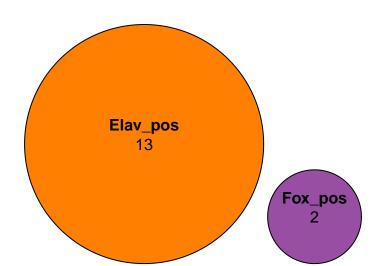
Nvec_vc1.1_XM_048719664.1



Nvec_vc1.1_XM_048725869.1

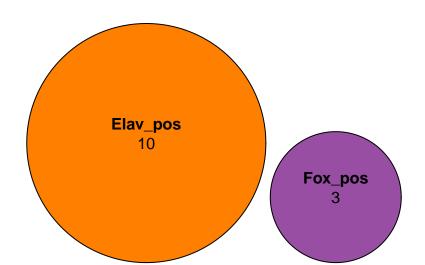


Nvec_vc1.1_XM_001632492.3 Alpha-(1,3)-fucosyltransf; Glyco_transf_10/

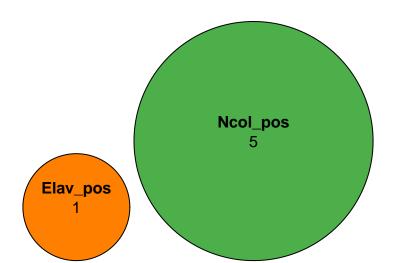


Nvec_vc1.1_XM_001639681.3

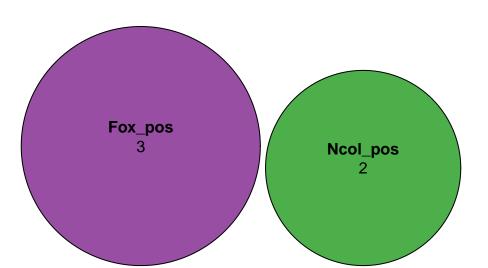
Potassium channel, subfam; lon_trans_2/lon_trans_2/



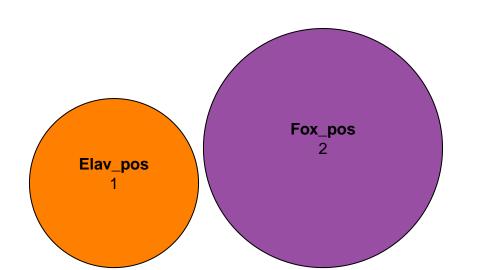
Nvec_vc1.1_XM_048729870.1 60 kDa lysophospholipase; Asparaginase/Ank_2/



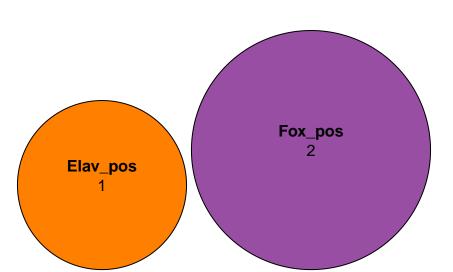
Nvec_NVE13928



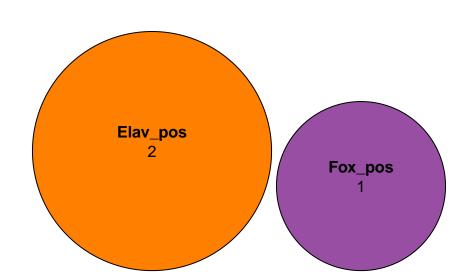
Nvec_vc1.1_XM_032364349.2 Aspartate aminotransferas; Aminotran_1_2/



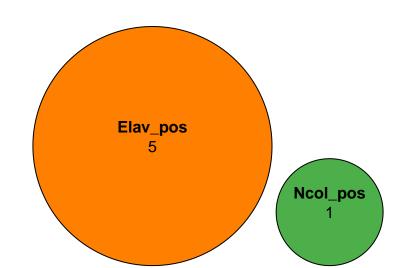
Nvec_vc1.1_XM_032376979.2



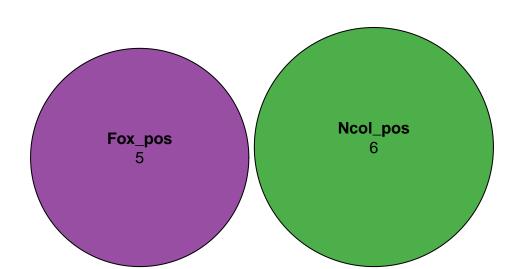
Nvec_vc1.1_XM_048730108.1 GPCR-rhodopsin:NA; 7tm_1/



