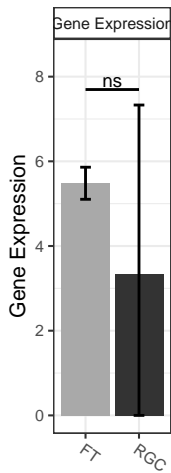
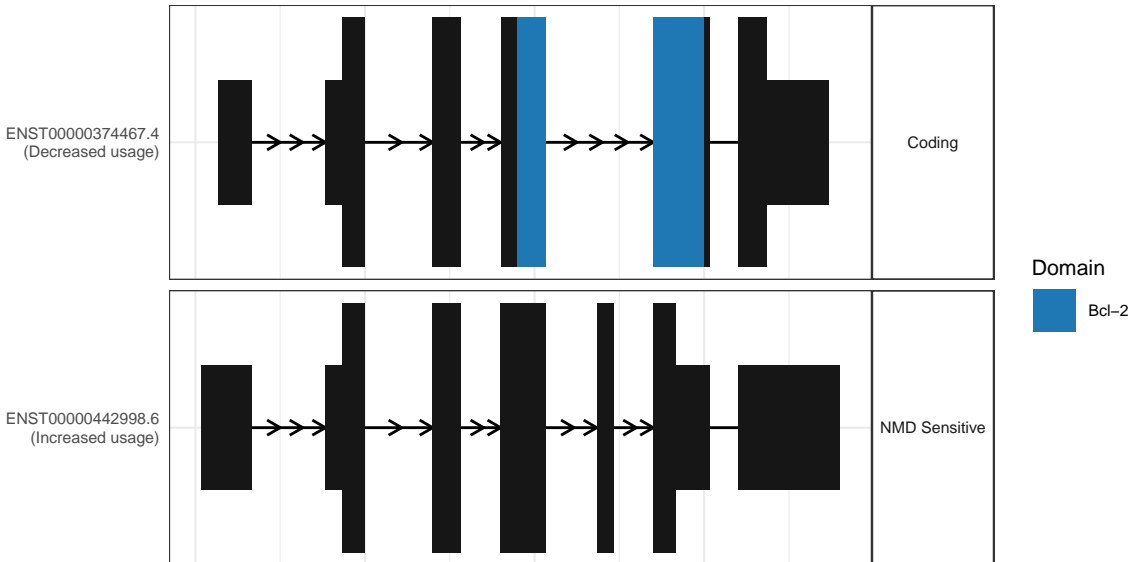
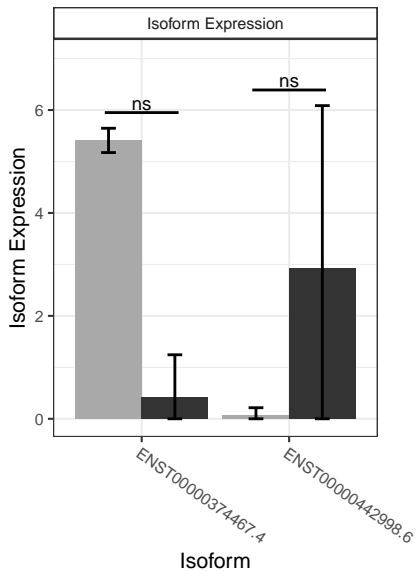


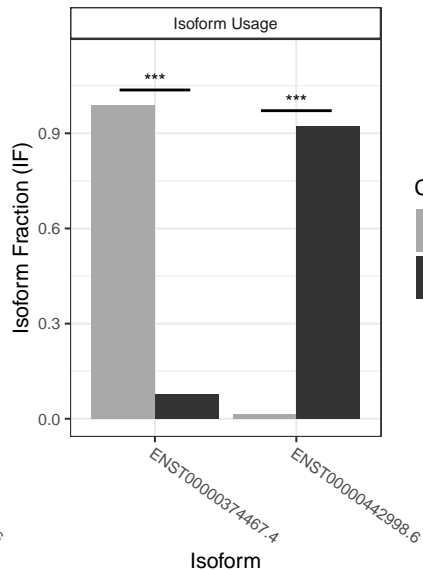
The isoform switch in BAK1 (FT vs RGC)



Condition



Isoform



Isoform

Condition

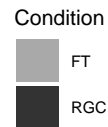
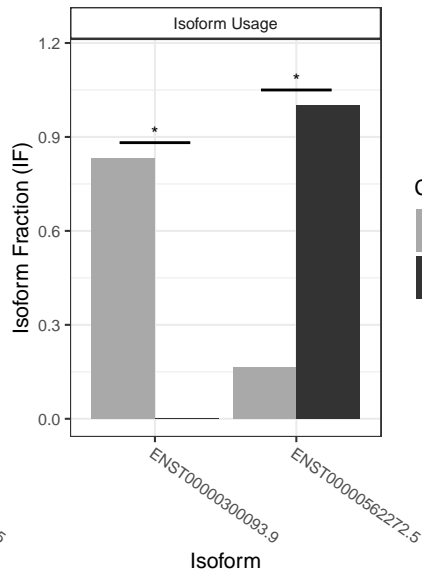
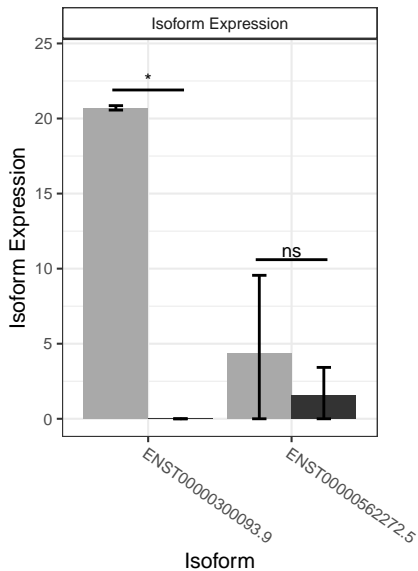
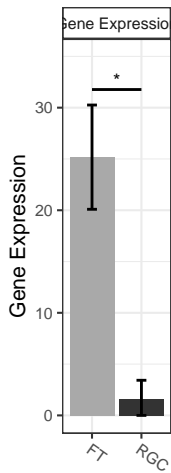
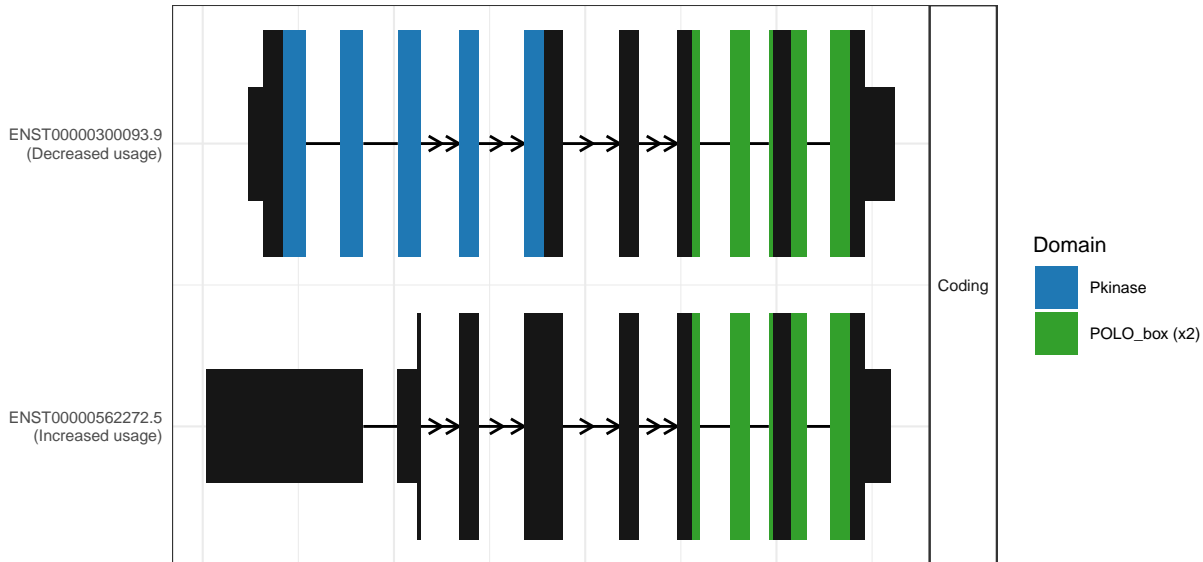


Domain



Bcl-2

The isoform switch in PLK1 (FT vs RGC)



Domain

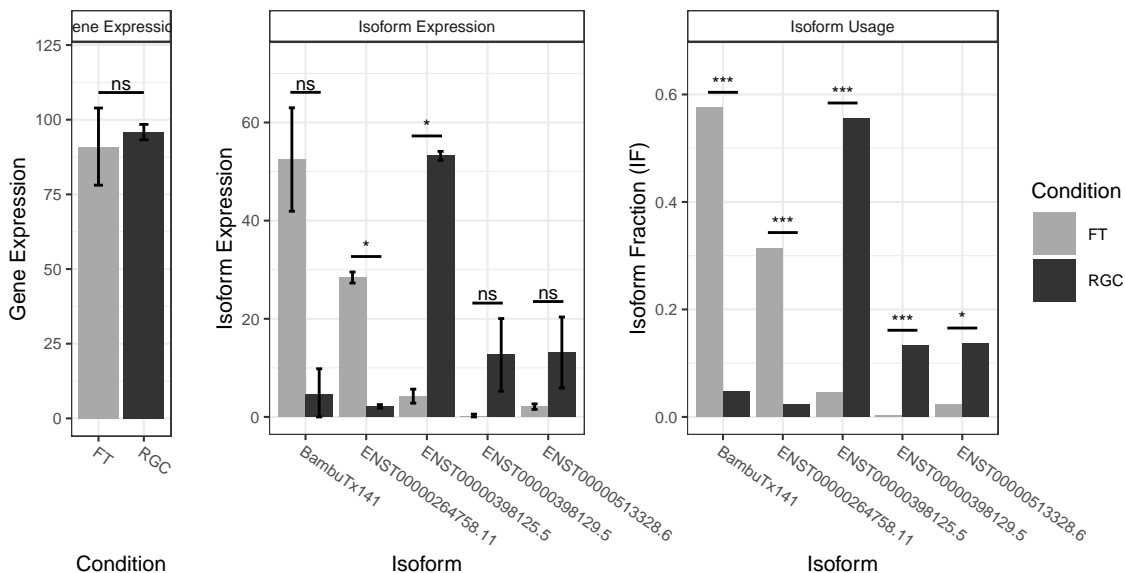
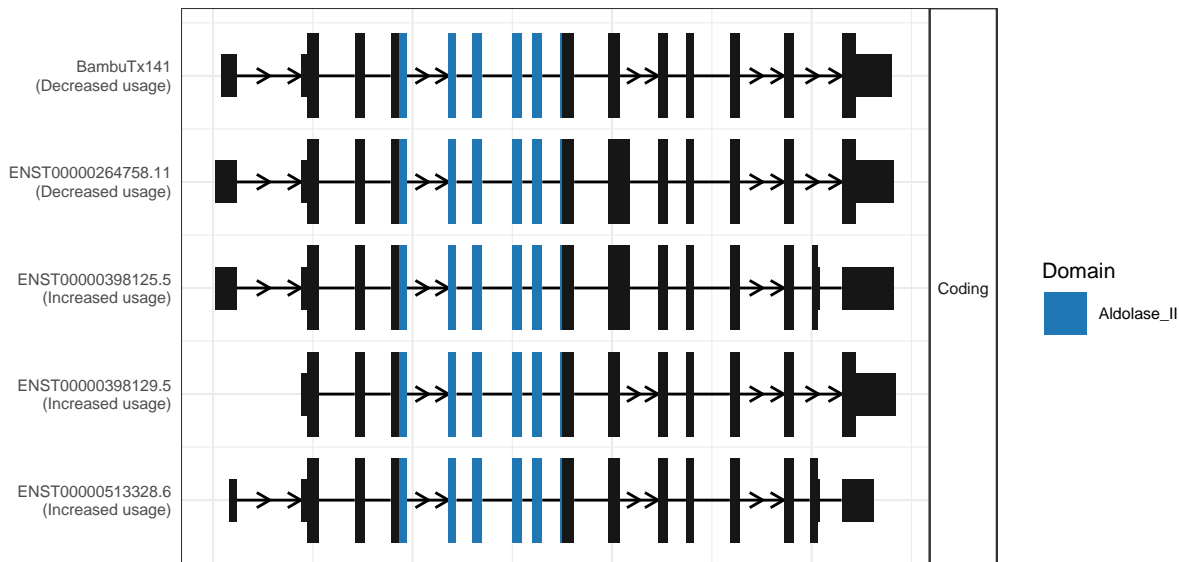


Kinase



POLO_box (x2)

The isoform switch in ADD1 (FT vs RGC)

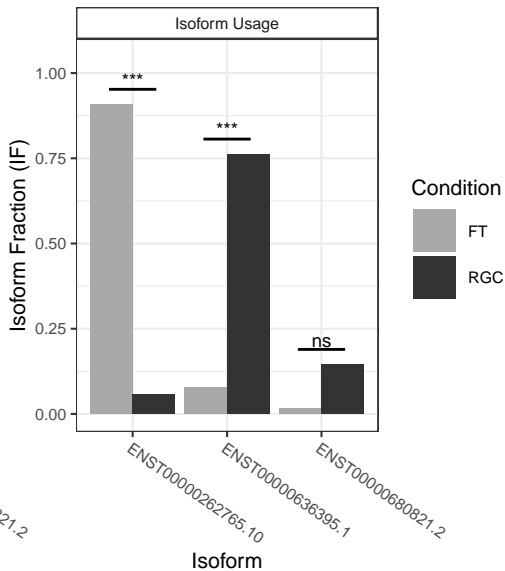
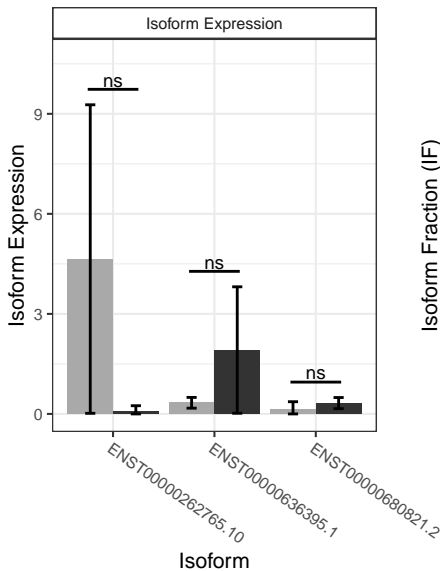
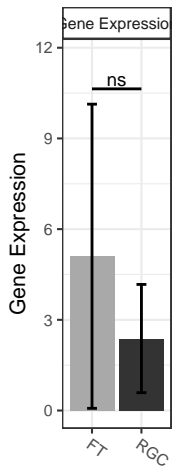
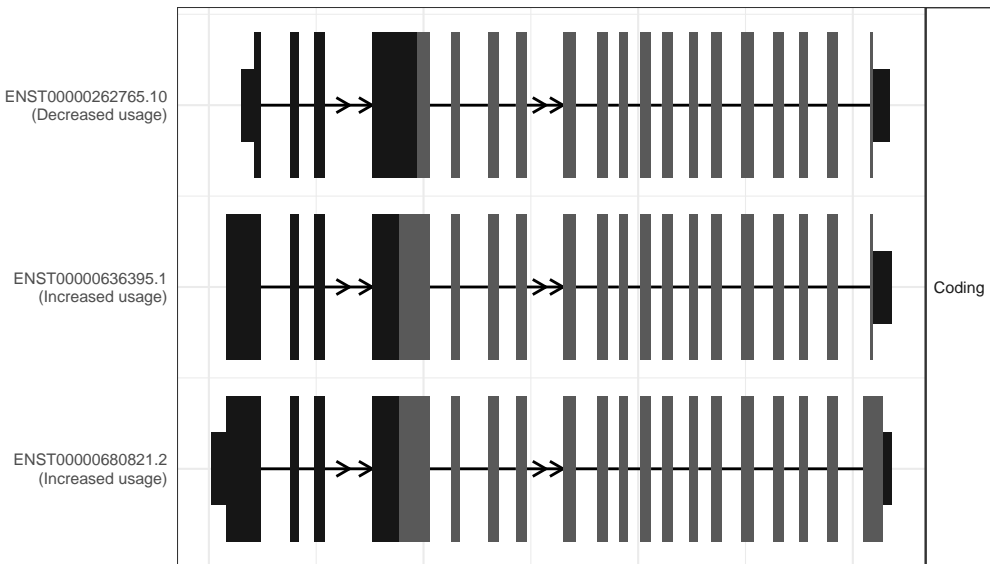


Domain



Aldolase_II

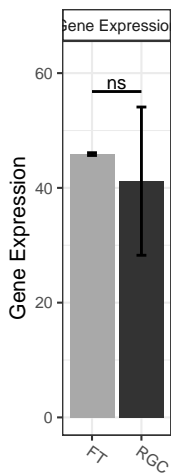
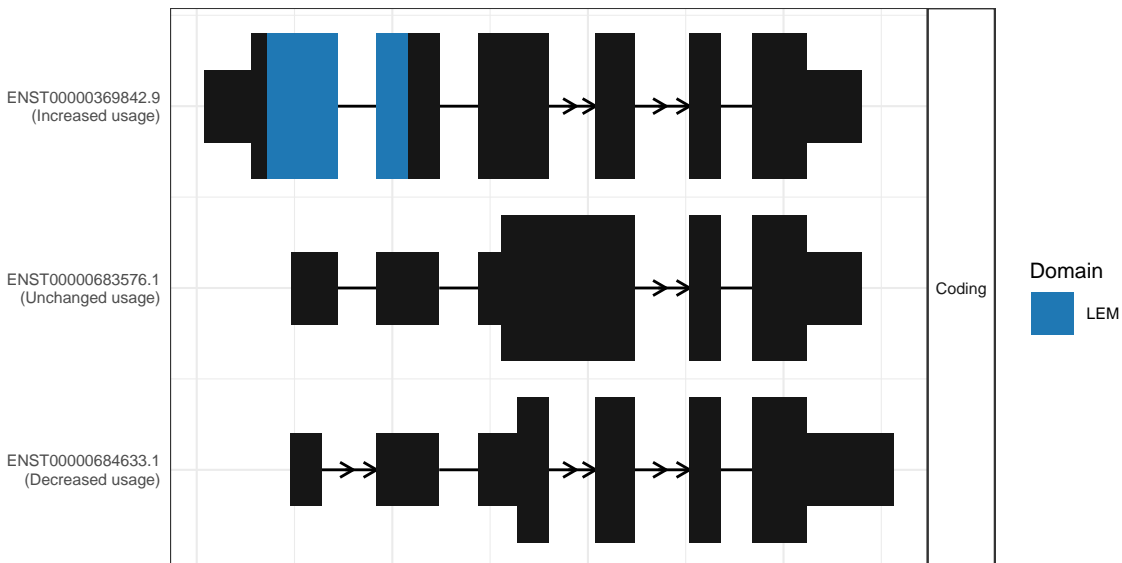
The isoform switch in QRICH2 (FT vs RGC)



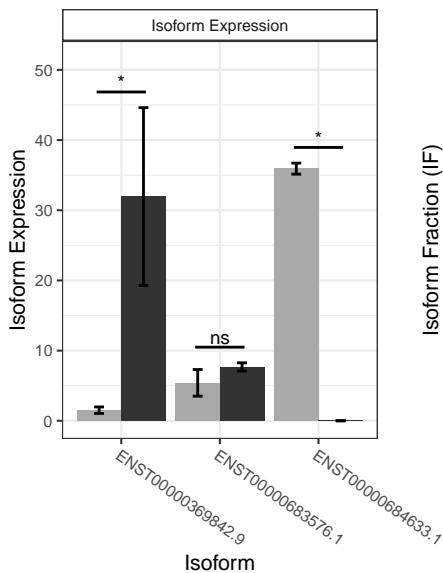
Condition



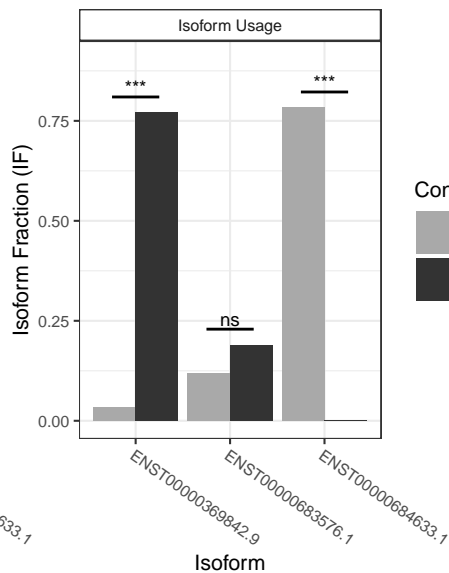
The isoform switch in EMD (FT vs RGC)



Condition



Isoform



Isoform

Condition

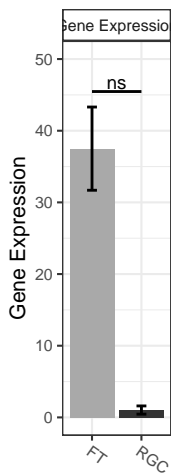
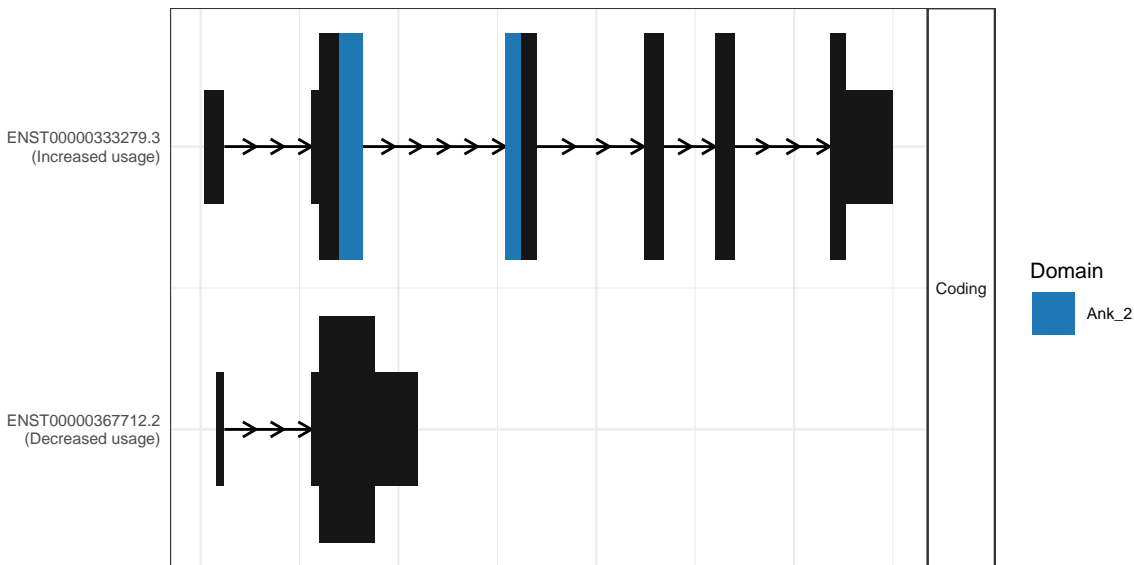