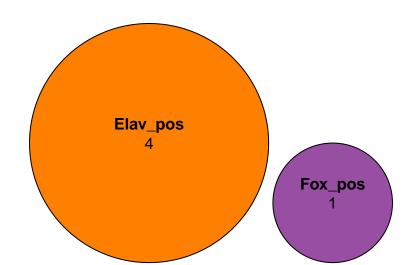
Nvec_vc1.1_XM_032382511.2 Serine incorporator 1; Serinc/Serinc/



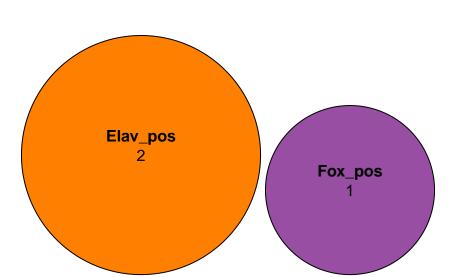
Nvec_vc1.1_XM_032382117.2 GPCR-rhodopsin:like:GPR17; 7tm_1/



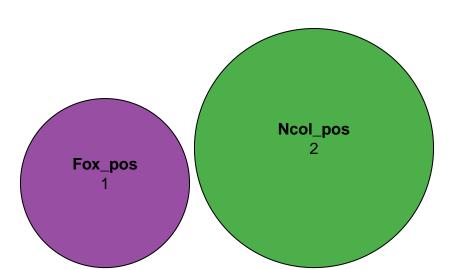
Nvec_vc1.1_XM_032386298.2 GPCR-rhodopsin:NA; 7tm_1/



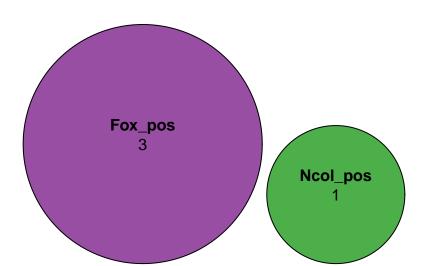
Nvec_vc1.1_XM_032362299.2



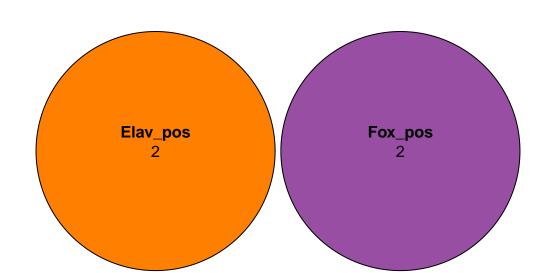
Nvec_vc1.1_XM_001633615.3 HCG2044780; Kua-UEV1_localn/



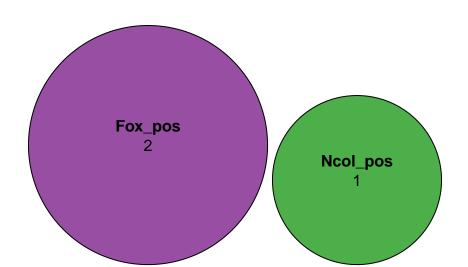
Nvec_vc1.1_XM_001636545.3 ; RNase_P_Rpp14/



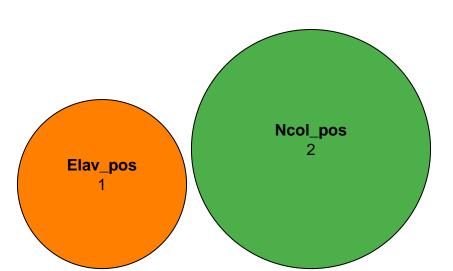
Nvec_vc1.1_XM_048720443.1 C-Maf-inducing protein;



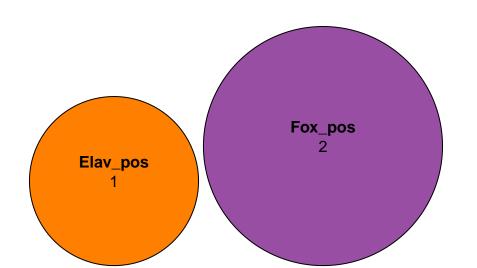
Nvec_vc1.1_XM_032384695.2 Monocarboxylate transport; MFS_1/



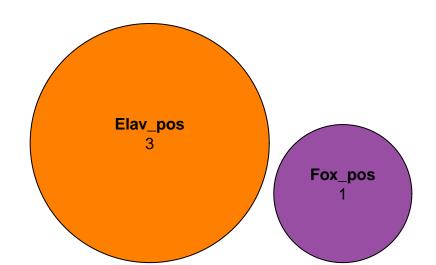
Nvec_vc1.1_XM_048731044.1



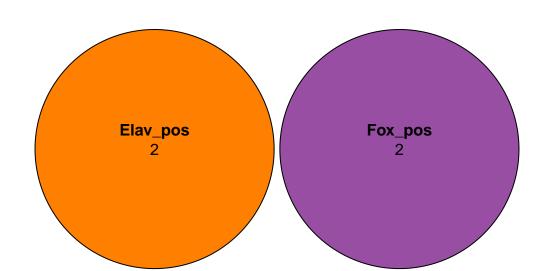
Nvec_vc1.1_XM_001629129.3 Probable aminopeptidase N; Peptidase_M17/



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

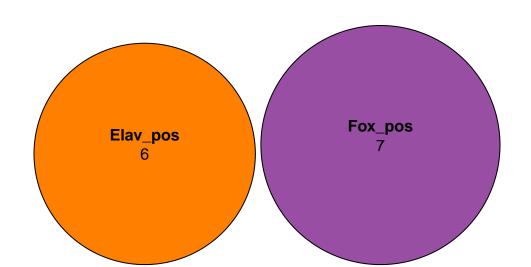


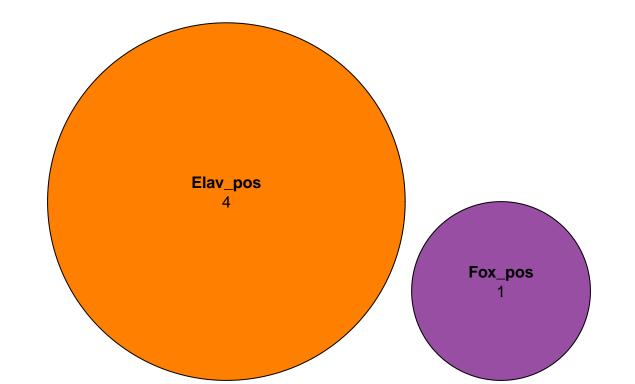
Nvec_vc1.1_XM_032373148.2 S1:SUPT6H; S1.HG5.0:SUPT6H