Domain

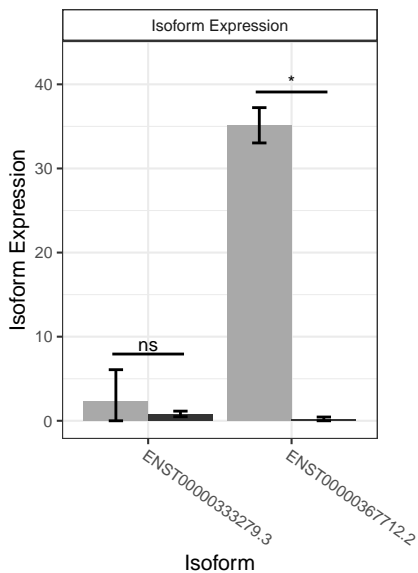


LEM

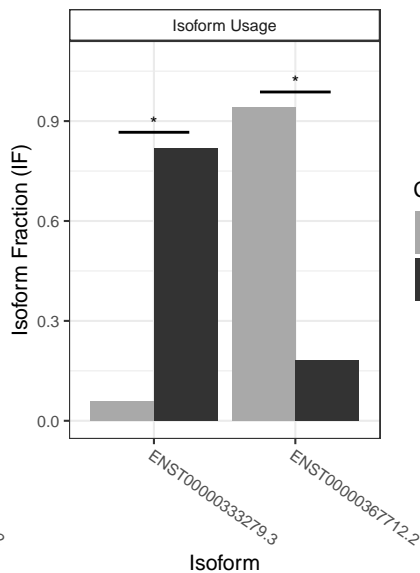
The isoform switch in ANKRD45 (FT vs RGC)



Condition



Isoform



Isoform

Condition

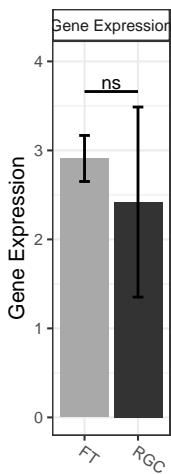
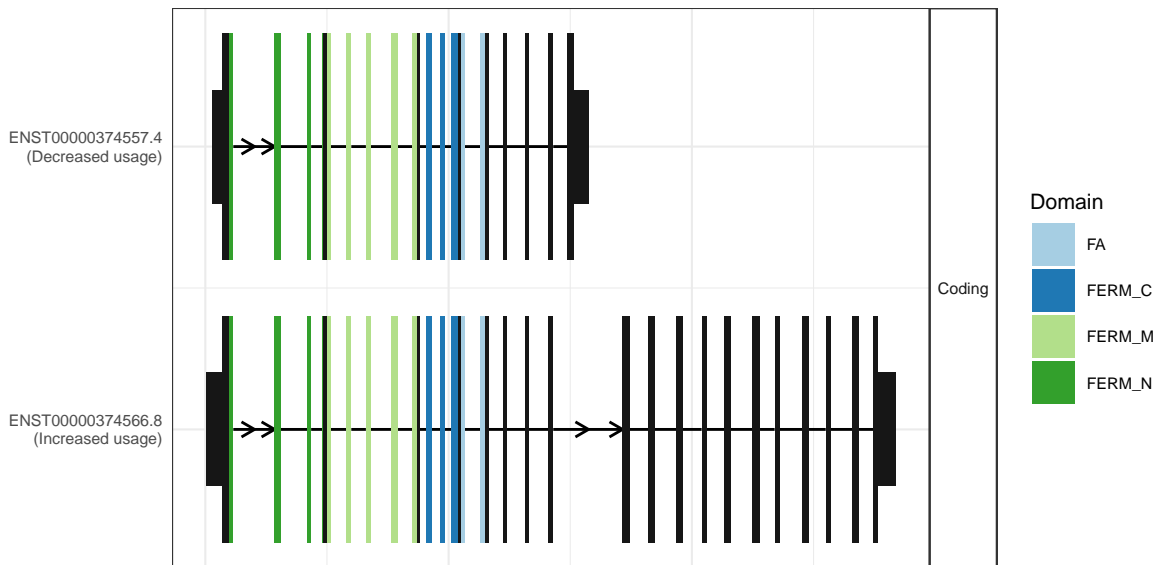


Domain

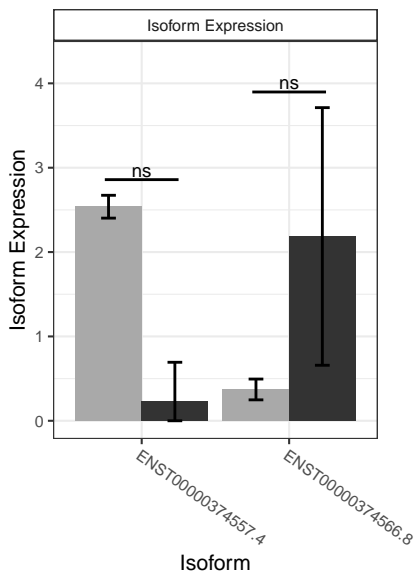


Ank_2

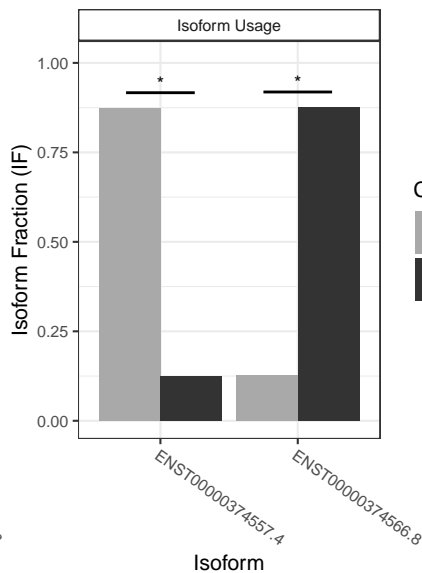
The isoform switch in EPB41L4B (FT vs RGC)



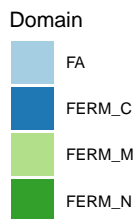
Condition



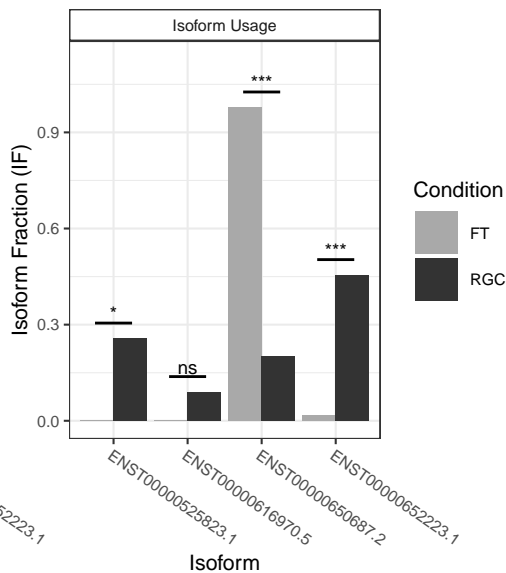
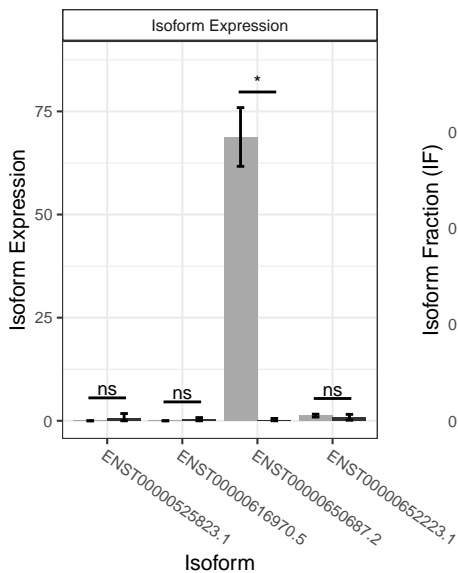
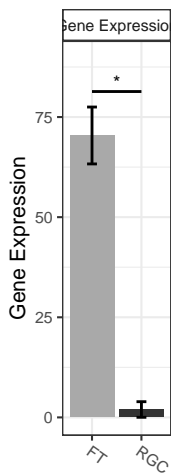
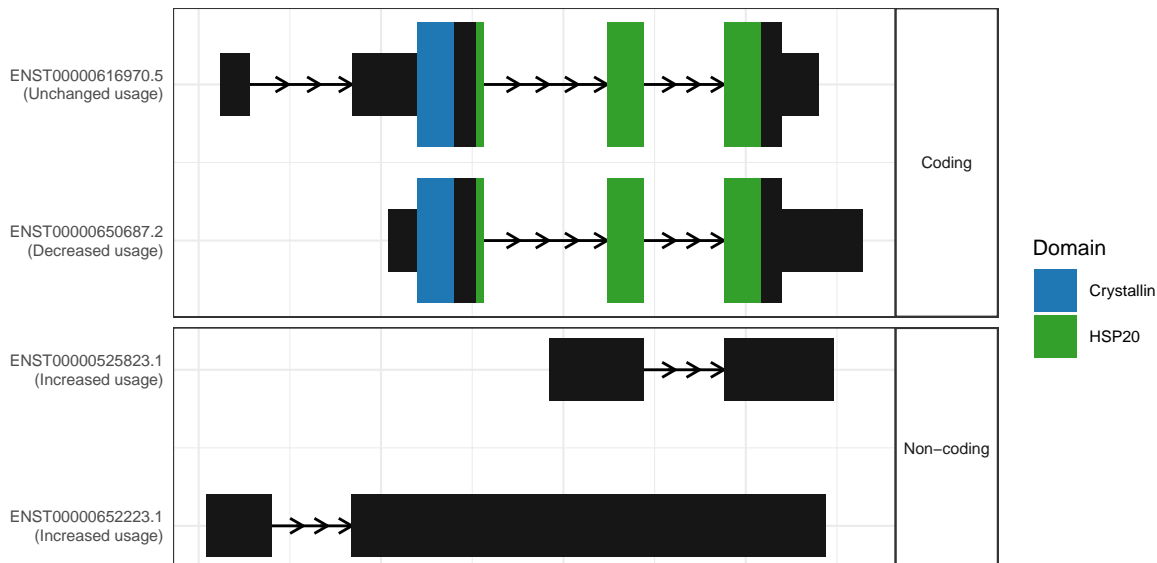
Isoform



Isoform



The isoform switch in CRYAB (FT vs RGC)



Domain

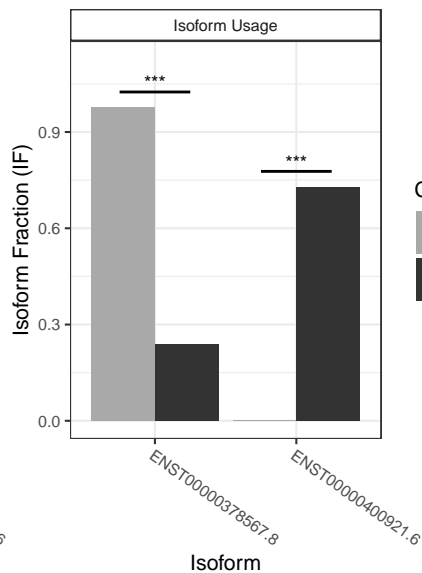
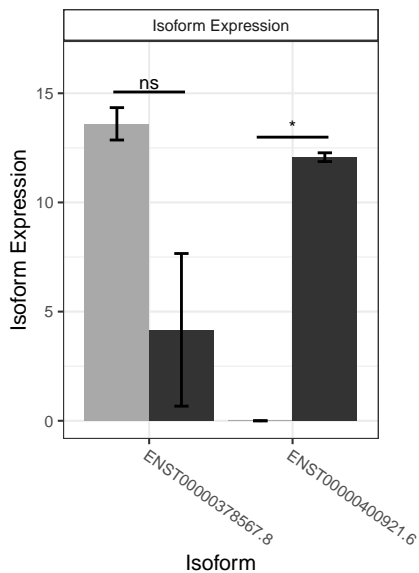
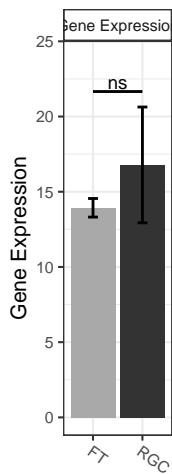
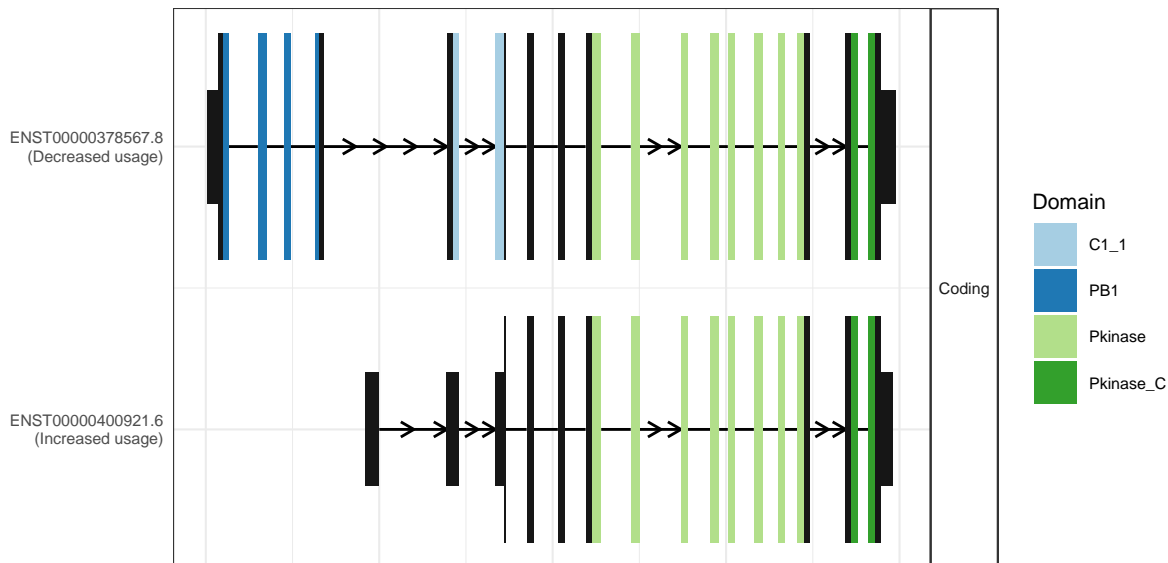


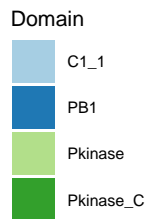
Crystallin



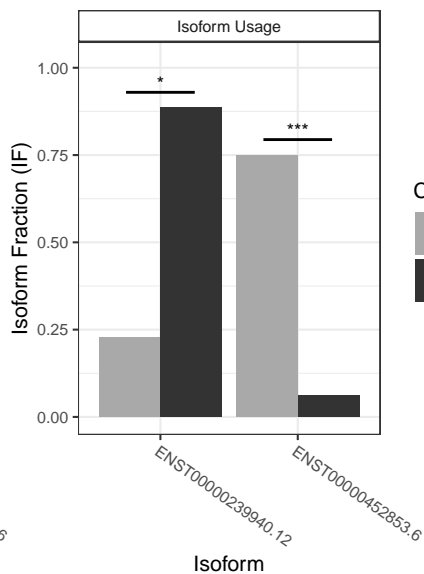
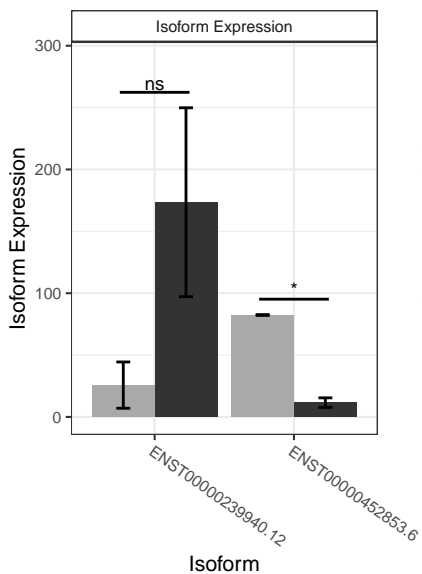
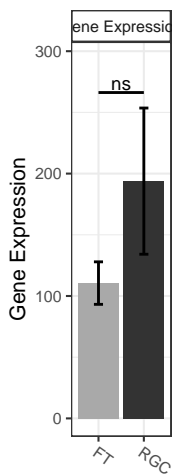
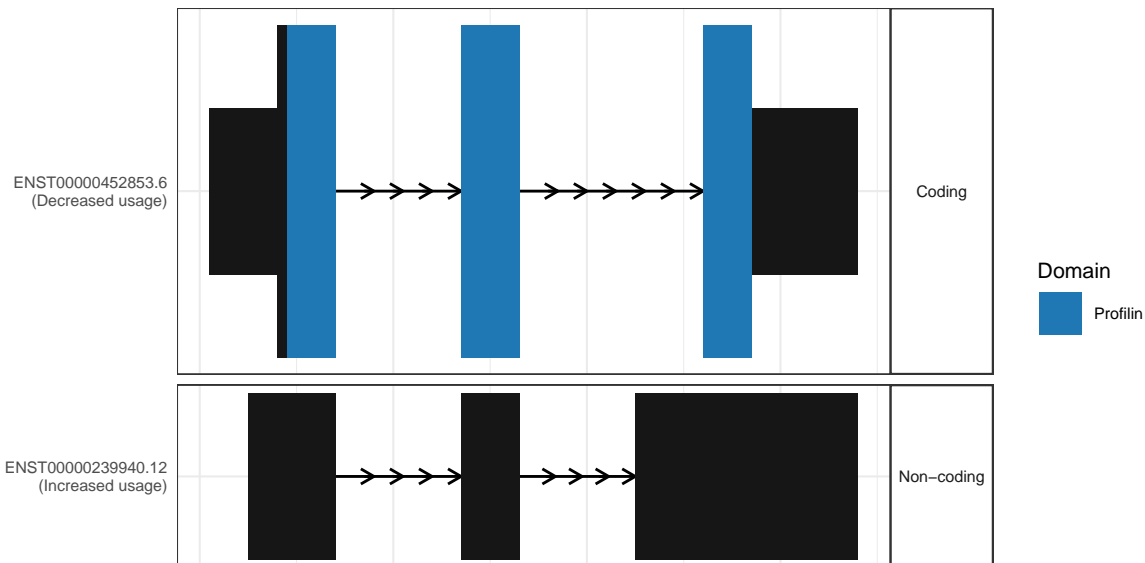
HSP20

The isoform switch in PRKCZ (FT vs RGC)





The isoform switch in PFN2 (FT vs RGC)

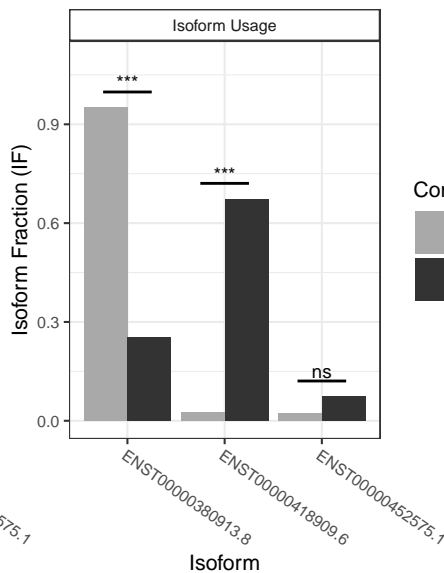
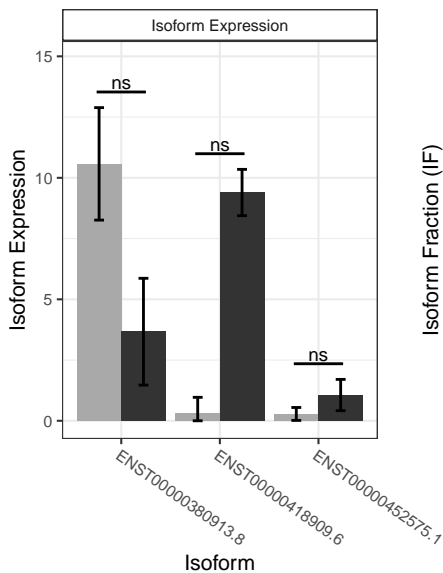
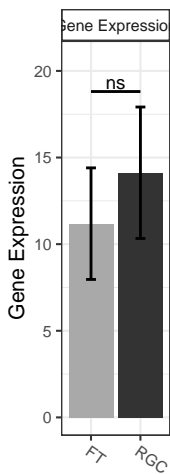
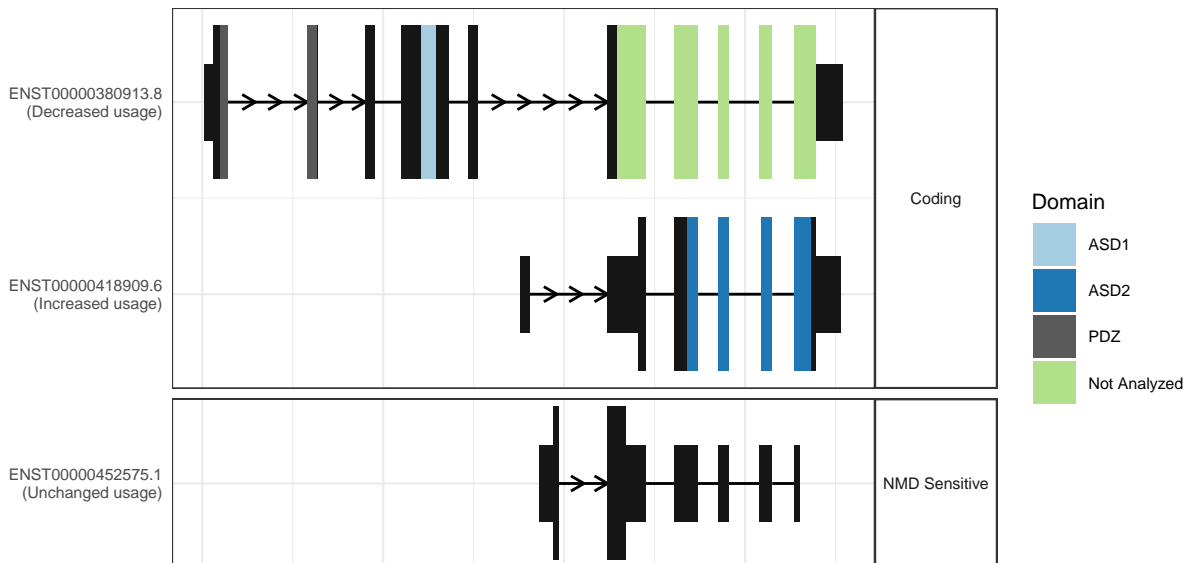