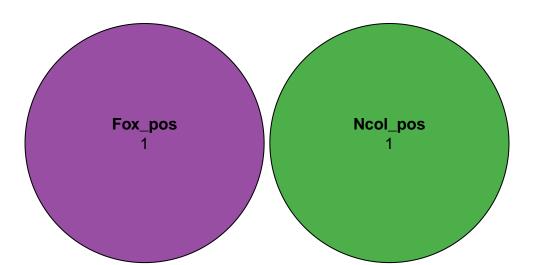
Nvec_vc1.1_XM_032381433.2

Guanylate cyclase soluble; HNOB/HNOBA/Guanylate_cyc/

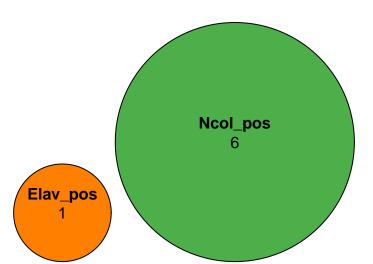




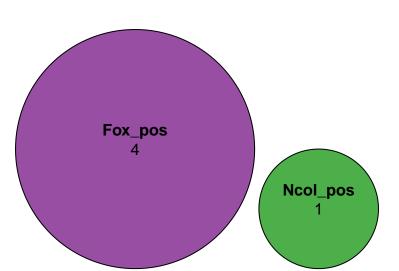
Nvec_NVE7893



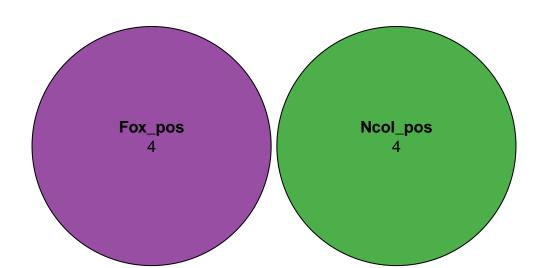
Nvec_vc1.1_XM_032372214.2



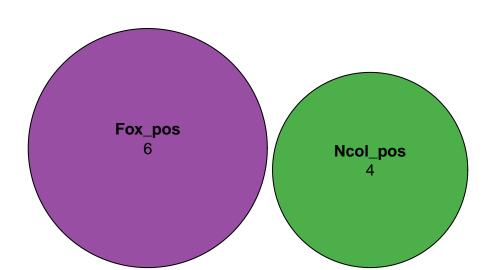
Nvec_vc1.1_XM_048730962.1



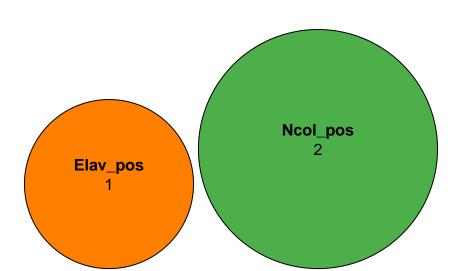
Nvec_vc1.1_XM_001637616.3 Fukutin related protein; LicD/



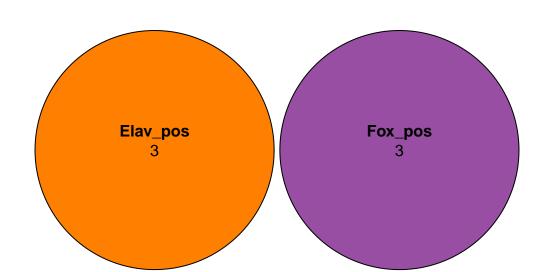
Nvec_vc1.1_XM_001628059.3 Fatty acid amide hydrolas; Amidase/



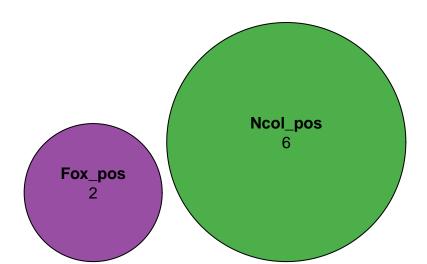
Nvec_vc1.1_XM_048730184.1 Thyroid receptor–interact;



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

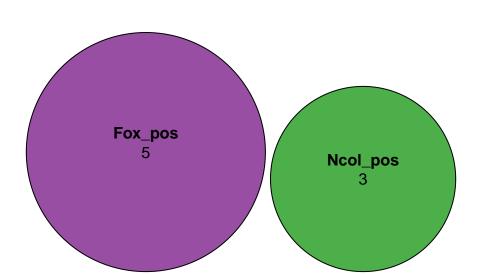


Nvec_vc1.1_XM_001641783.3 mTERF.HG1.0:MTERF4; MTERF4

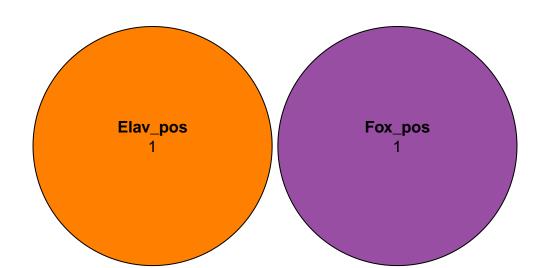


Nvec_vc1.1_XM_032374380.2

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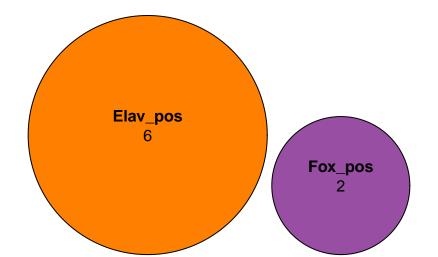