Domain

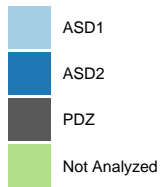


Profilin

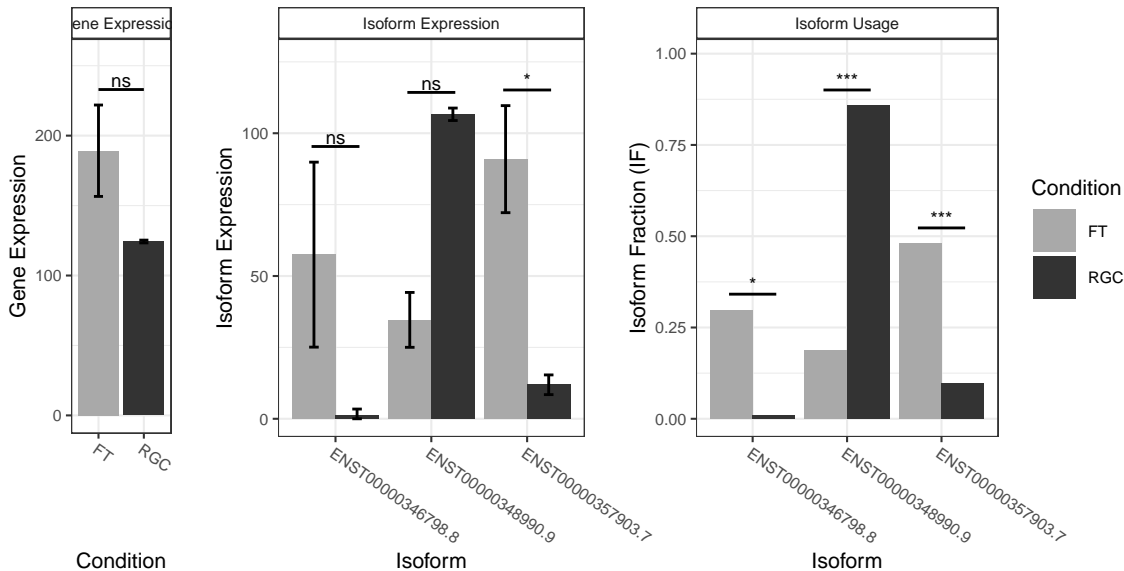
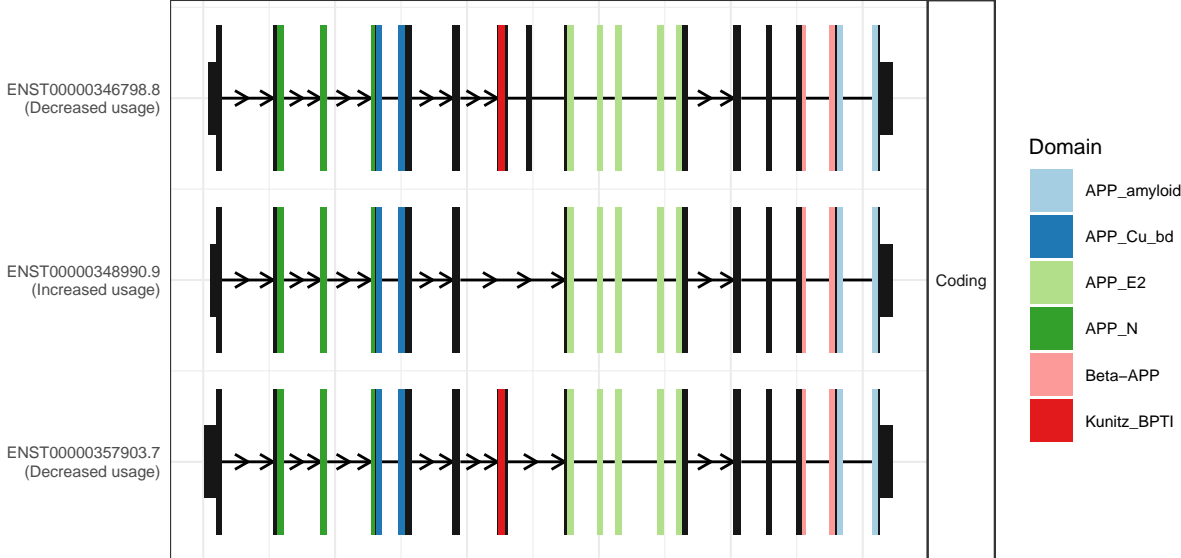
The isoform switch in SHROOM2 (FT vs RGC)









Domain



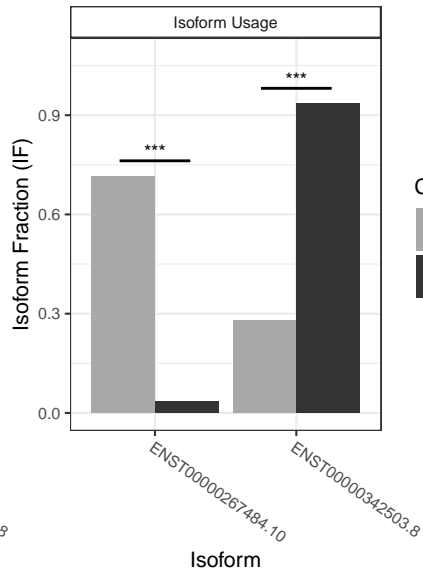
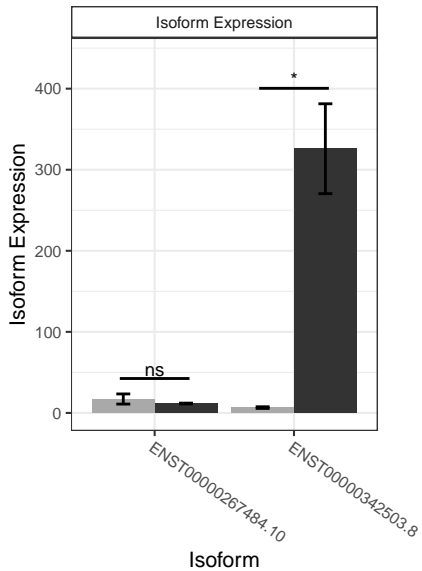
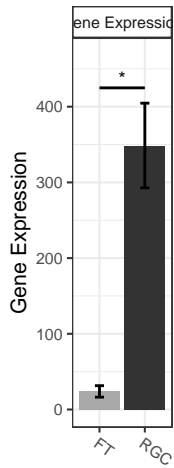
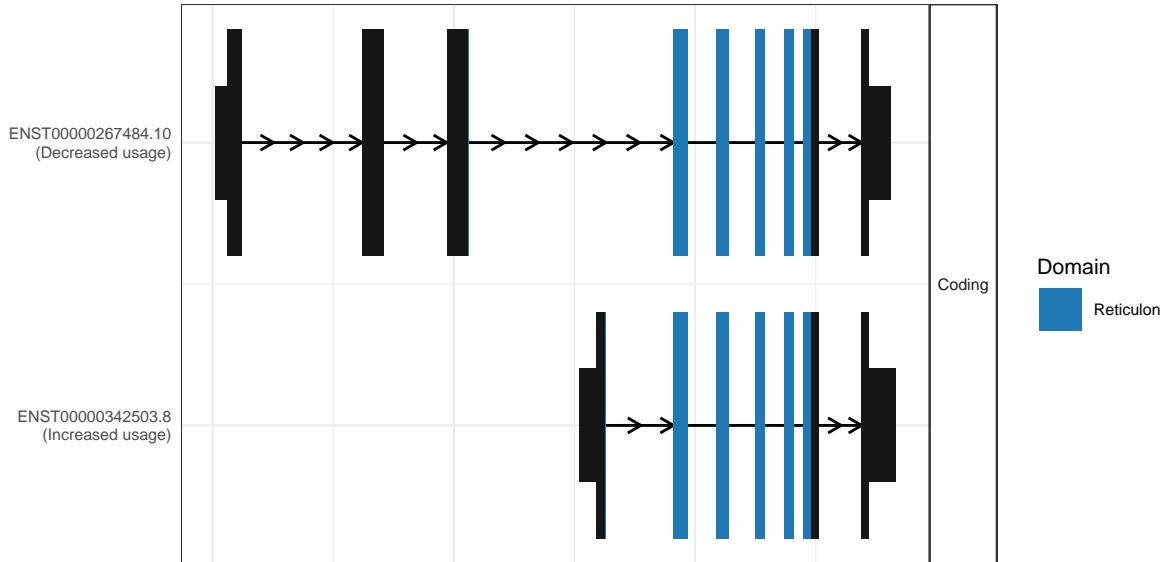
The isoform switch in APP (FT vs RGC)



Domain

	APP_amyloid
	APP_Cu_bd
	APP_E2
	APP_N
	Beta-APP
	Kunitz_BPTI

The isoform switch in RTN1 (FT vs RGC)

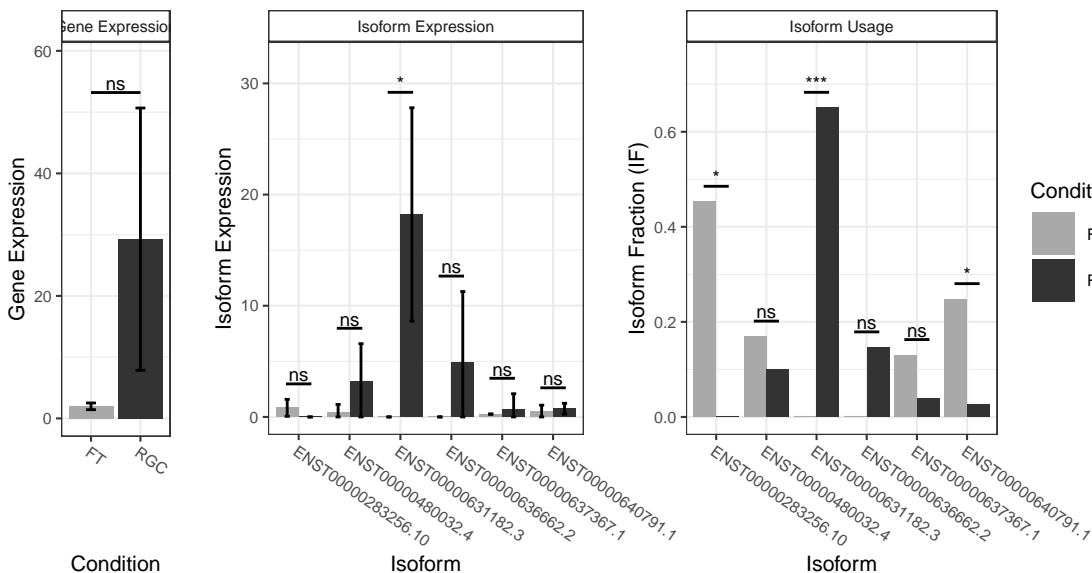
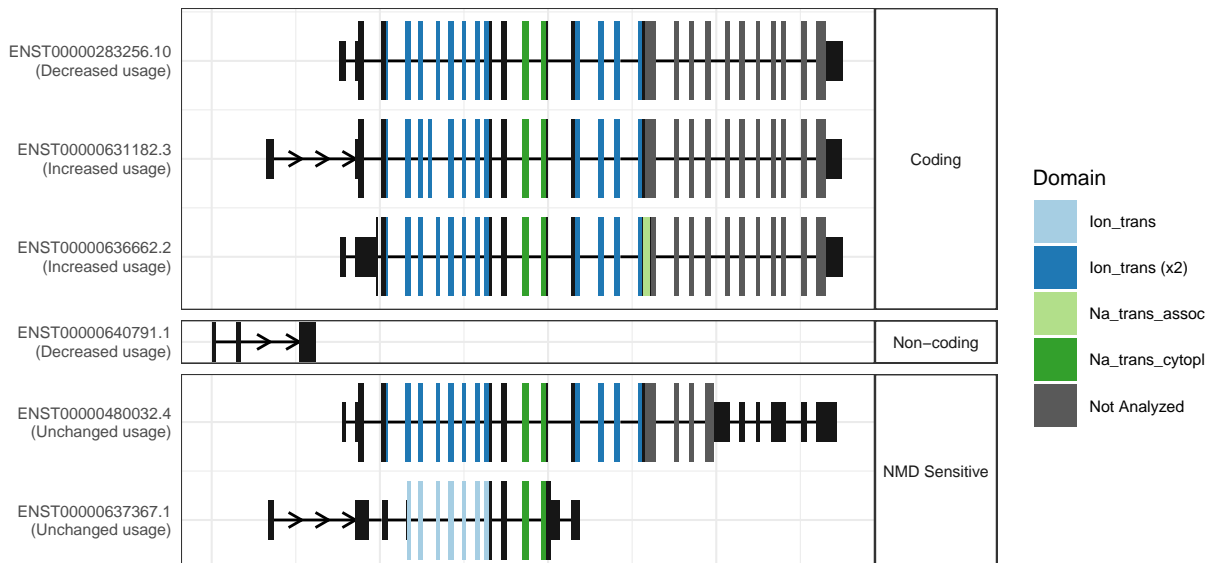


Domain

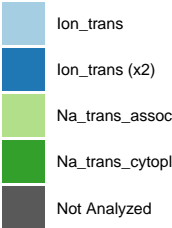


Reticulon

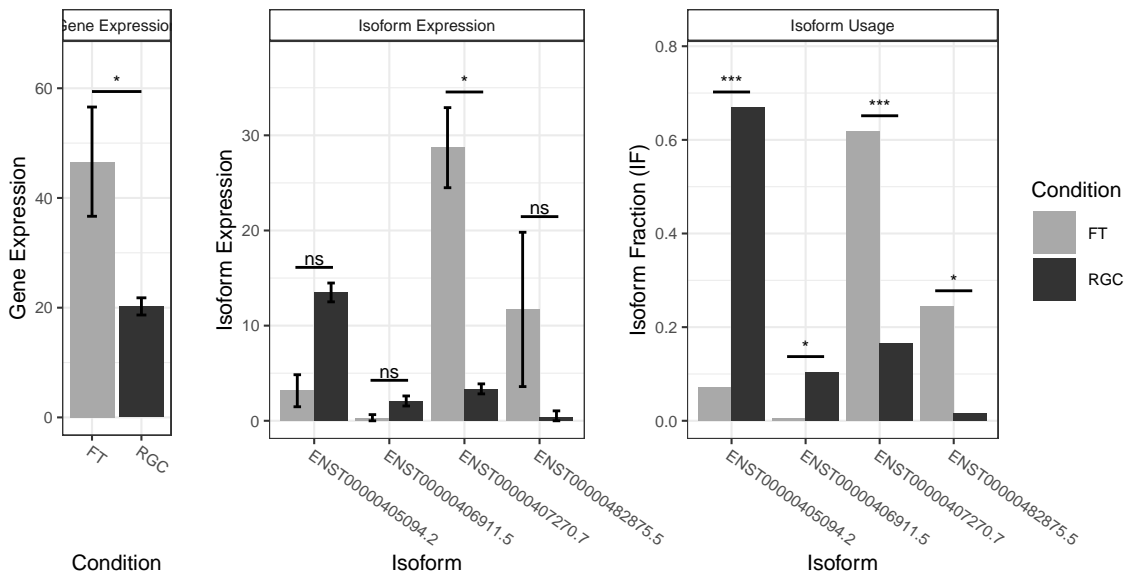
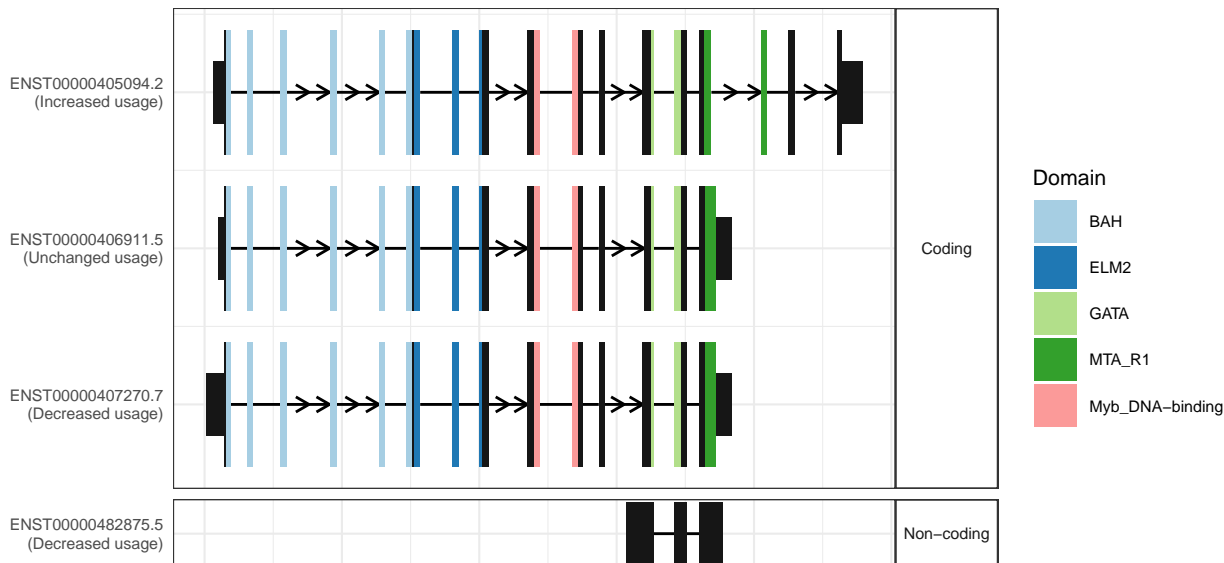
The isoform switch in SCN2A (FT vs RGC)



Domain



The isoform switch in MTA3 (FT vs RGC)



Domain



BAH



ELM2



GATA

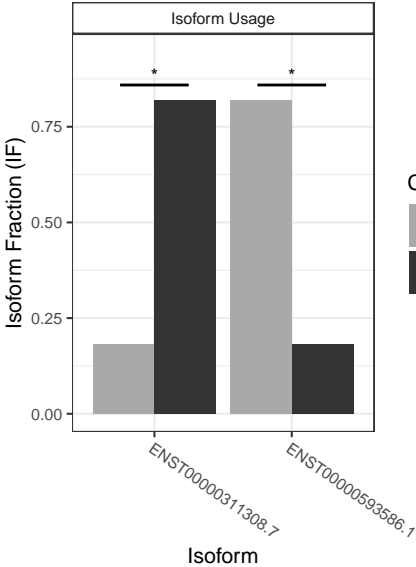
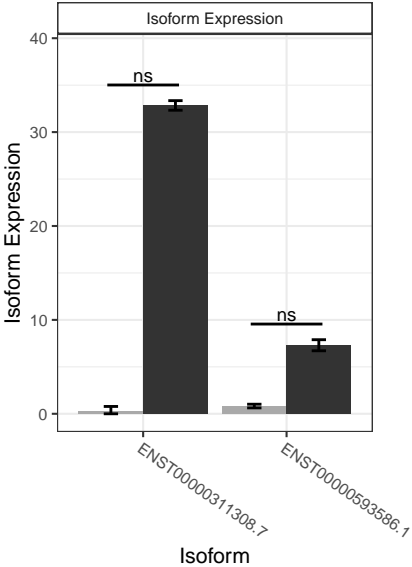
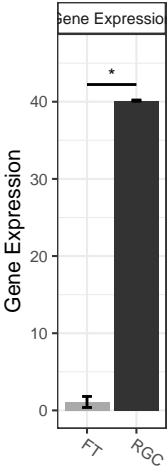
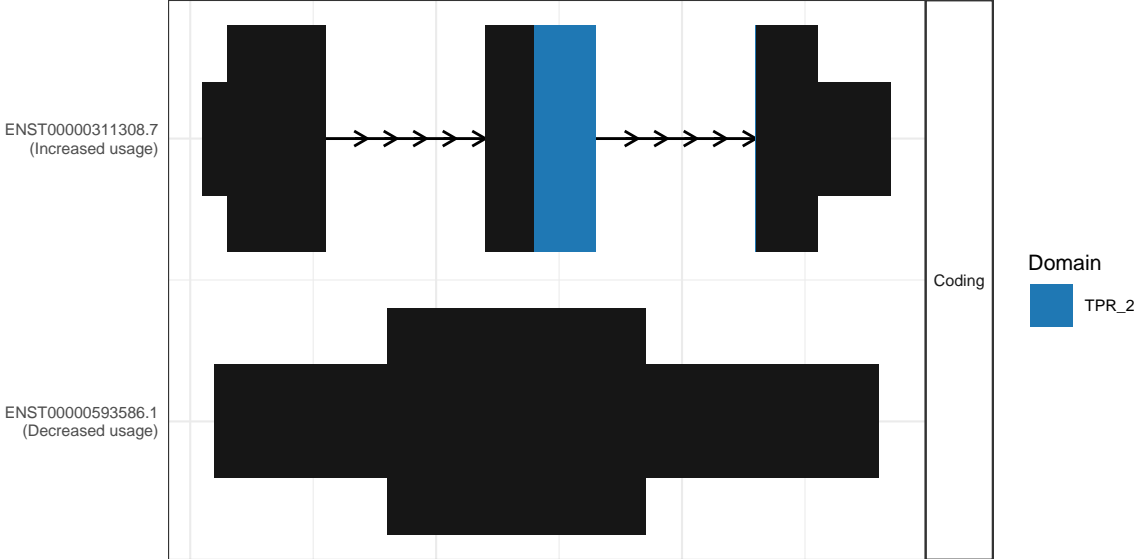


MTA_R1



Myb_DNA-binding

The isoform switch in TTC9B (FT vs RGC)