Nvec_vc1.1_XM_032384990.2 Protein odr-4 homolog; ODR4-like/ODR4-like/



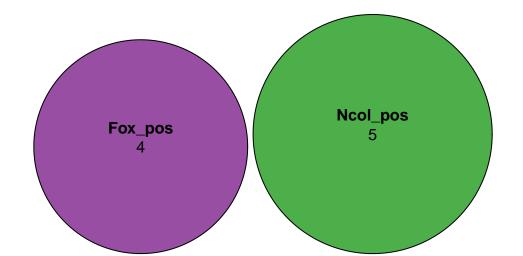
Nvec_vc1.1_XM_048729479.1

TAF5-like RNA polymerase ; TFIID_90kDa/WD40/WD4

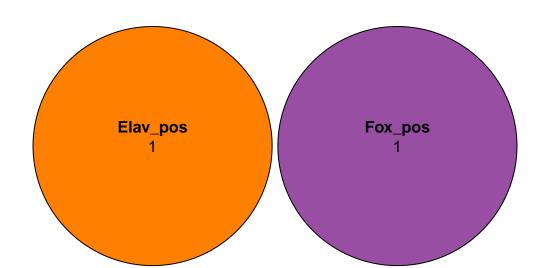


Nvec_vc1.1_XM_001639817.3

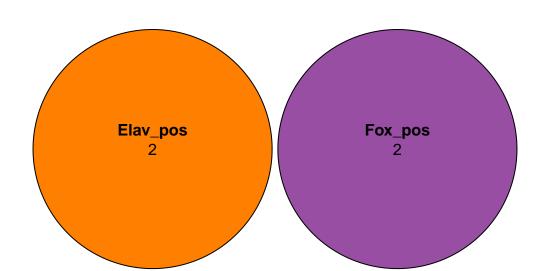
Ephrin:AL691442/EFNA1/EFN; Ephrin.HG1.0:AL691442/EFN



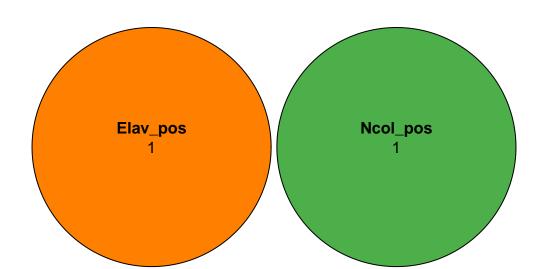
Nvec_vc1.1_XM_032366492.2 N-acetylated-alpha-linked; TFR_dimer/



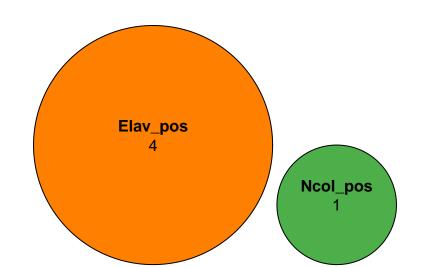
Nvec_vc1.1_XM_048729868.1 E3 ubiquitin-protein liga;



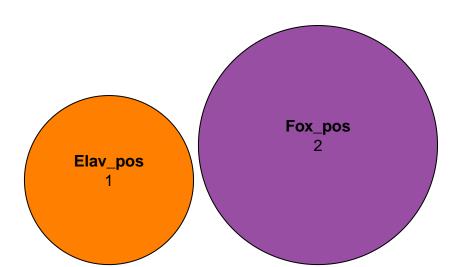
Nvec_vc1.1_XM_048734712.1 Very long-chain-specific;