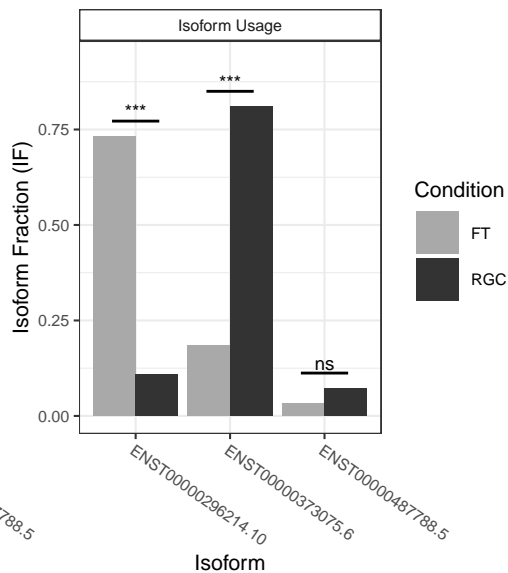
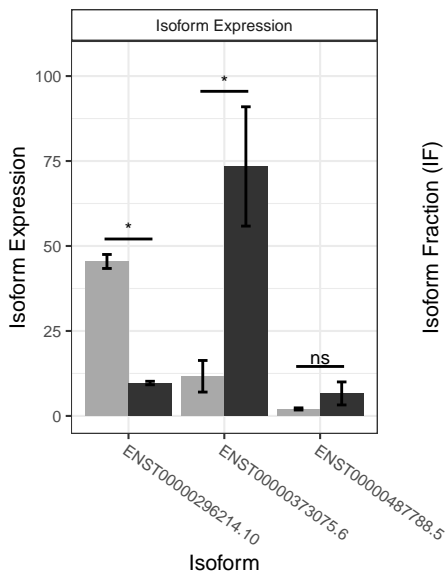
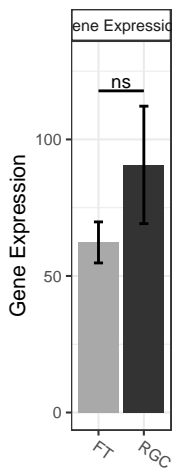
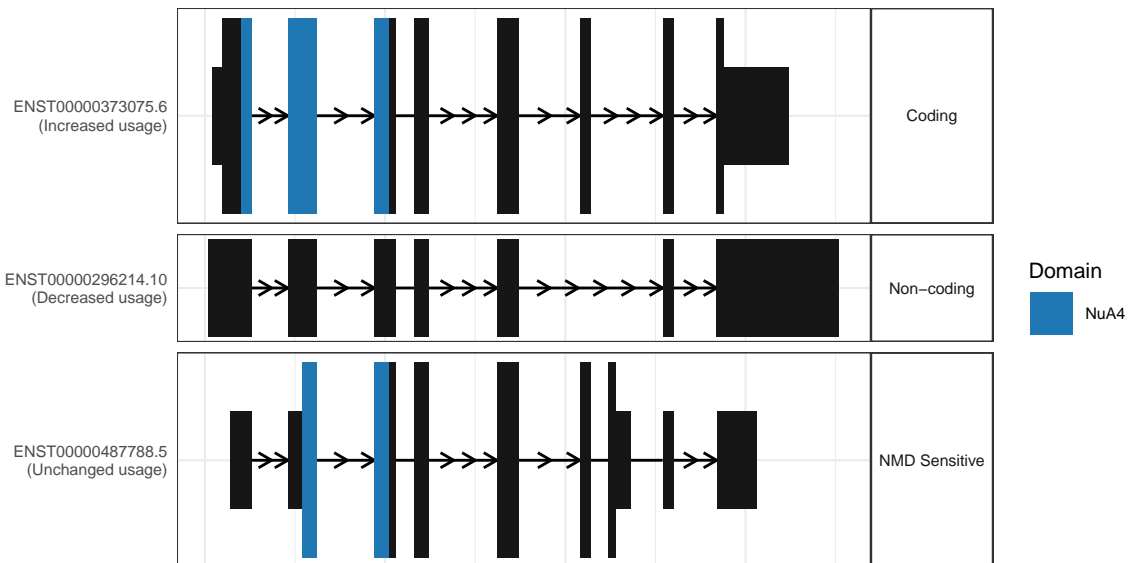


Domain



TPR_2

The isoform switch in MEAF6 (FT vs RGC)

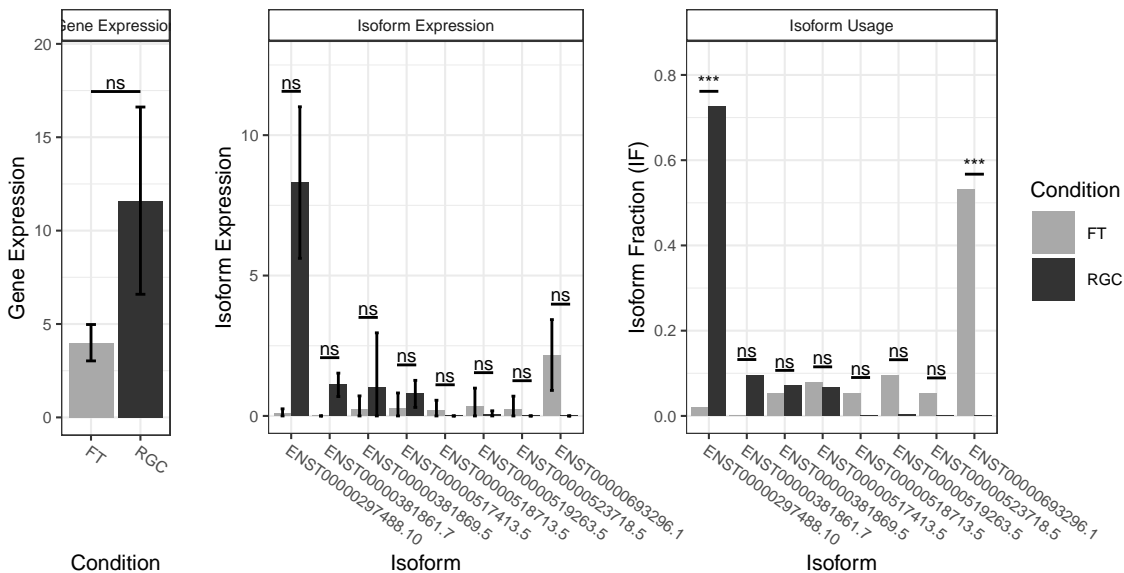
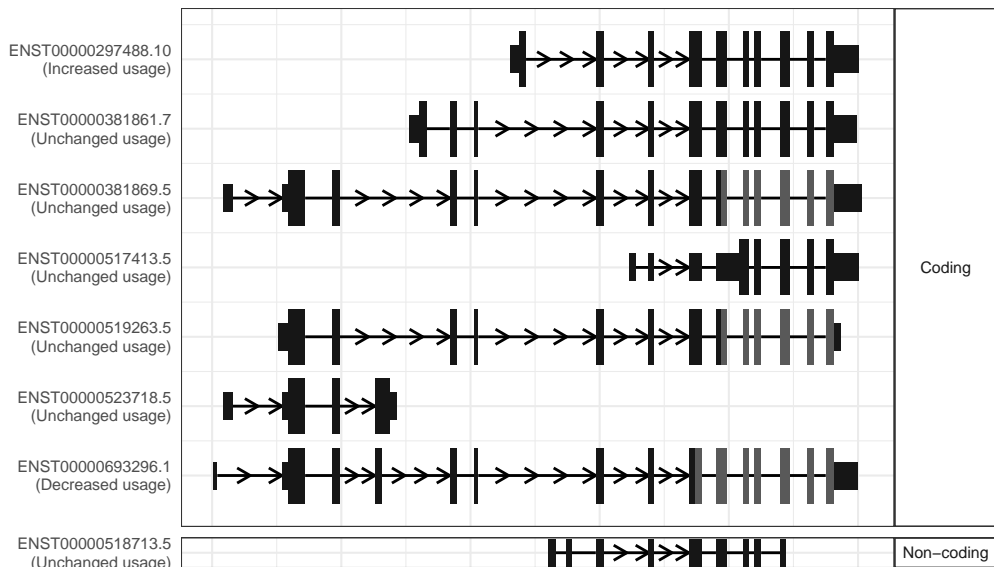


Domain

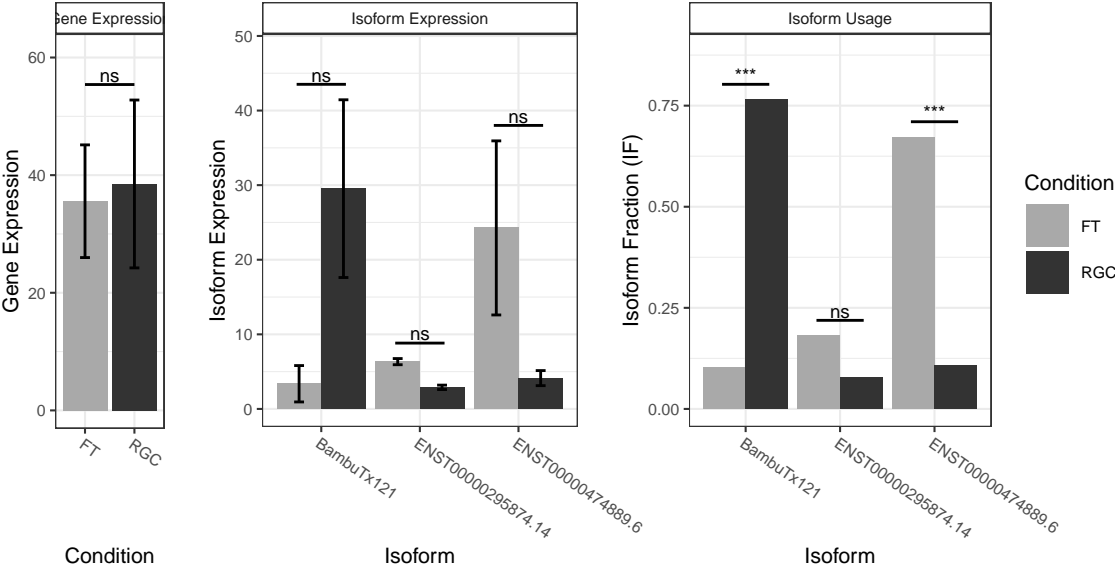
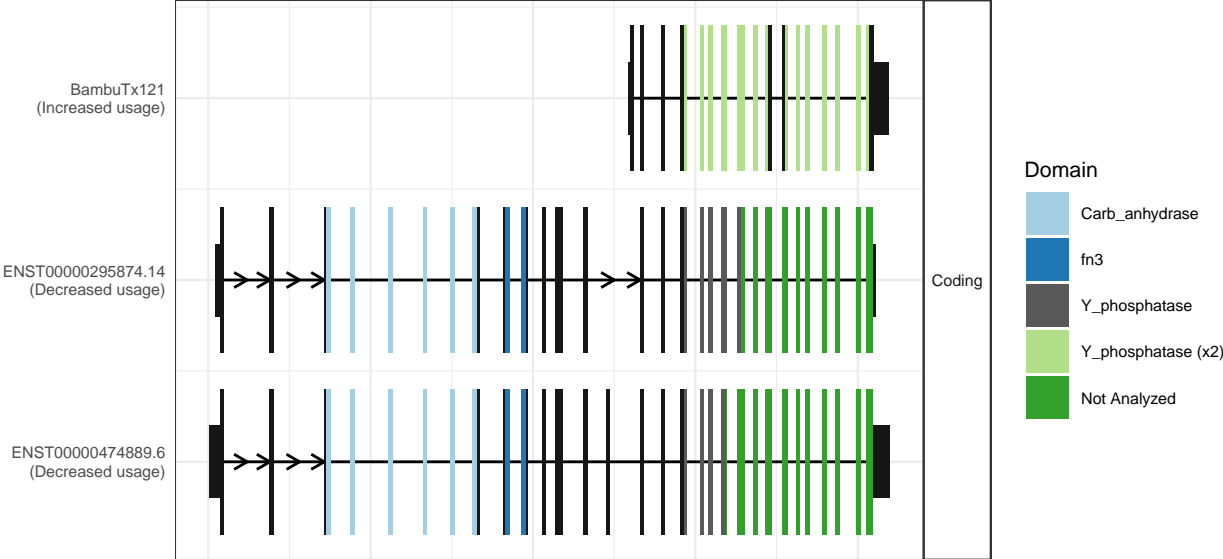


NuA4






The isoform switch in MTUS1 (FT vs RGC)



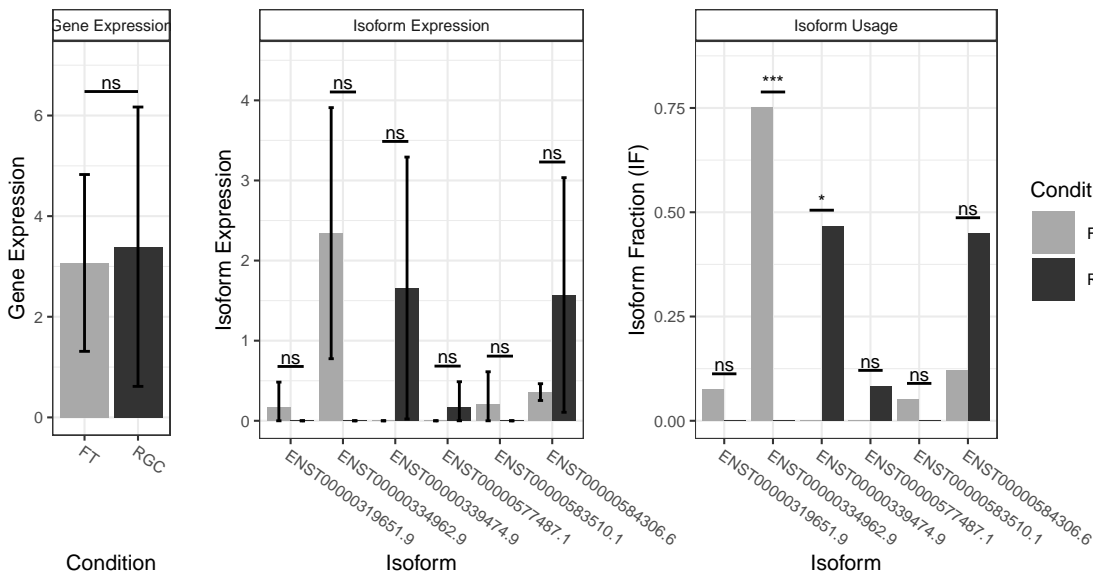
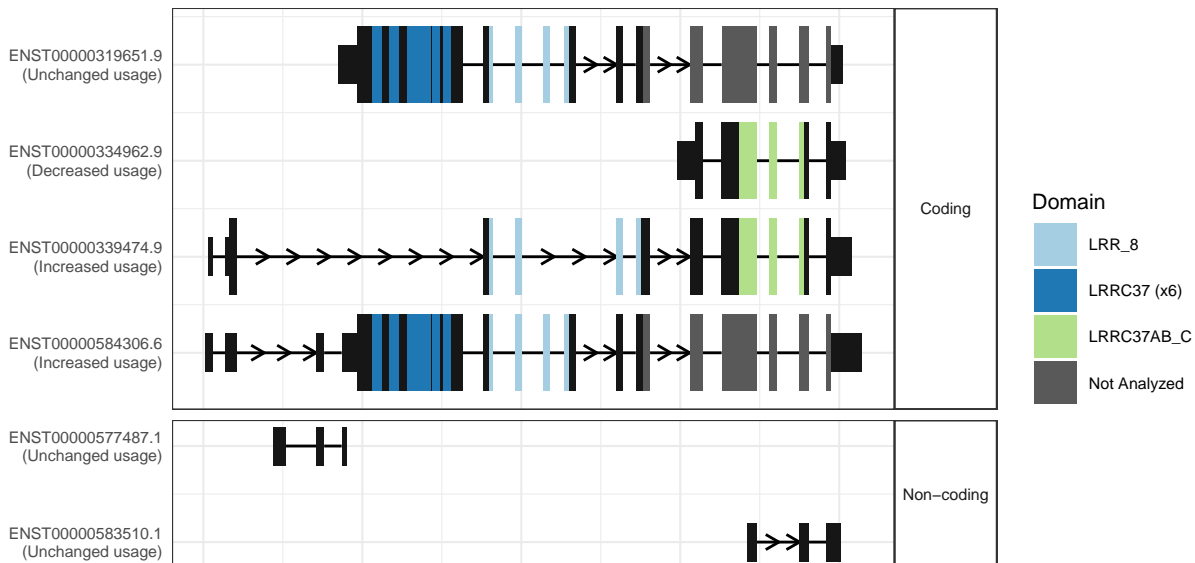
The isoform switch in PTPRG (FT vs RGC)



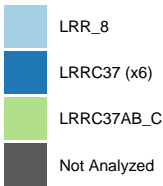
Domain

	Carb_anhydrase
	fn3
	Y_phosphatase
	Y_phosphatase (x2)
	Not Analyzed

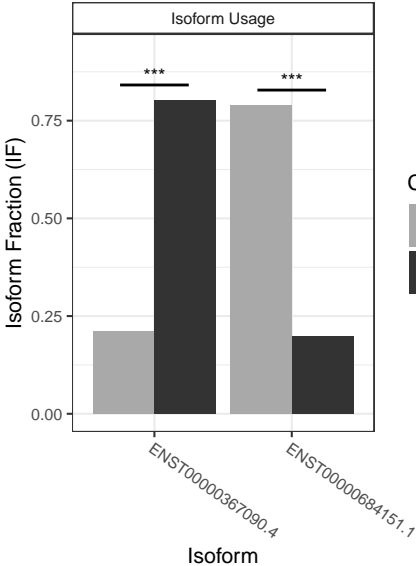
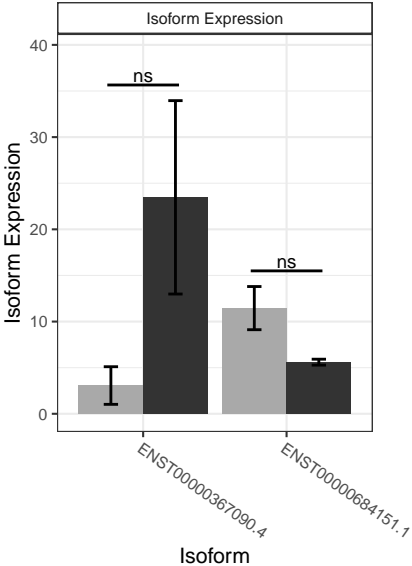
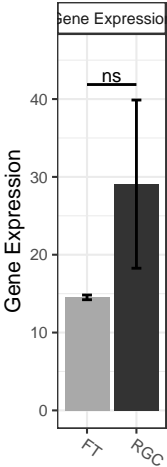
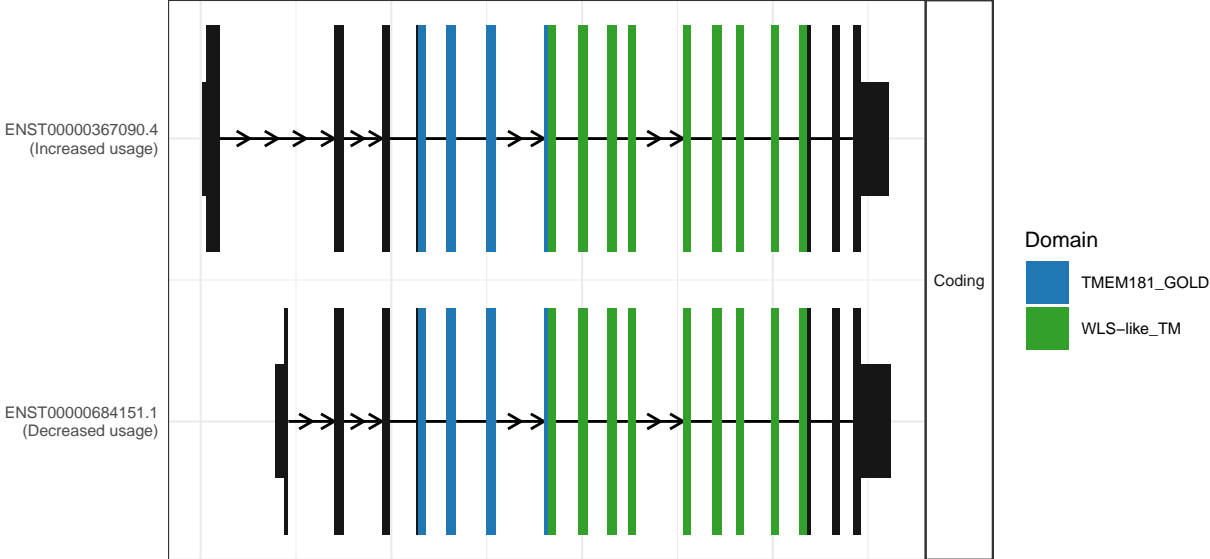
The isoform switch in LRRC37A3 (FT vs RGC)



Domain



The isoform switch in TMEM181 (FT vs RGC)



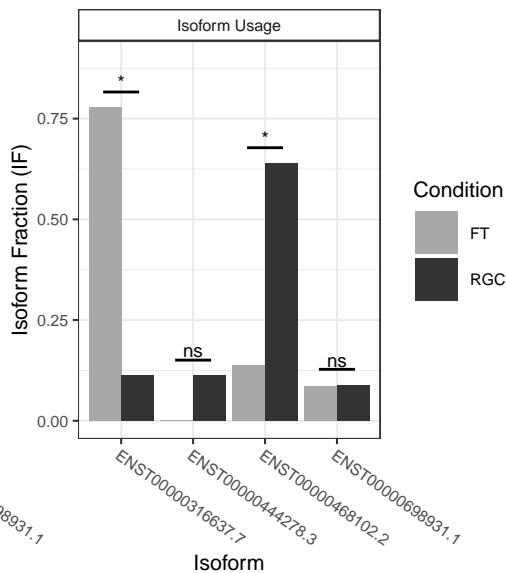
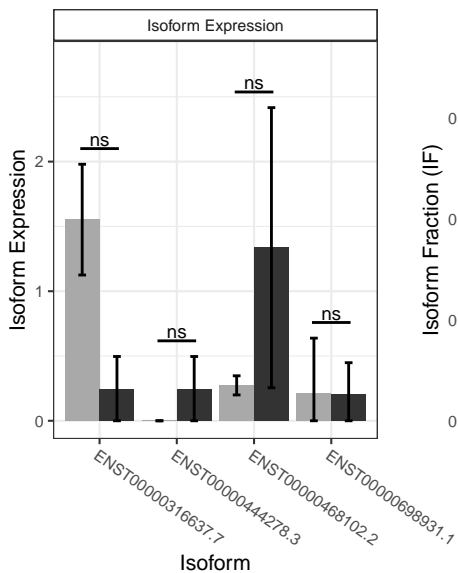
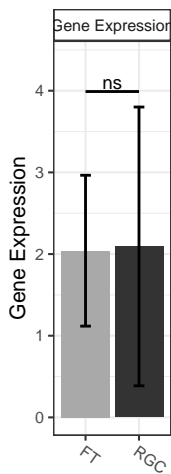
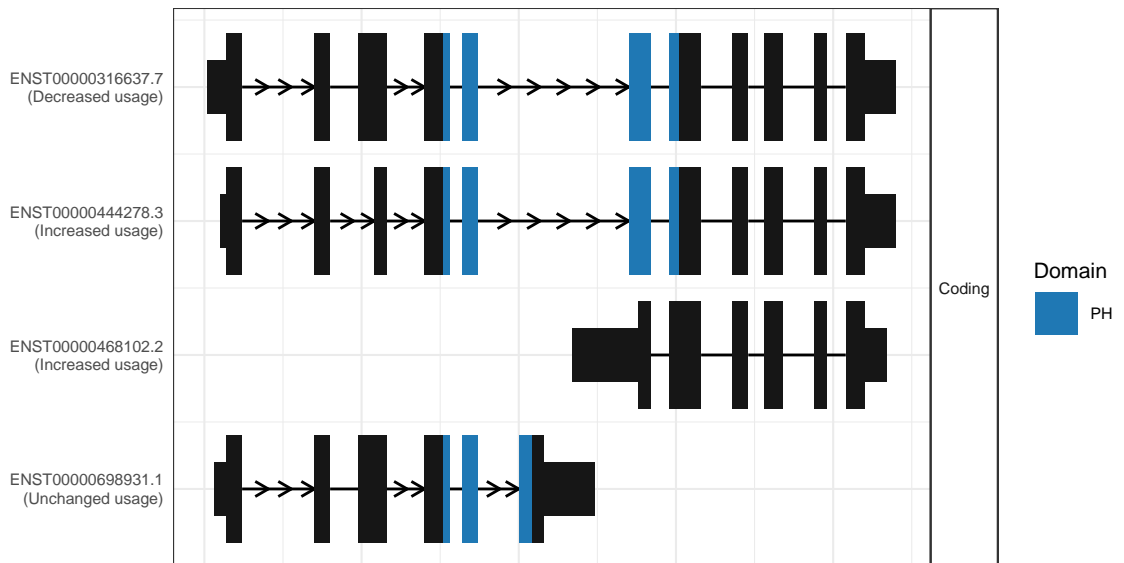
Domain



TMEM181_GOLD

WLS-like_TM

The isoform switch in DEF6 (FT vs RGC)

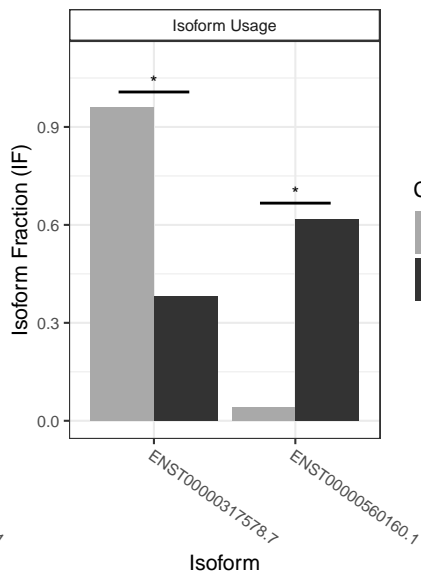
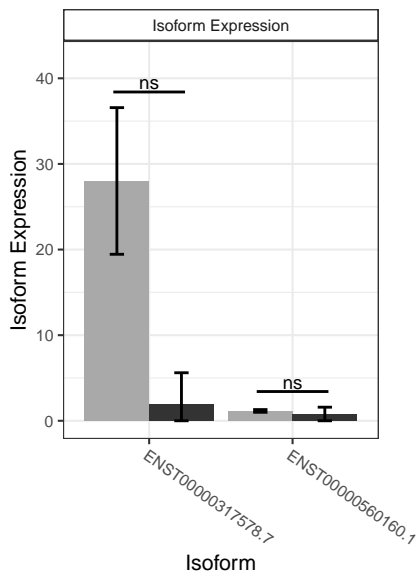
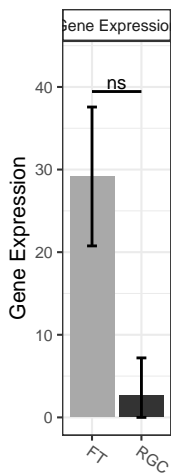
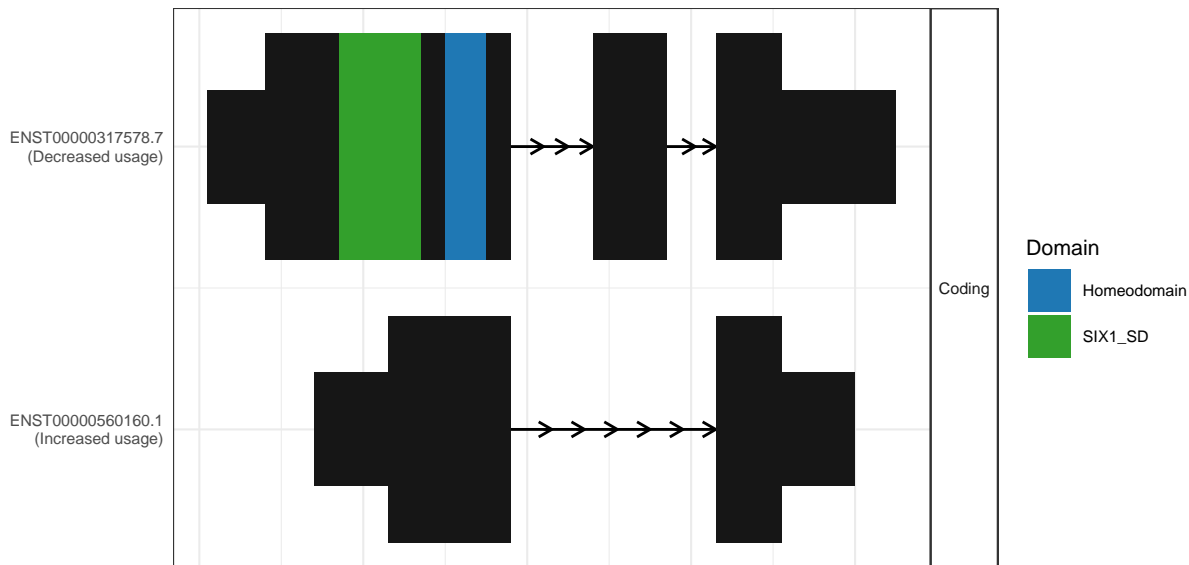


Domain



PH

The isoform switch in SIX5 (FT vs RGC)



Domain

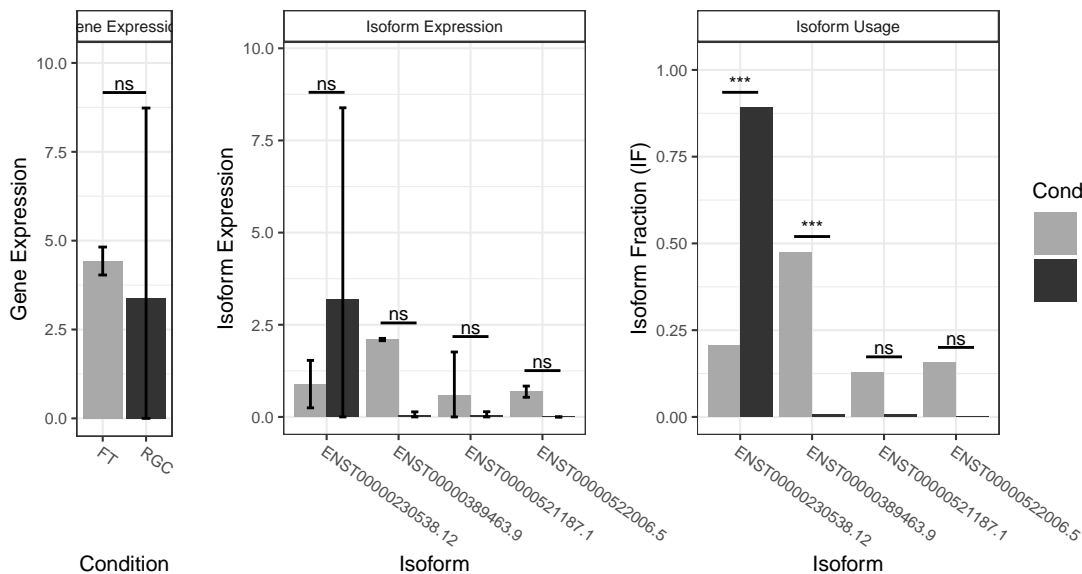
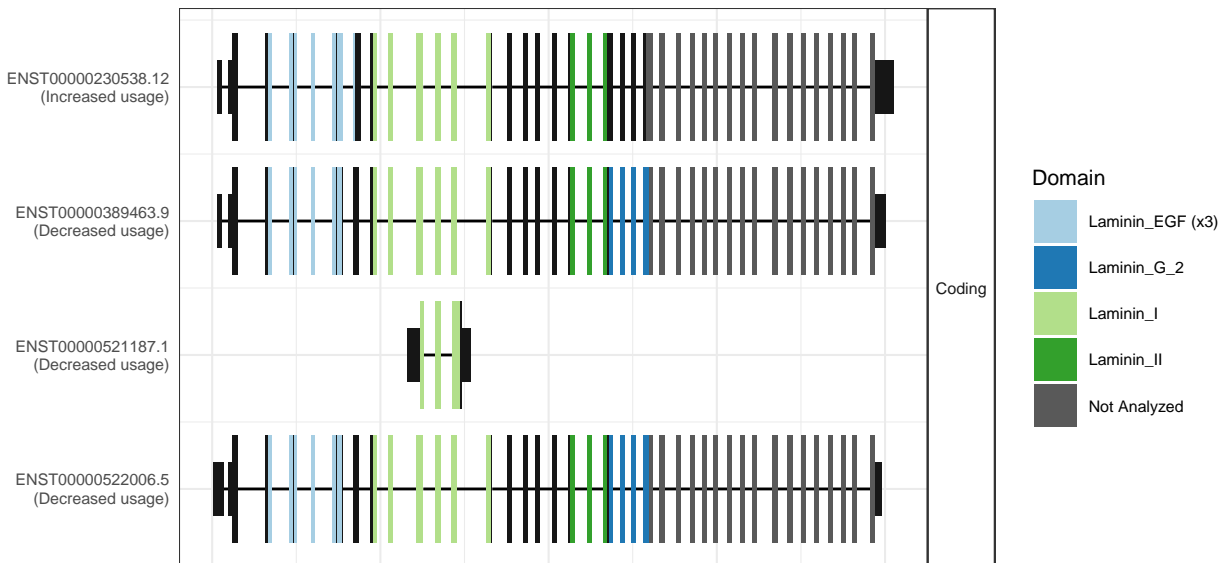


Homeodomain







SIX1_SD

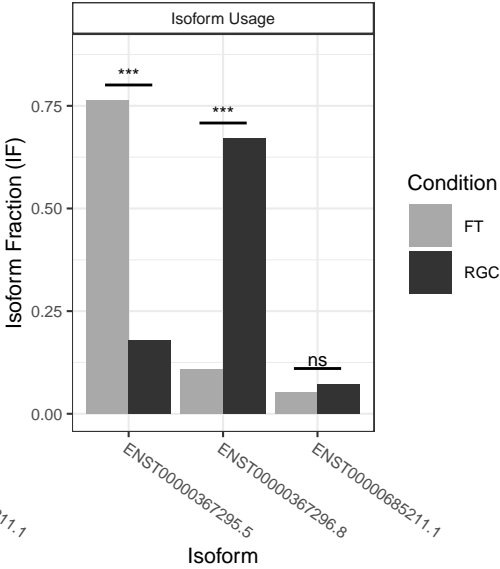
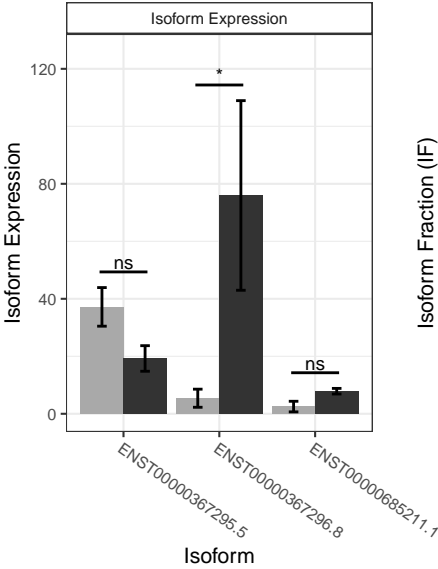
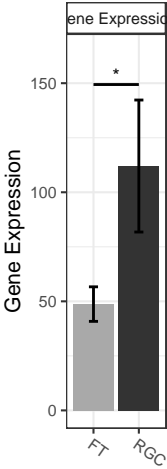
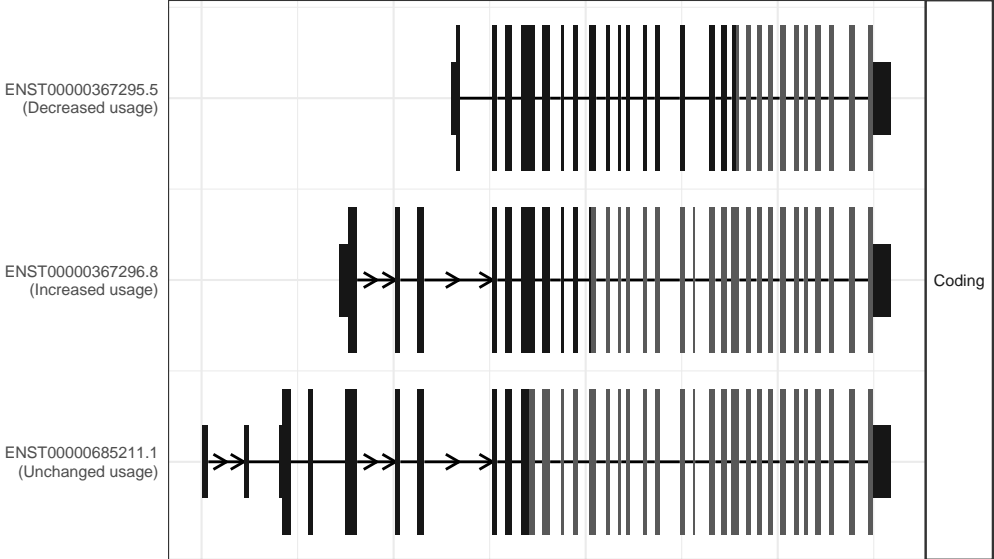
The isoform switch in LAMA4 (FT vs RGC)



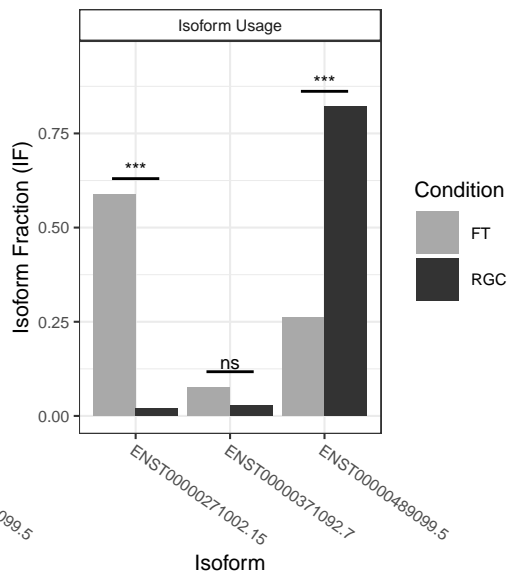
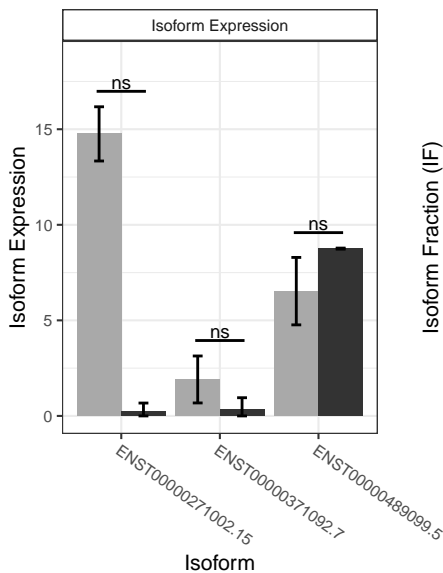
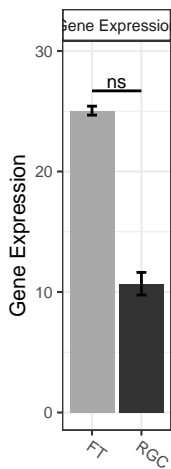
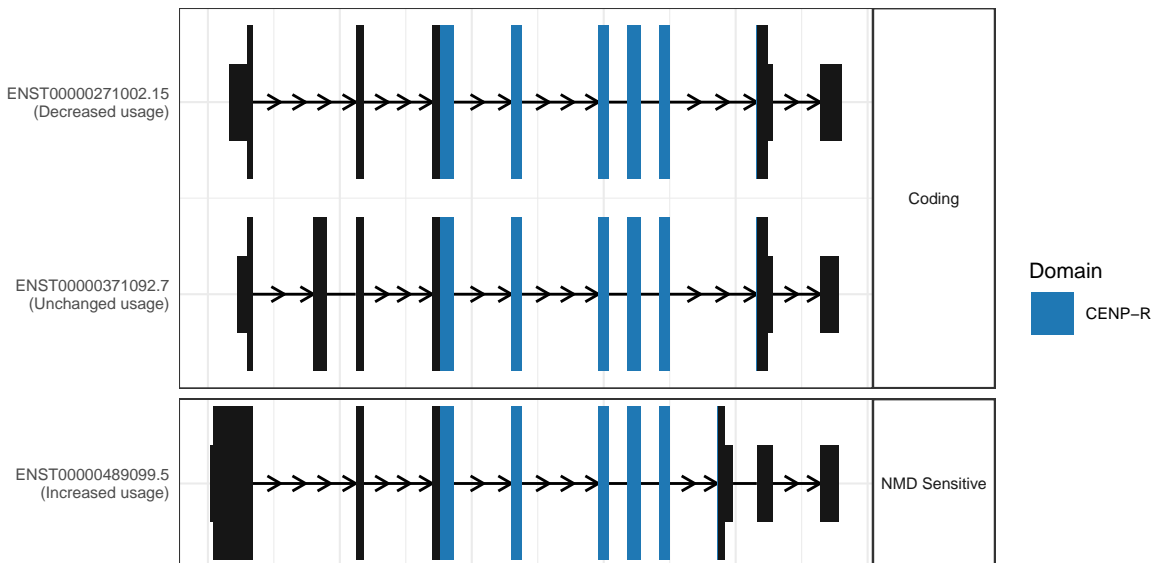
Domain

	Laminin_EGF (x3)
	Laminin_G_2
	Laminin_I
	Laminin_II
	Not Analyzed

The isoform switch in NAV1 (FT vs RGC)



The isoform switch in ITGB3BP (FT vs RGC)

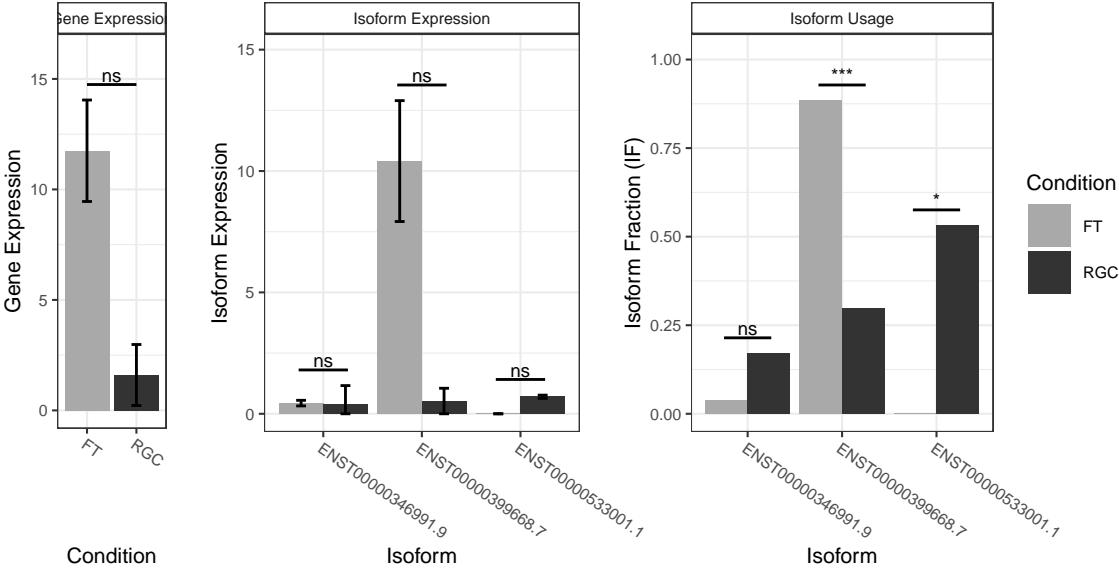
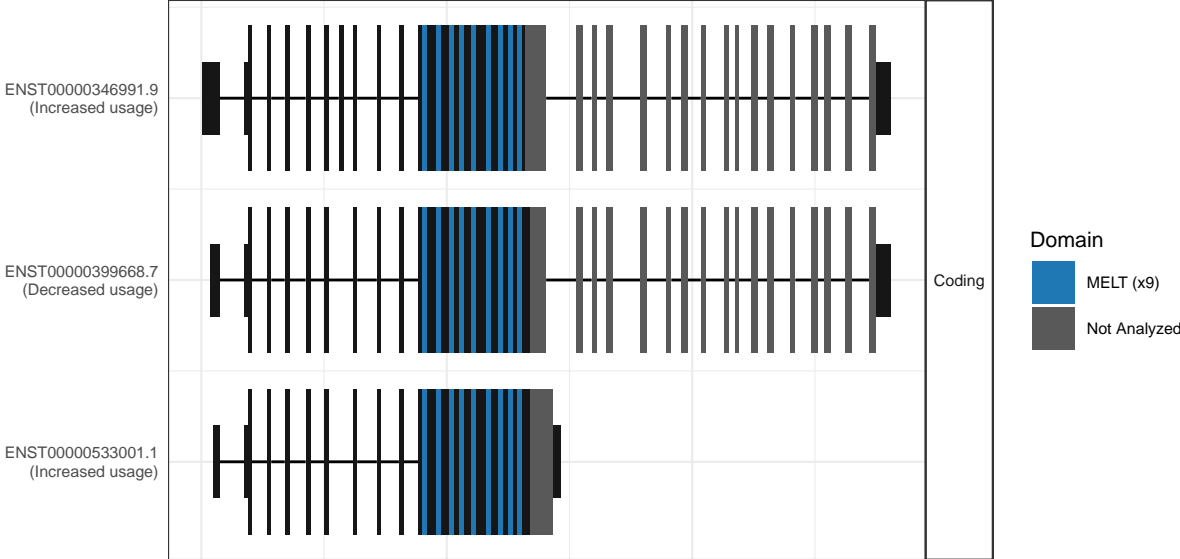


Domain



CENP-R

The isoform switch in KNL1 (FT vs RGC)