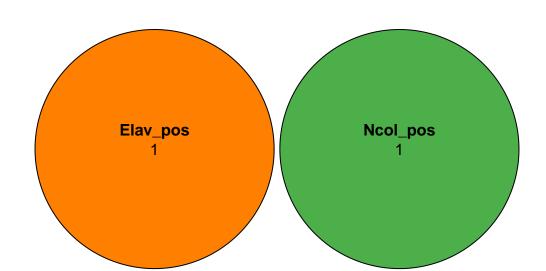
Nvec_vc1.1_XM_001628363.3 cDNA, FLJ94698, highly si; Sybindin/



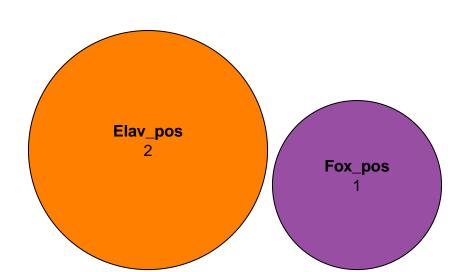
Nvec_vc1.1_XM_032387176.2 LMBR1 domain-containing p; LMBR1/



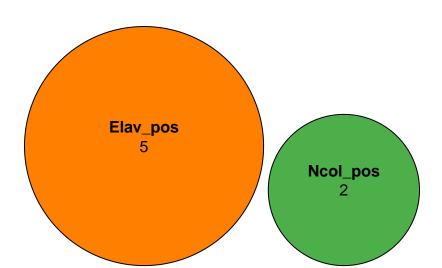
Nvec_vc1.1_XM_001631292.3 ADP-ribosylation factor r;



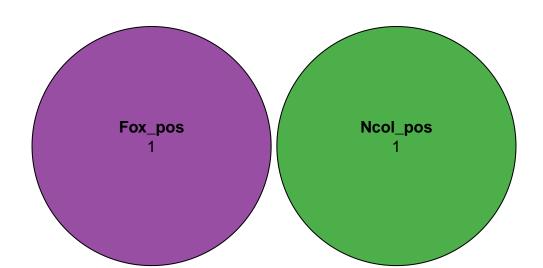
Nvec_vc1.1_XM_001631286.2 NNMT protein; NNMT_PNMT_TEMT/



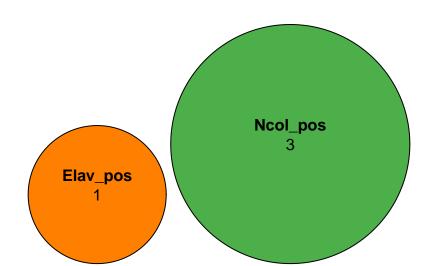
Nvec_vc1.1_XM_001624997.3 ; F-box-like/



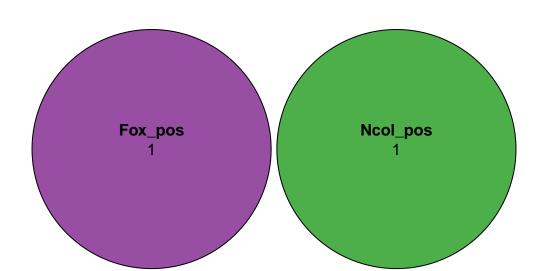
Nvec_vc1.1_XM_001641286.3 cDNA FLJ51917, highly sim; TIMP/



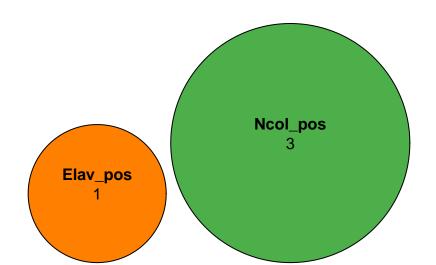
Nvec_vc1.1_XM_032368559.2 Protein prenyltransferase; PPTA/



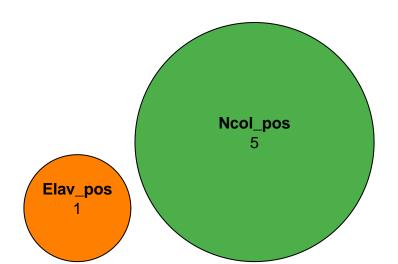
Nvec_vc1.1_XM_048727242.1 Zinc transporter 7; Cation_efflux/