Domain

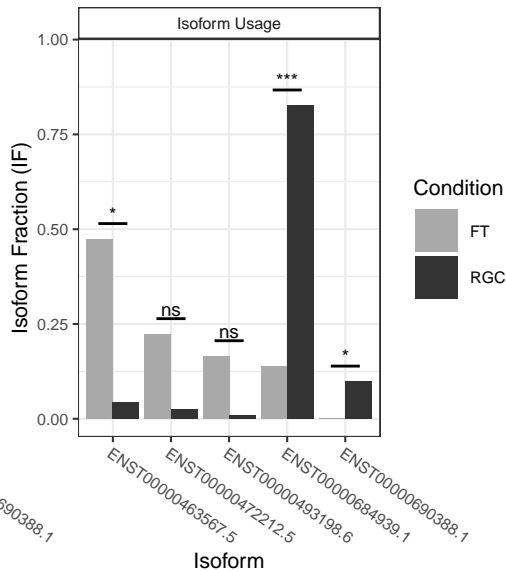
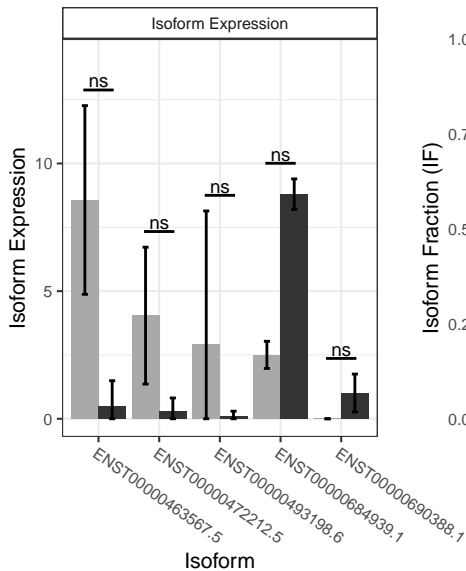
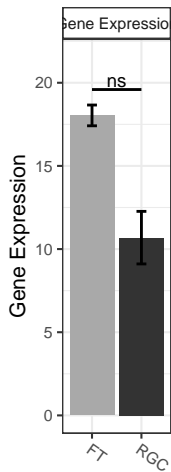
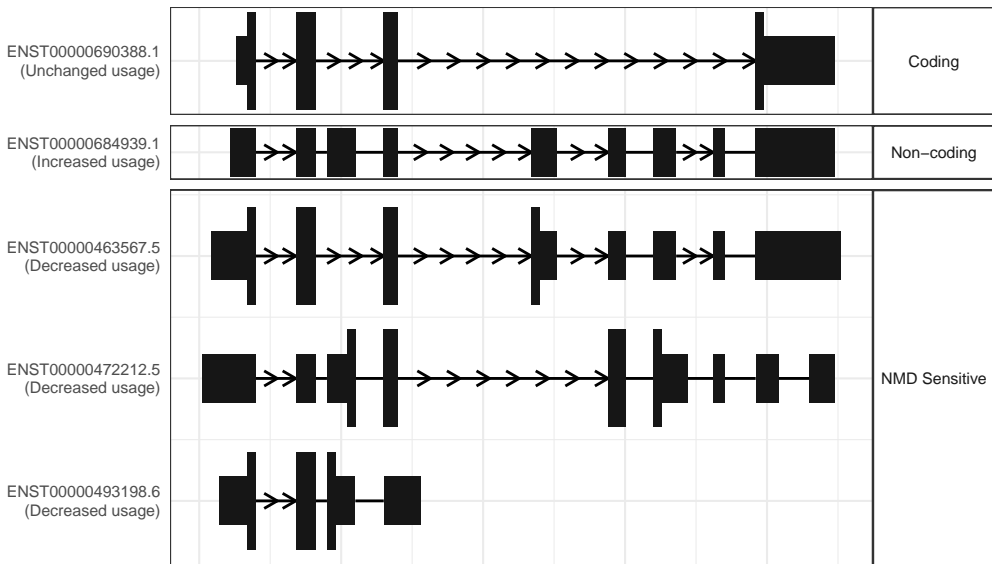


MELT (x9)



Not Analyzed

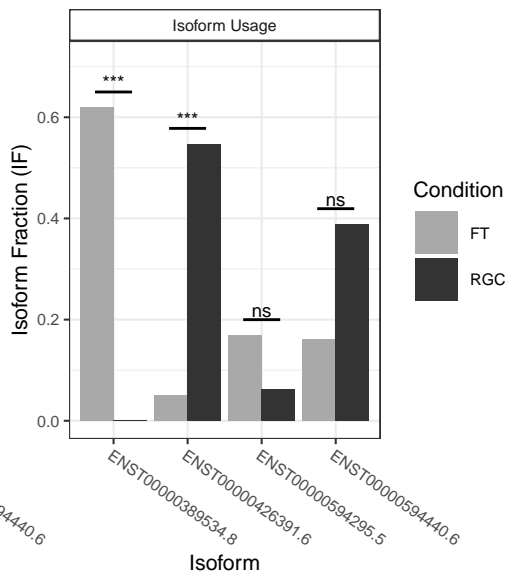
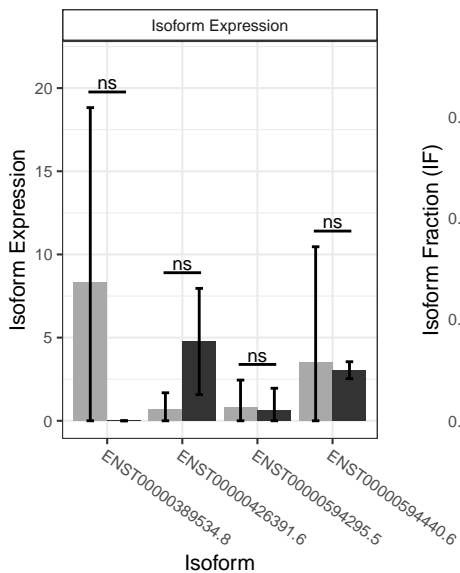
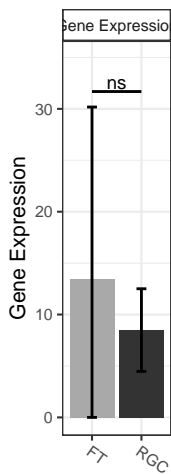
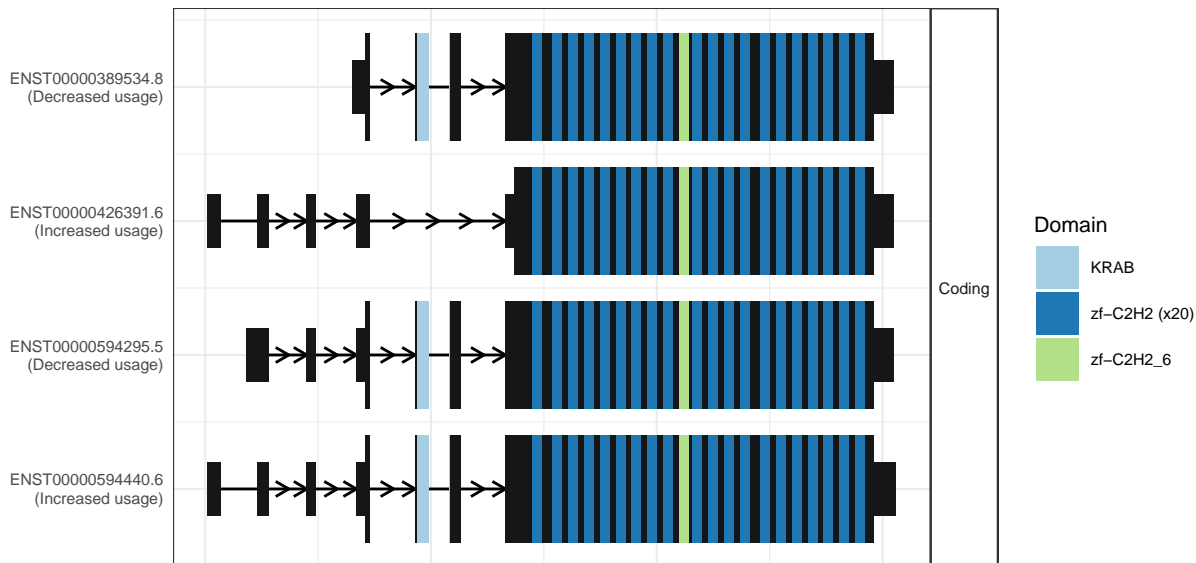
The isoform switch in (FT vs RGC)



Condition



The isoform switch in ZNF841 (FT vs RGC)



Domain



KRAB

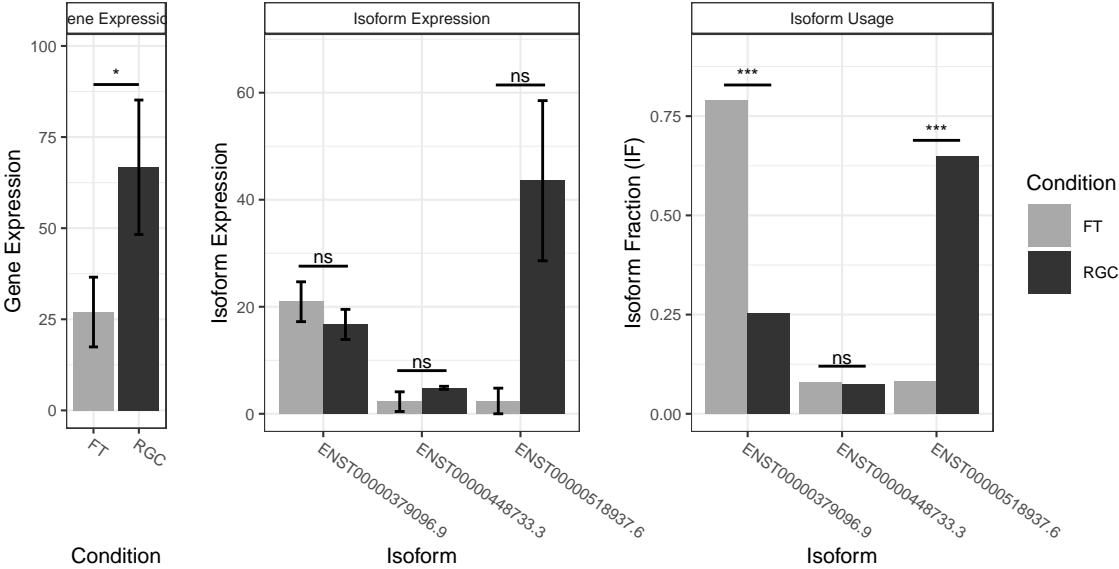
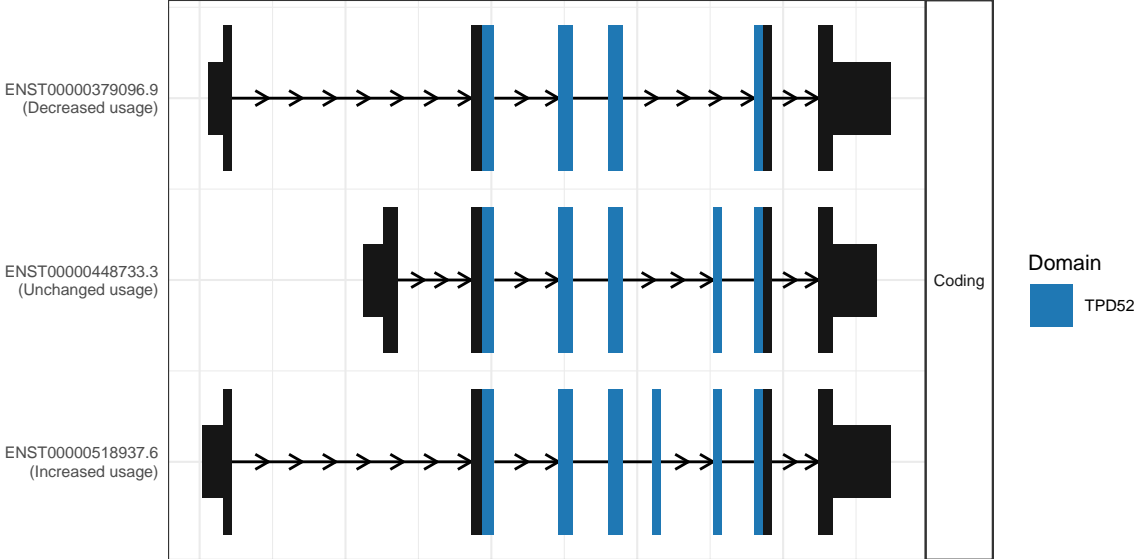


zf-C2H2 (x20)



zf-C2H2_6

The isoform switch in TPD52 (FT vs RGC)

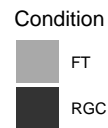
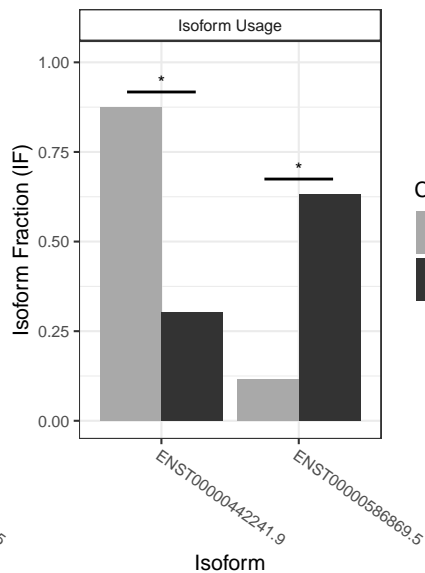
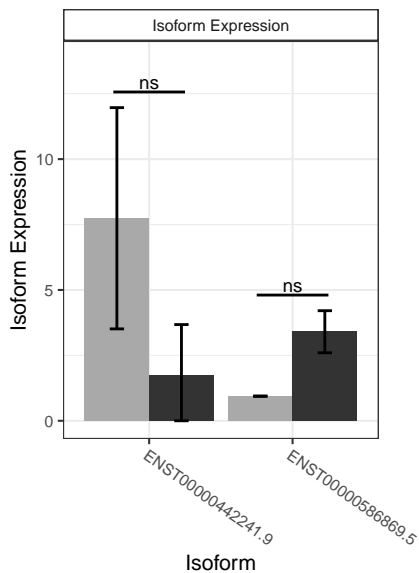
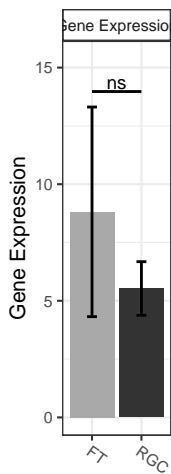
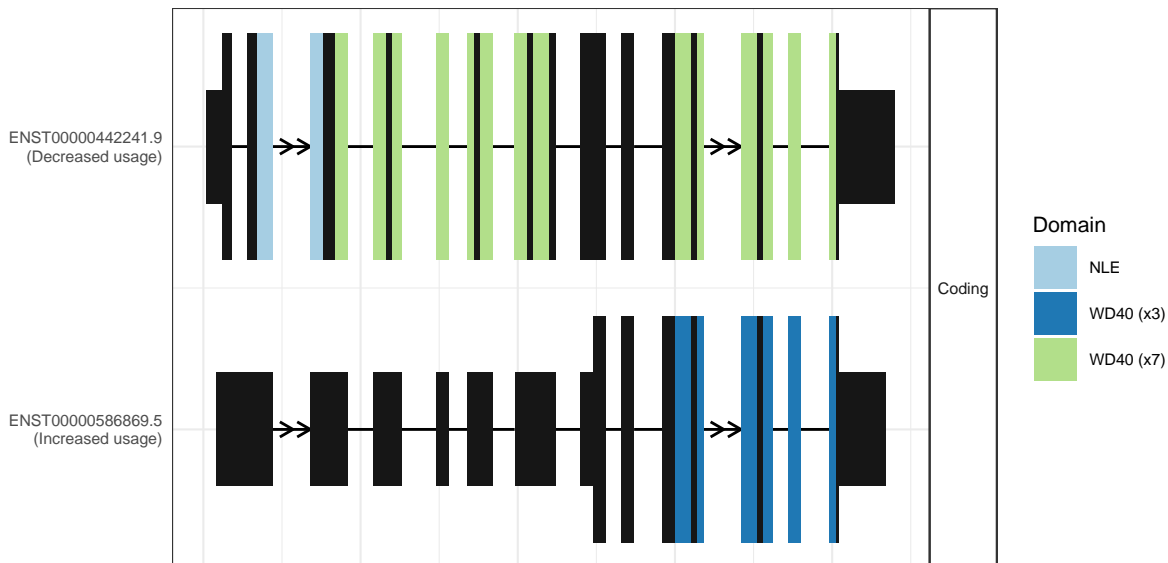


Domain



TPD52

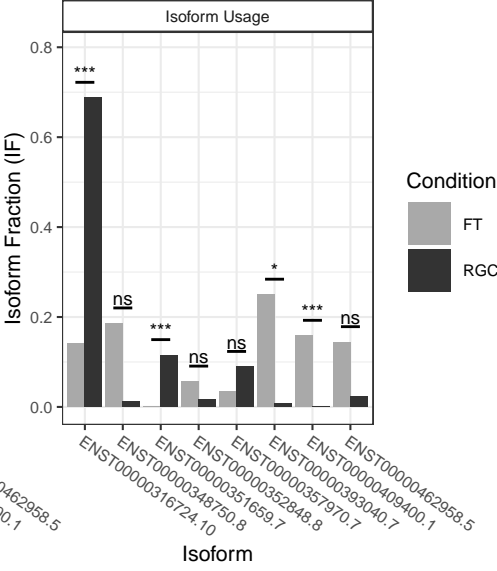
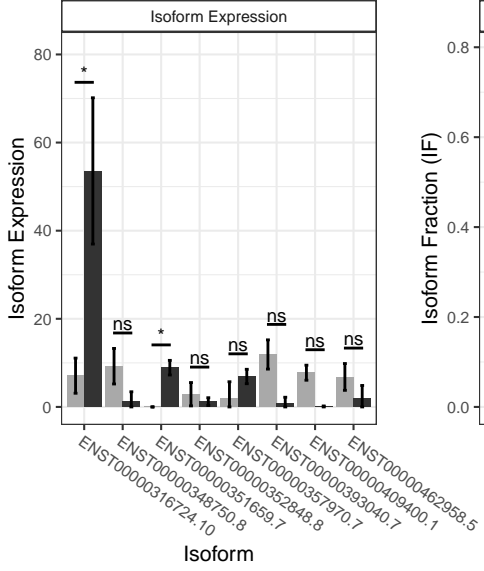
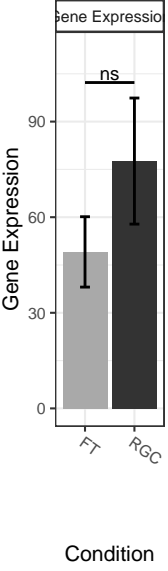
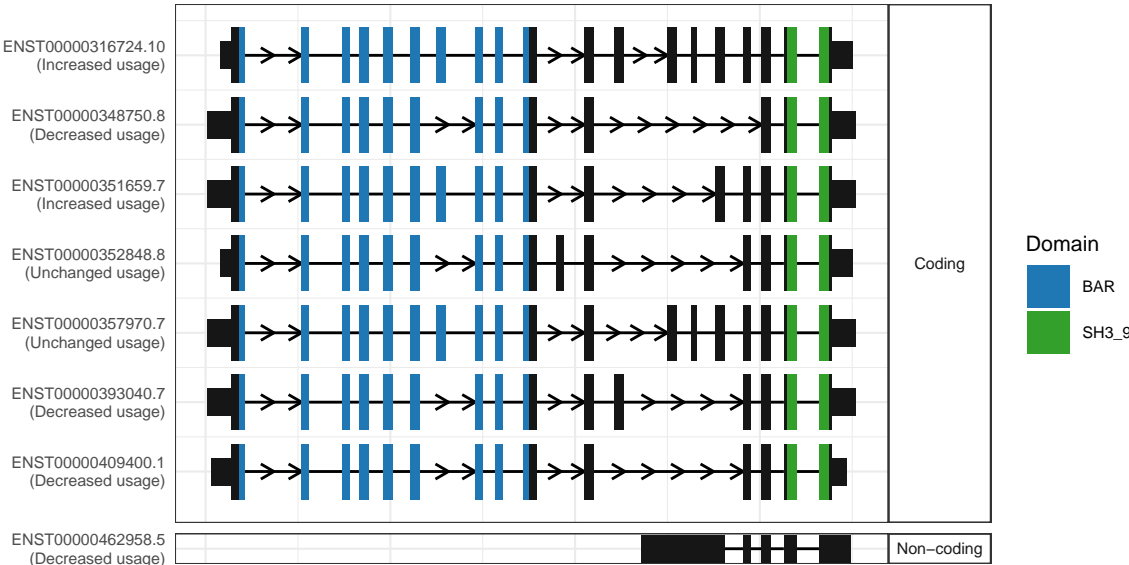
The isoform switch in NLE1 (FT vs RGC)



Domain

	NLE
	WD40 (x3)
	WD40 (x7)

The isoform switch in BIN1 (FT vs RGC)



Domain

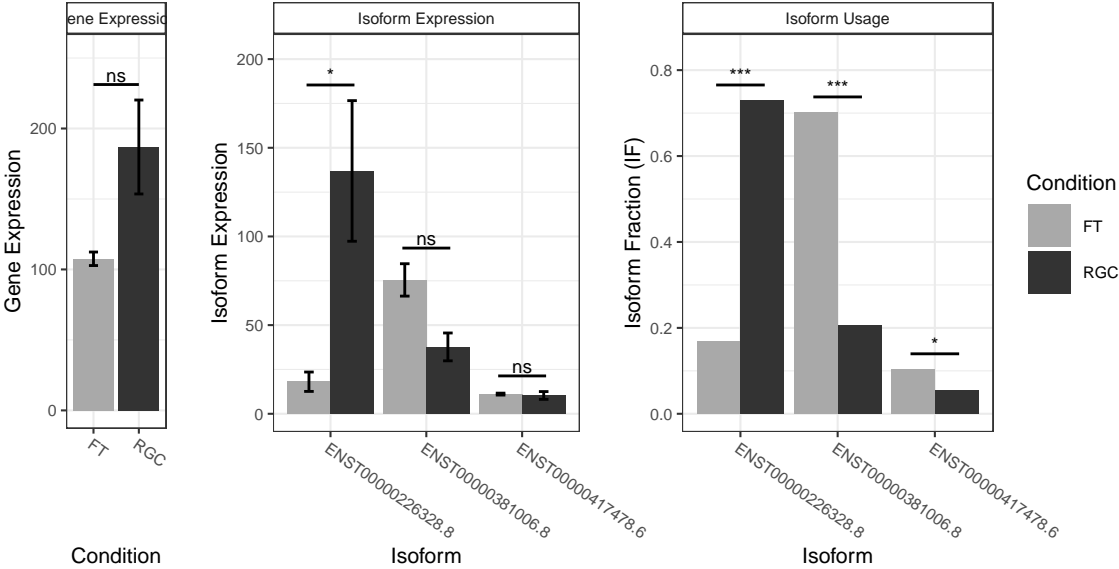
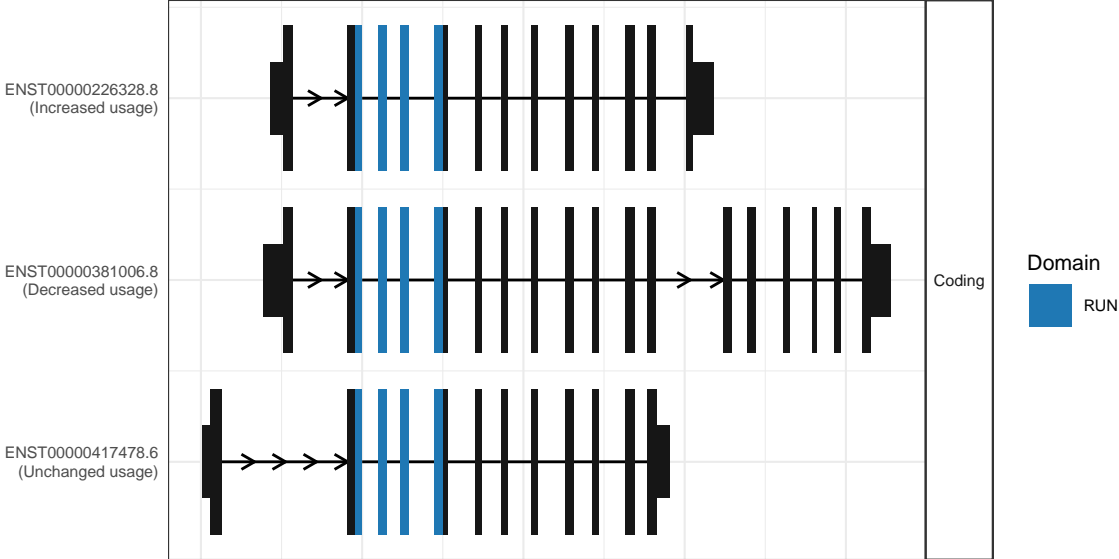