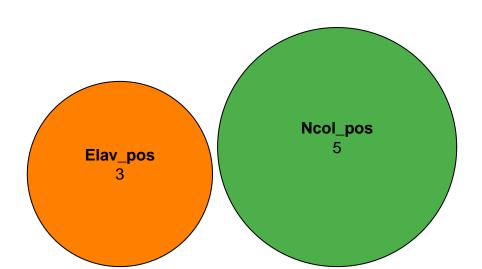
Nvec_vc1.1_XM_032372191.2 Abhydrolase domain contai; Abhydrolase_6/



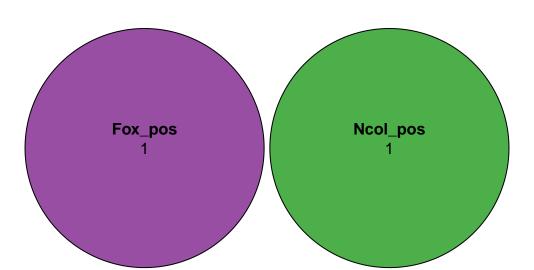
Nvec_vc1.1_XM_048730264.1 Component of oligomeric g; COG2/DUF3510/



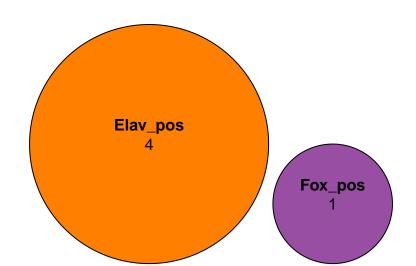
Nvec_vc1.1_XM_032375751.2 Transmembrane protein 115; DUF1751/



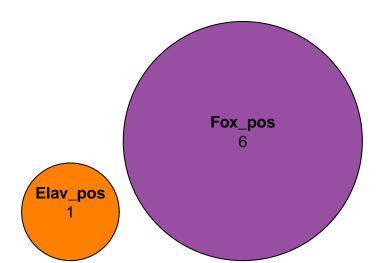
Nvec_vc1.1_XM_048725016.1



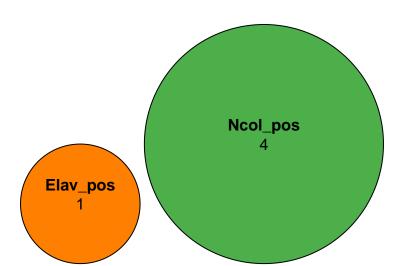
Nvec_vc1.1_XM_001638793.3 Zinc finger CCHC domain-c; PSP/



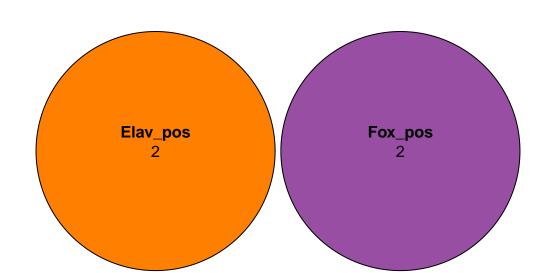
Nvec_vc1.1_XM_032363581.2 GPCR-rhodopsin:NA; 7tm_1/