BAR



SH3_9

The isoform switch in RUFY3 (FT vs RGC)

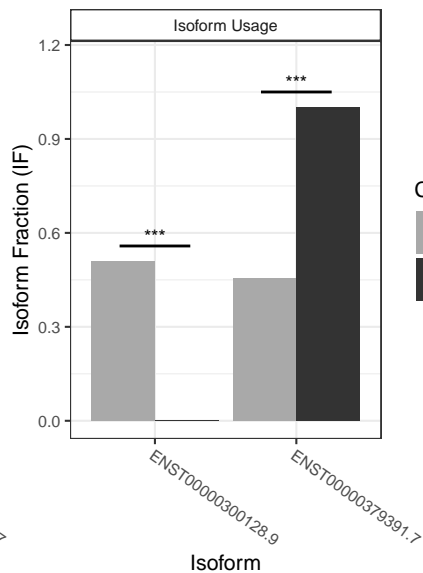
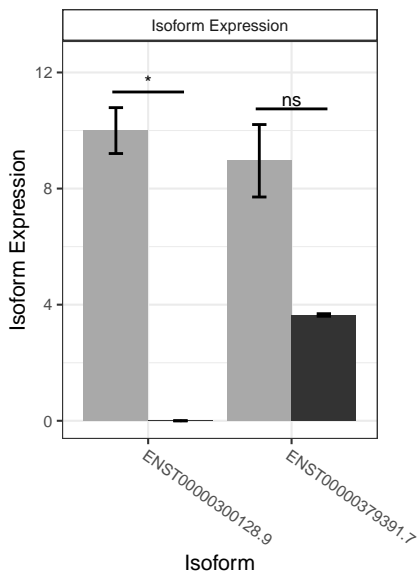
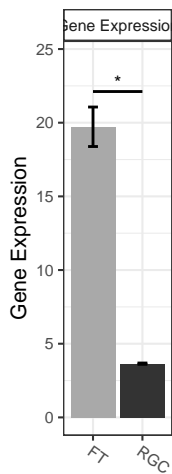
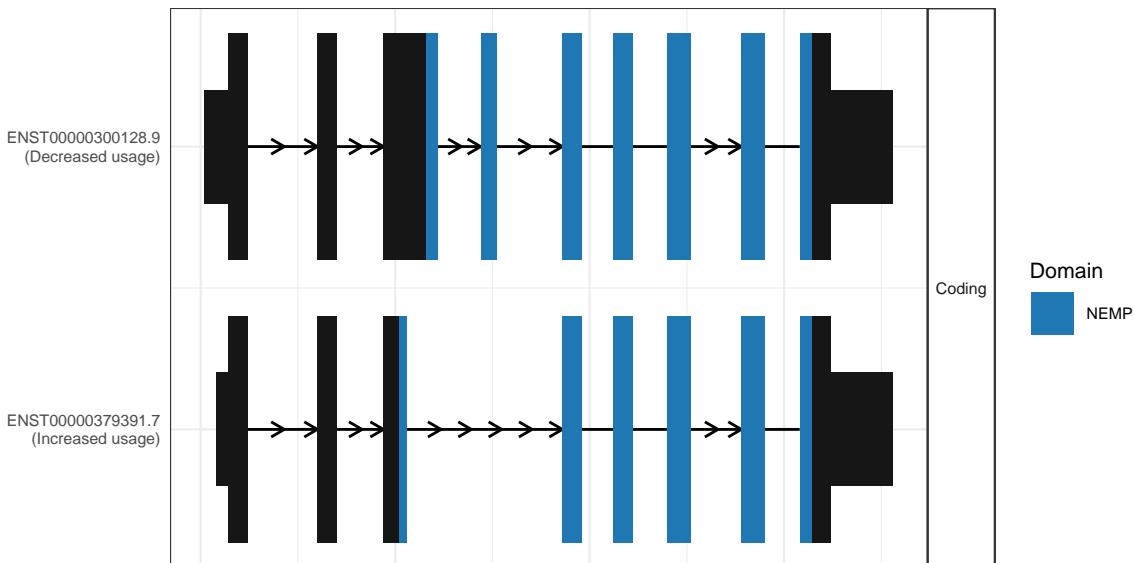


Domain



RUN

The isoform switch in NEMP1 (FT vs RGC)

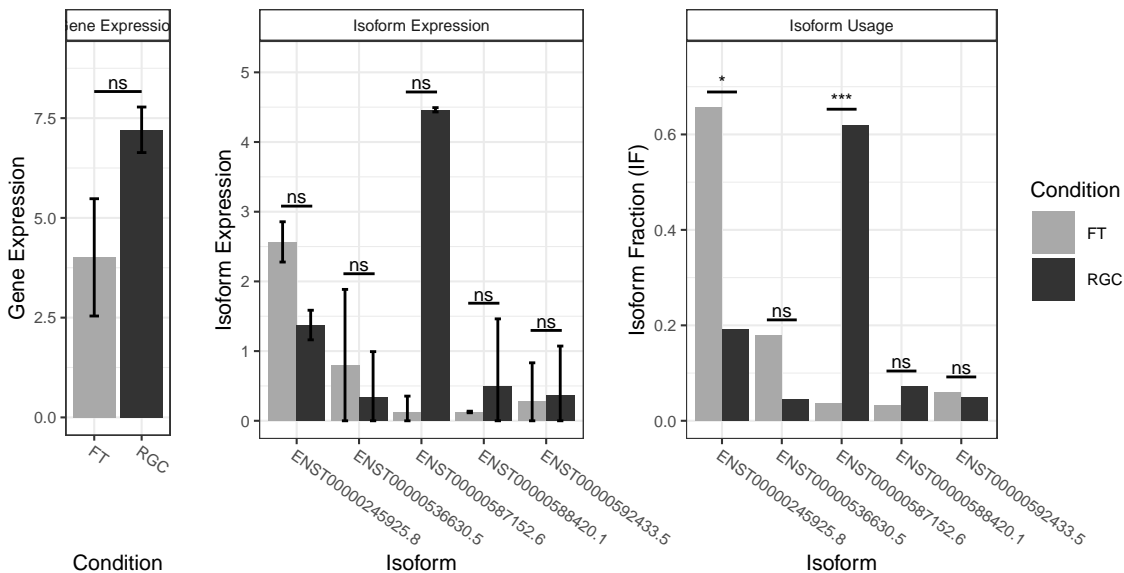
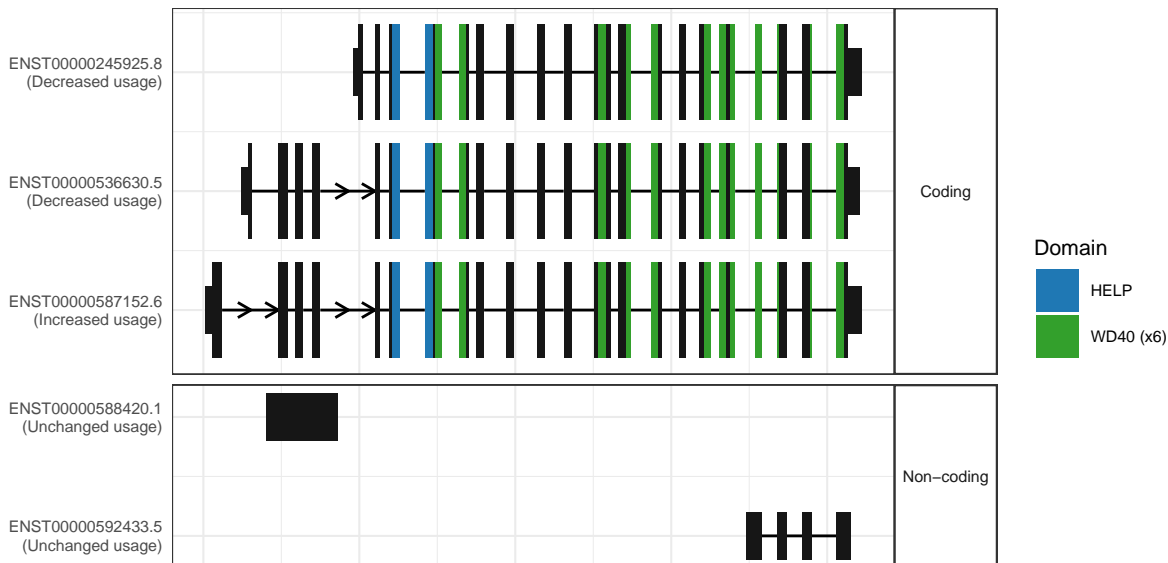


Domain



NEMP

The isoform switch in EML2 (FT vs RGC)



Domain

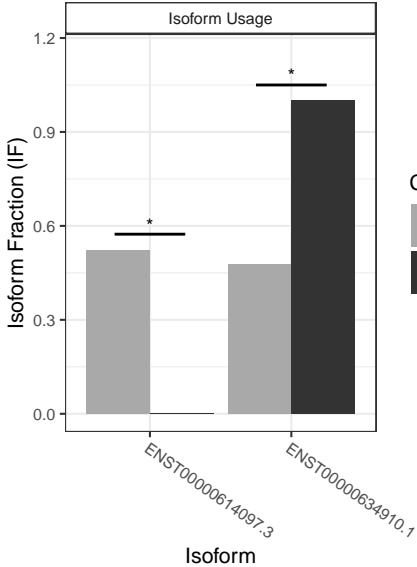
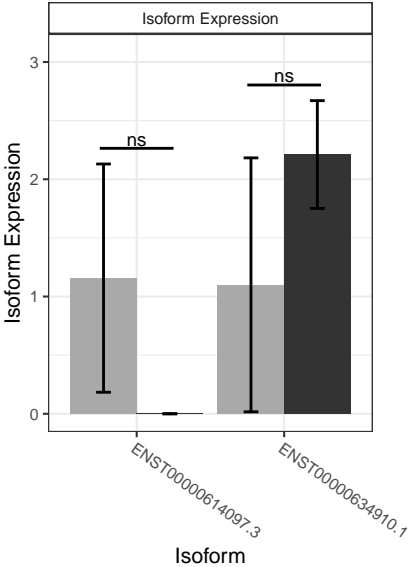
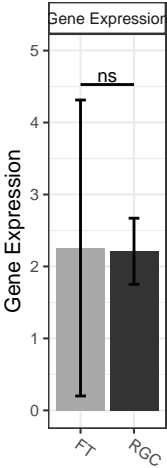
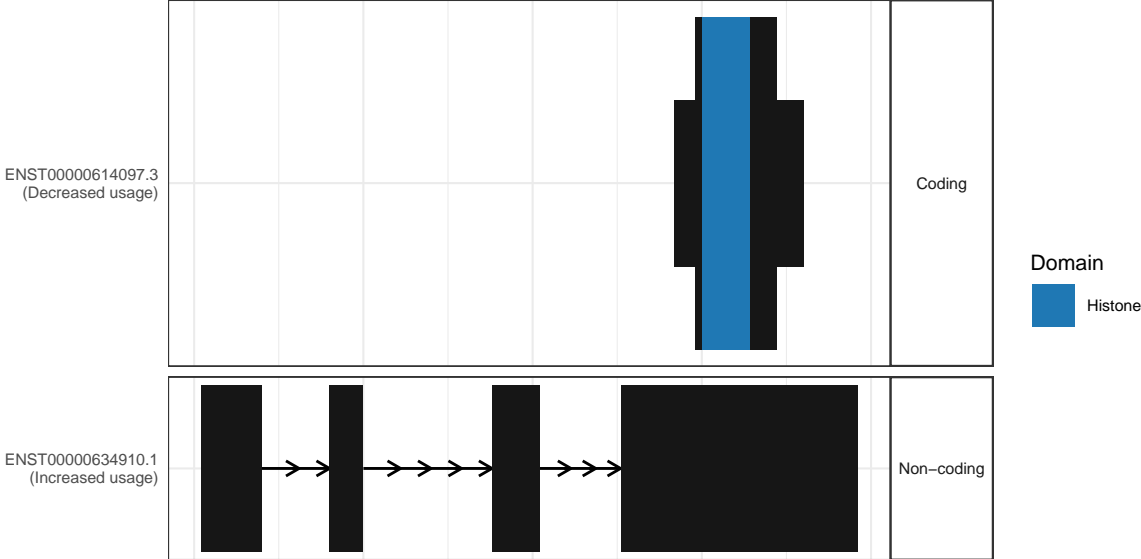


HELP



WD40 (x6)

The isoform switch in H2BC6 (FT vs RGC)

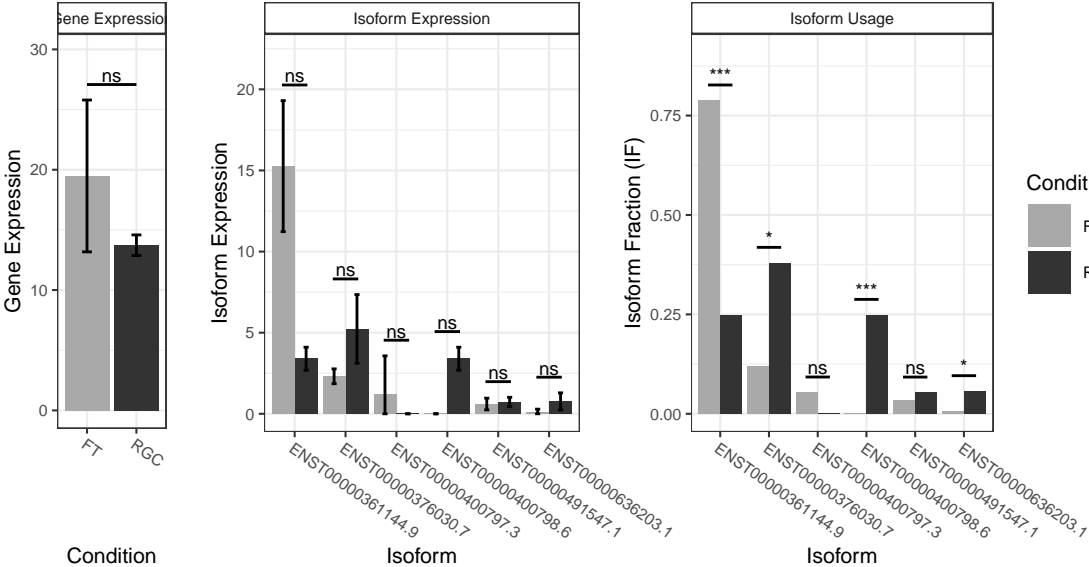
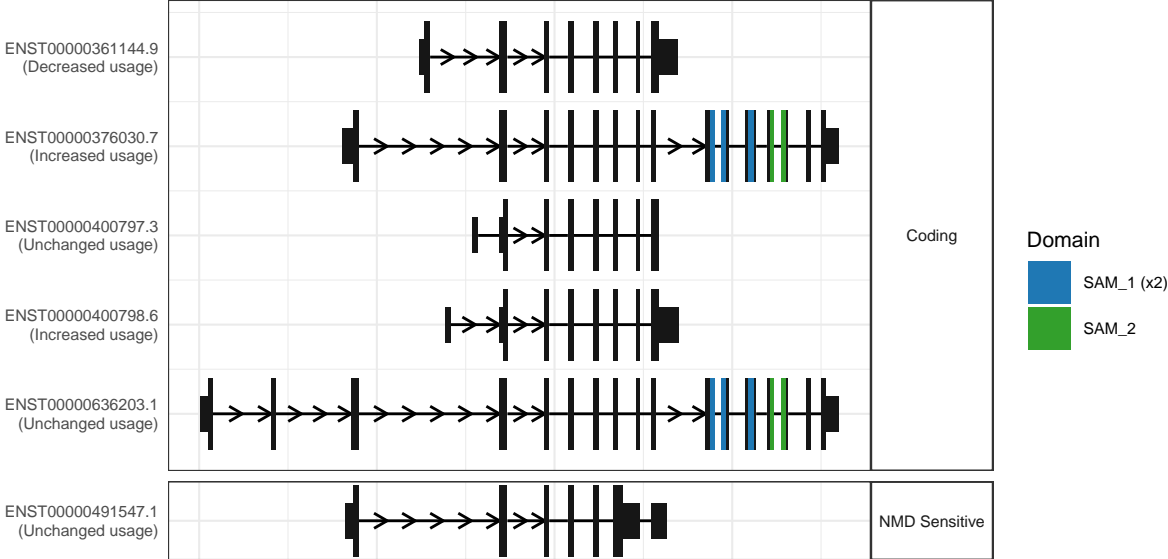


Domain



Histone

The isoform switch in KAZN (FT vs RGC)



Domain

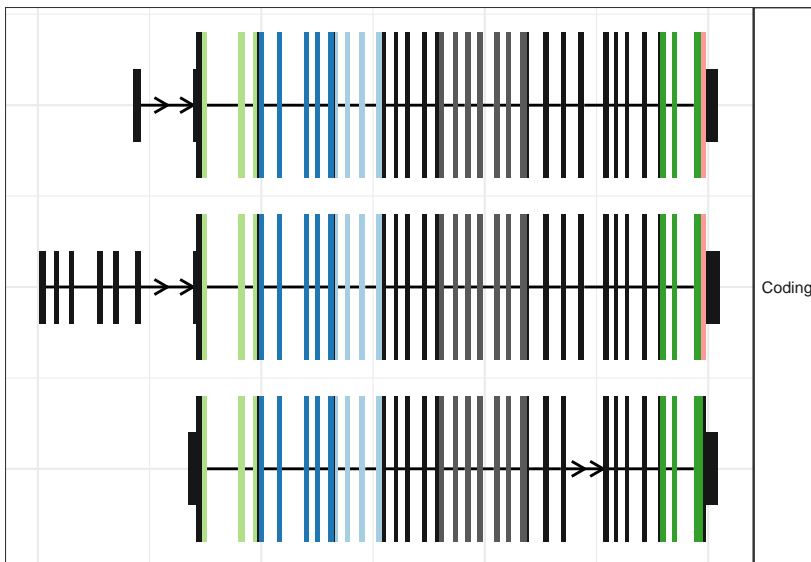


SAM_1 (x2)

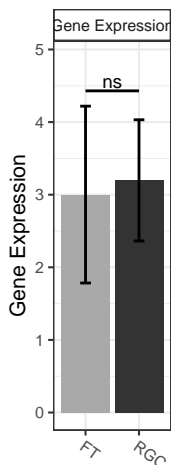
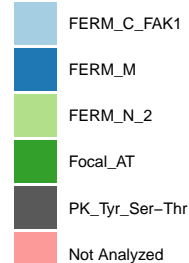


SAM_2

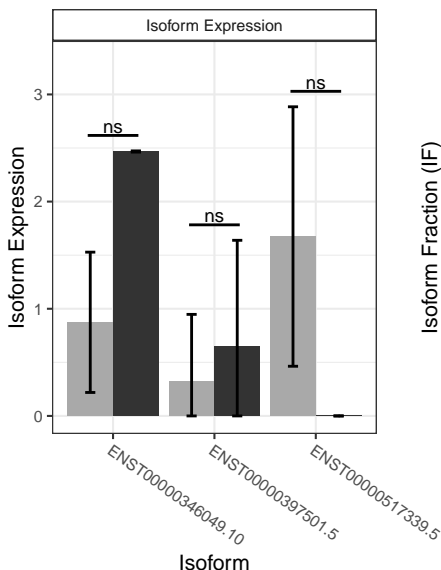
The isoform switch in PTK2B (FT vs RGC)



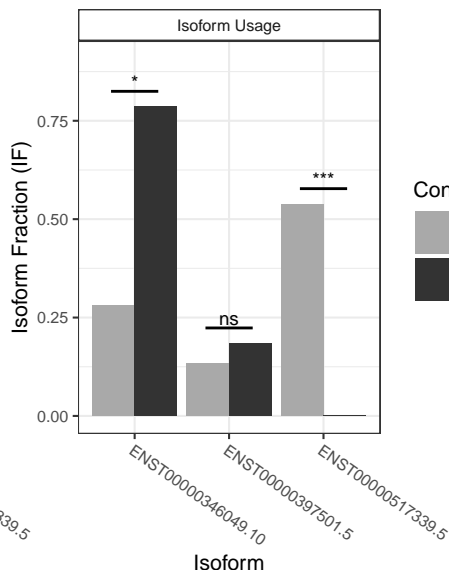
Domain



Condition



Isoform









Isoform

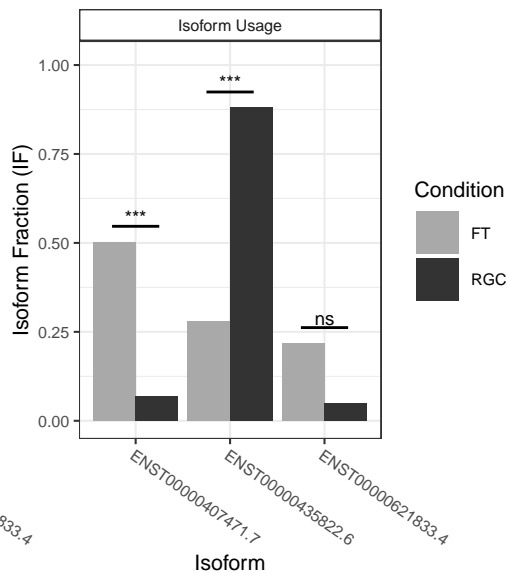
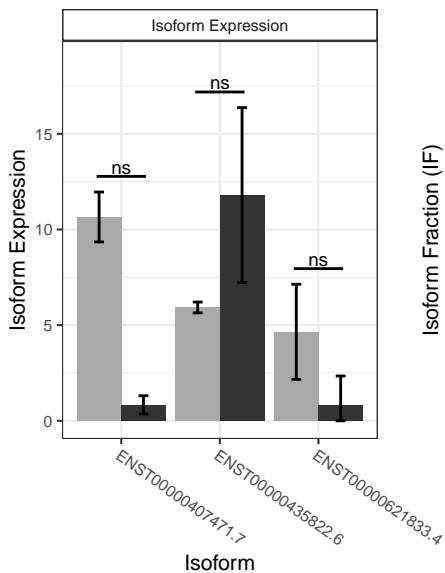
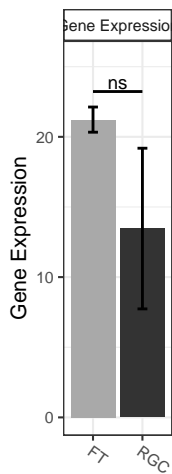
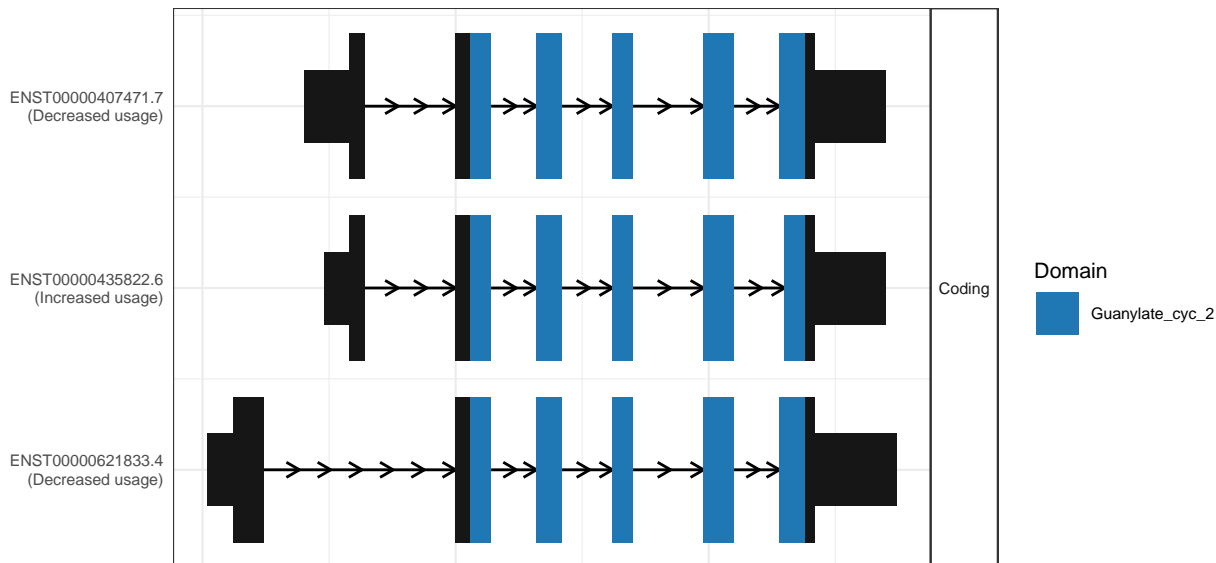
Condition