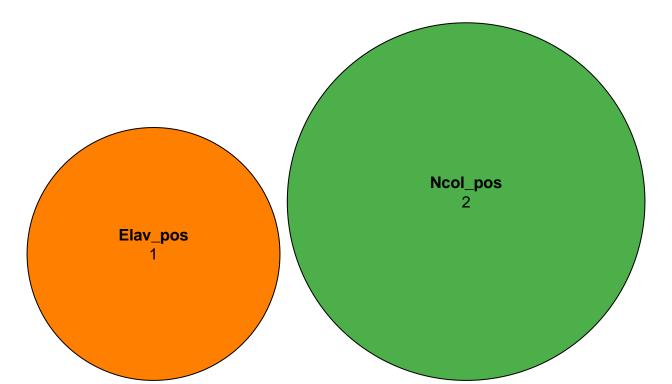


Nvec_vc1.1_XM_032364770.2 GPCR-rhodopsin:NA; 7tm_1/



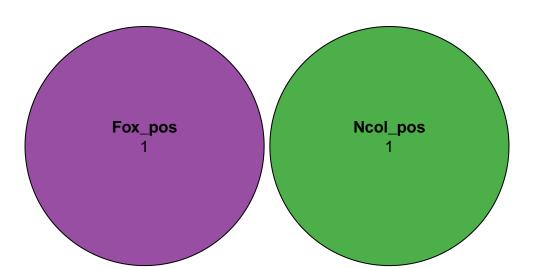
Nvec_vc1.1_XM_032367592.2 cDNA, FLJ79298, highly si;



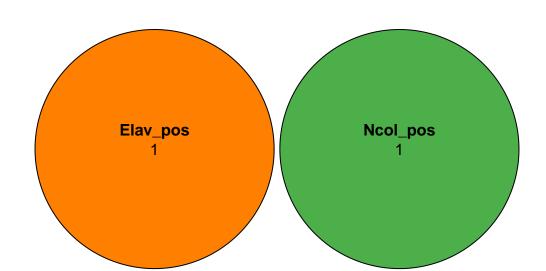


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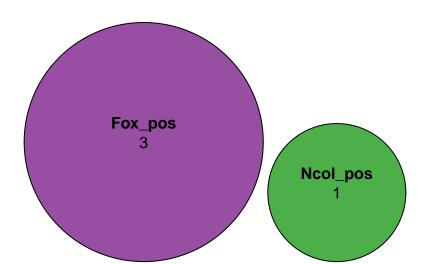
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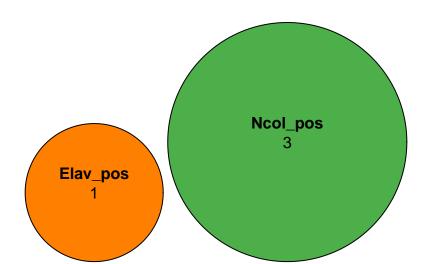
Nvec_vc1.1_XM_032385089.2 Ets.HG1.7:GABPA; GABPA



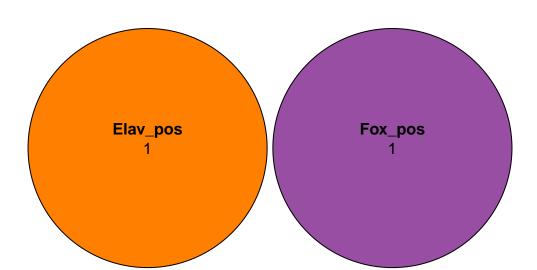
Nvec_vc1.1_XM_048720705.1 Ubiquitin carboxyl-termin;



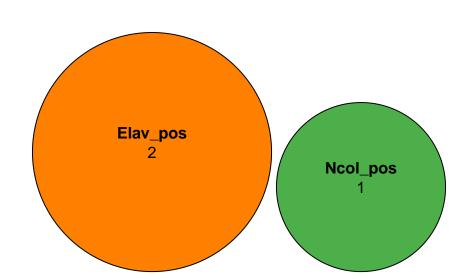
Nvec_vc1.1_XM_032362554.2 Synembryn-A; Ric8/



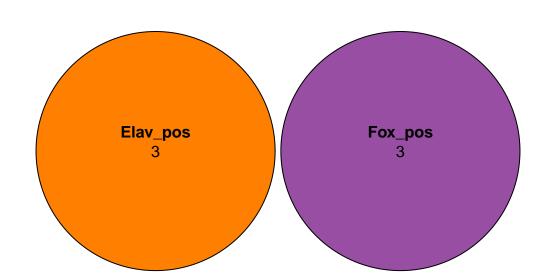
Nvec_vc1.1_XM_048726963.1



Nvec_vc1.1_XM_032382564.2 Uncharacterized protein; TMEM219/



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



Nvec_vc1.1_XM_048730070.1 FAM179B protein;