Domain

	FERM_C_FAK1
	FERM_M
	FERM_N_2
	Focal_AT
	PK_Tyr_Ser-Thr
	Not Analyzed

The isoform switch in GUCD1 (FT vs RGC)

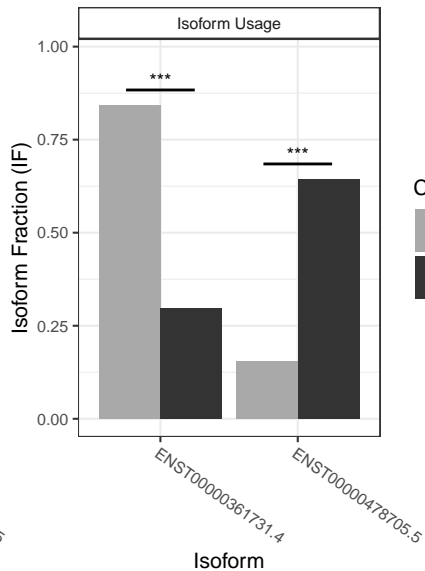
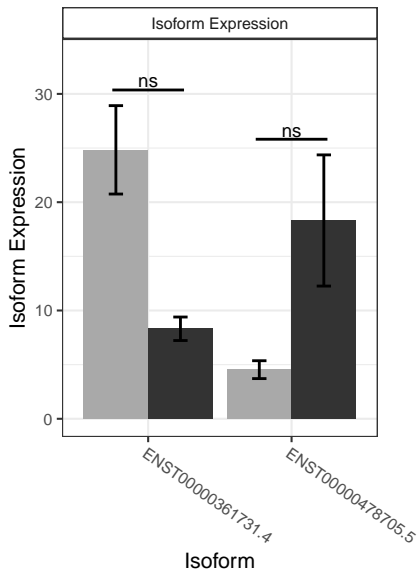
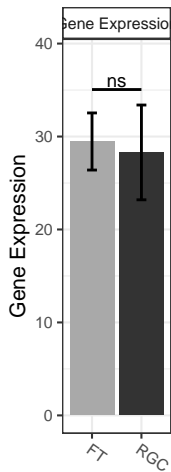
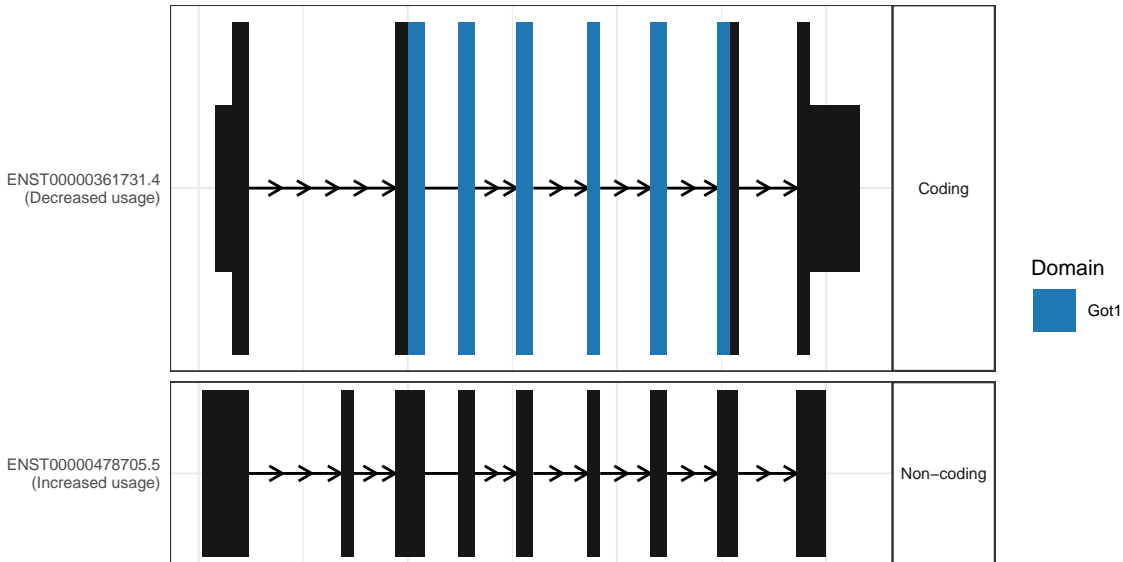


Domain



Guanylate_cyc_2

The isoform switch in SFT2D1 (FT vs RGC)

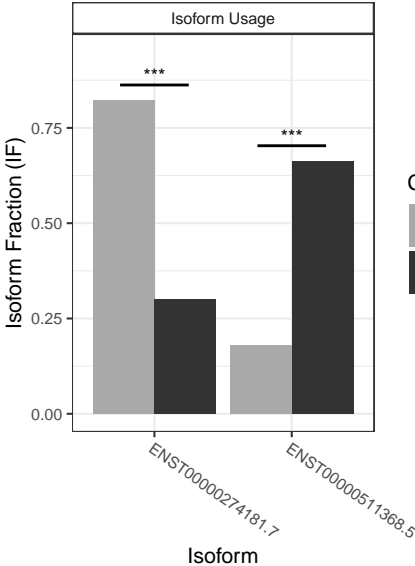
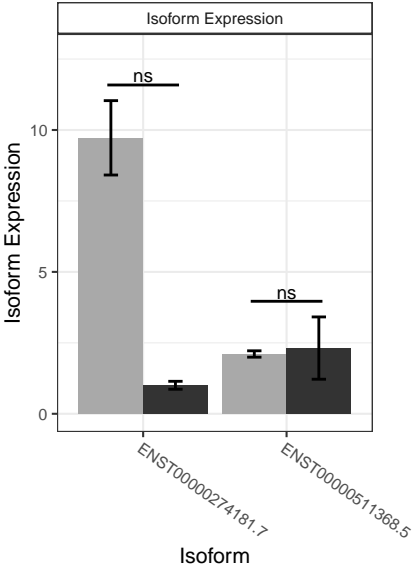
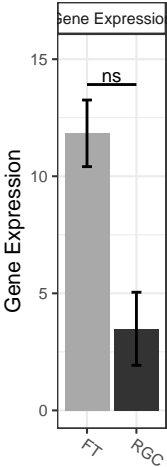
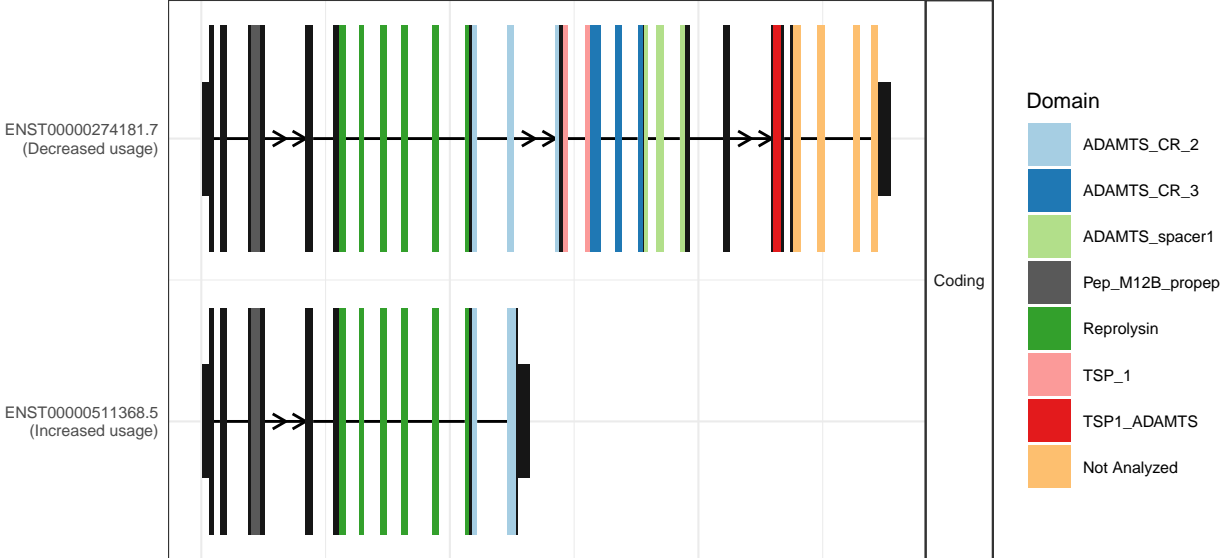


Domain











Got1

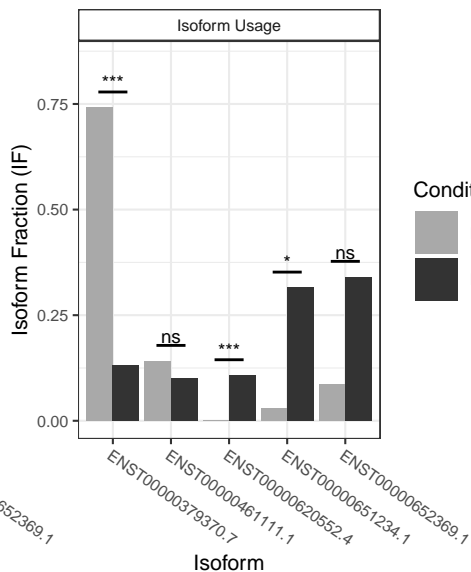
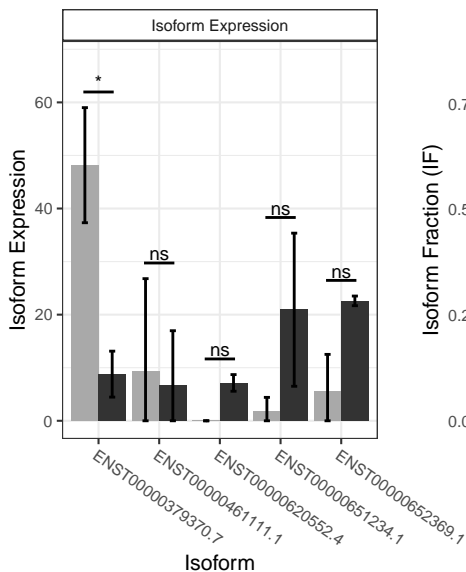
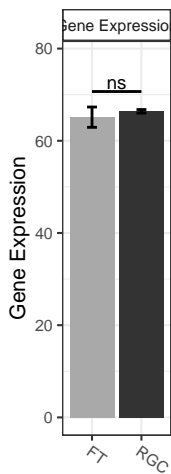
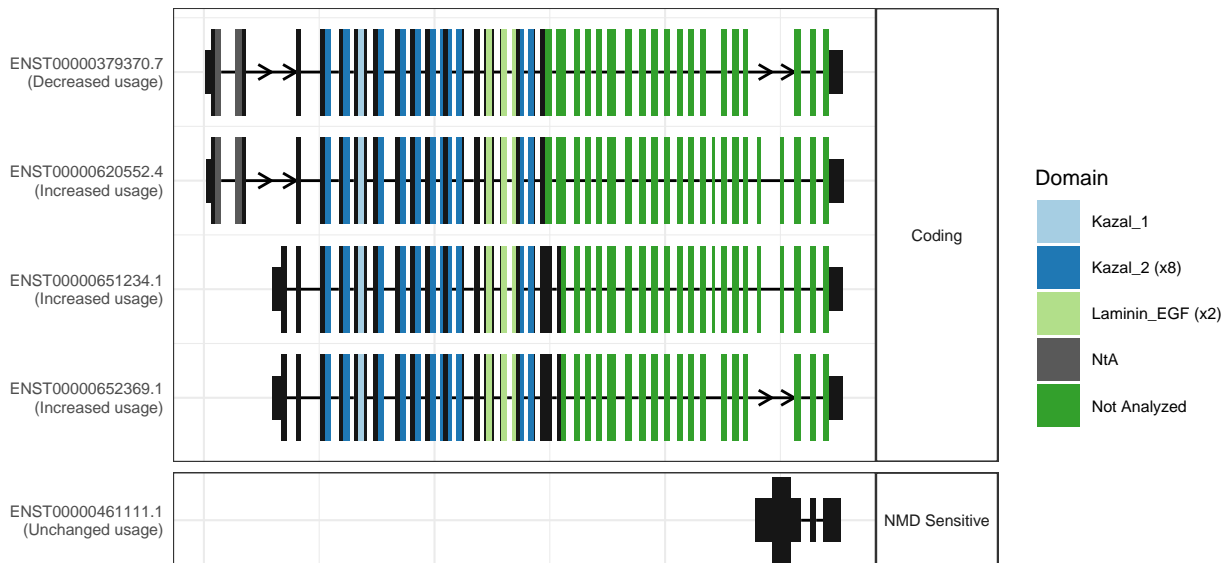
The isoform switch in ADAMTS16 (FT vs RGC)








Domain

	ADAMTS_CR_2
	ADAMTS_CR_3
	ADAMTS_spacer1
	Pep_M12B_propep
	Reprolysin
	TSP_1
	TSP1_ADAMTS
	Not Analyzed

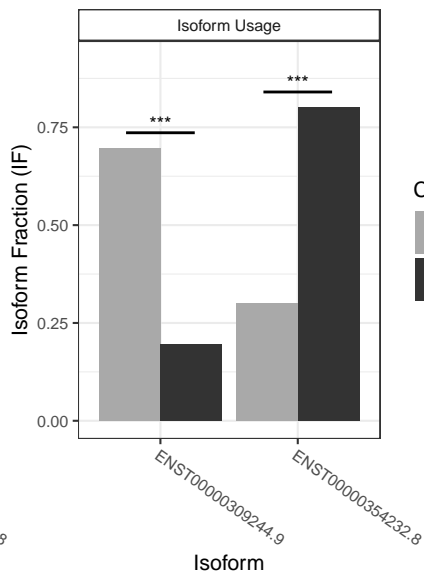
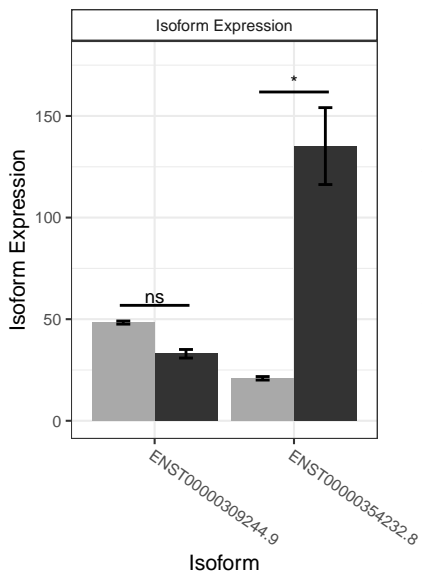
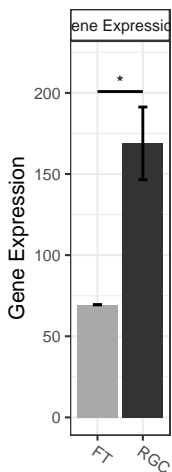
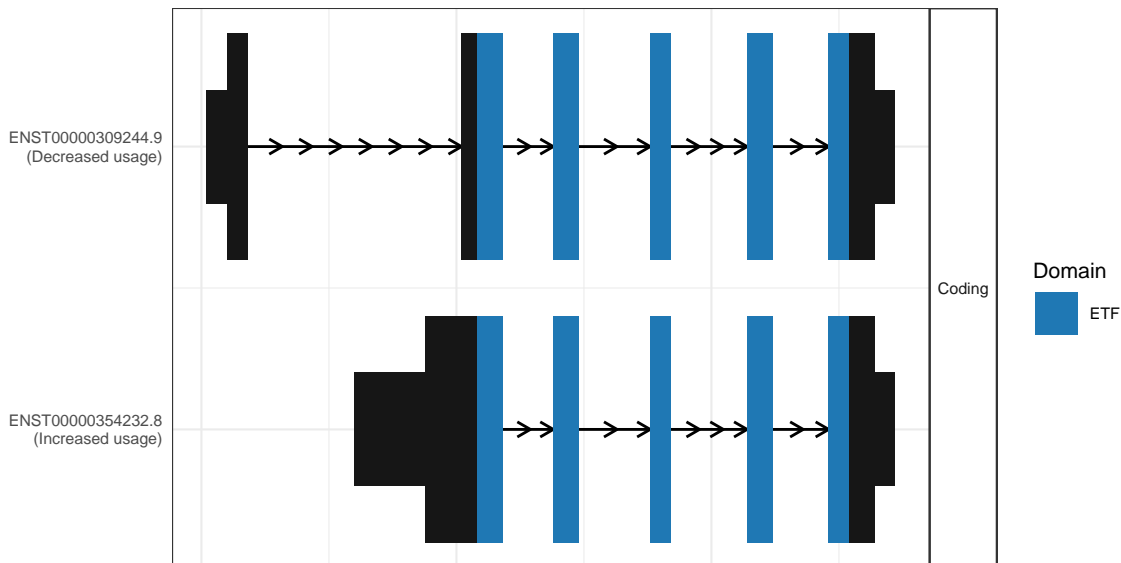
The isoform switch in AGRN (FT vs RGC)



Domain

	Kazal_1
	Kazal_2 (x8)
	Laminin_EGF (x2)
	NtA
	Not Analyzed

The isoform switch in ETFB (FT vs RGC)

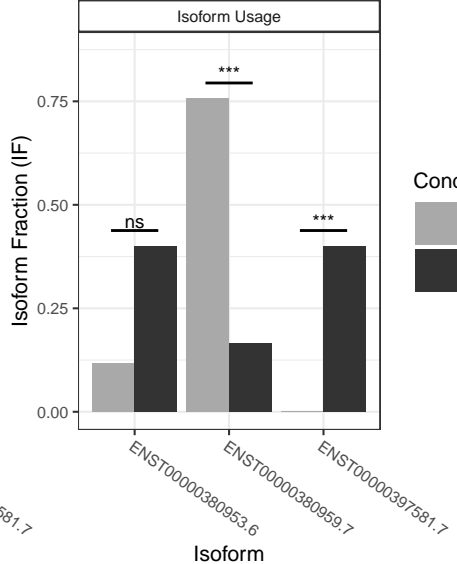
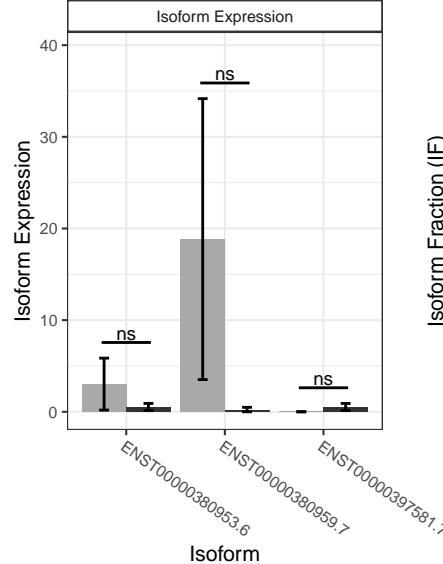
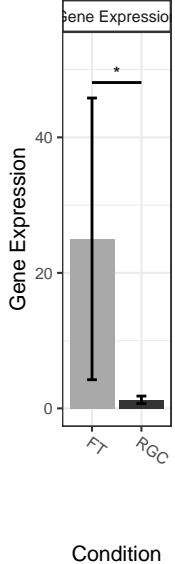
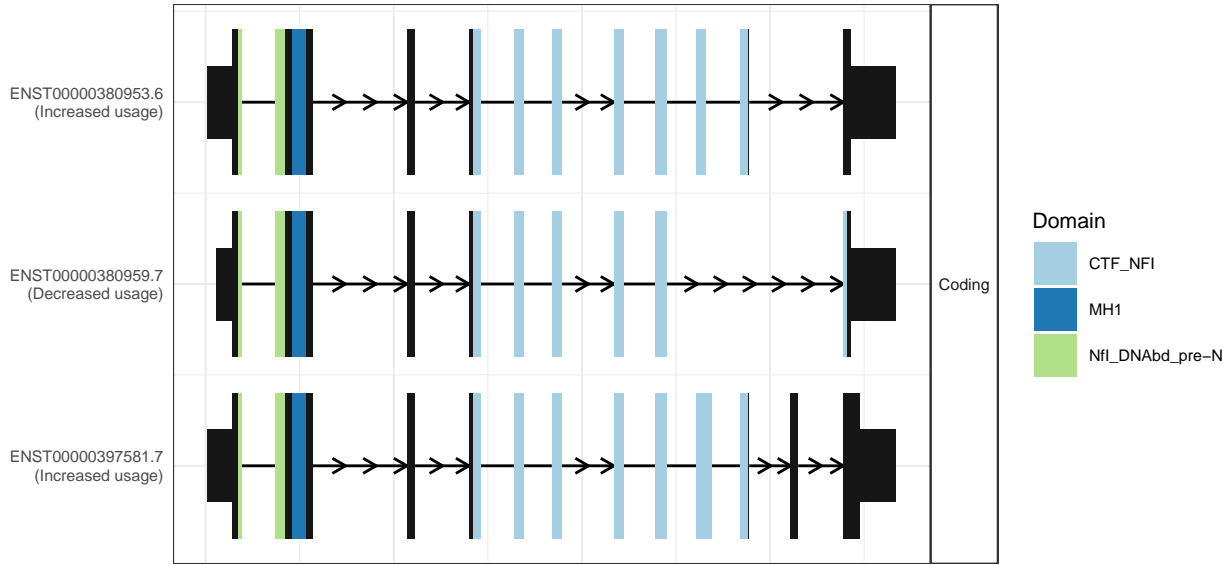


Domain



ETF

The isoform switch in NFIB (FT vs RGC)



Domain



CTF_NFI

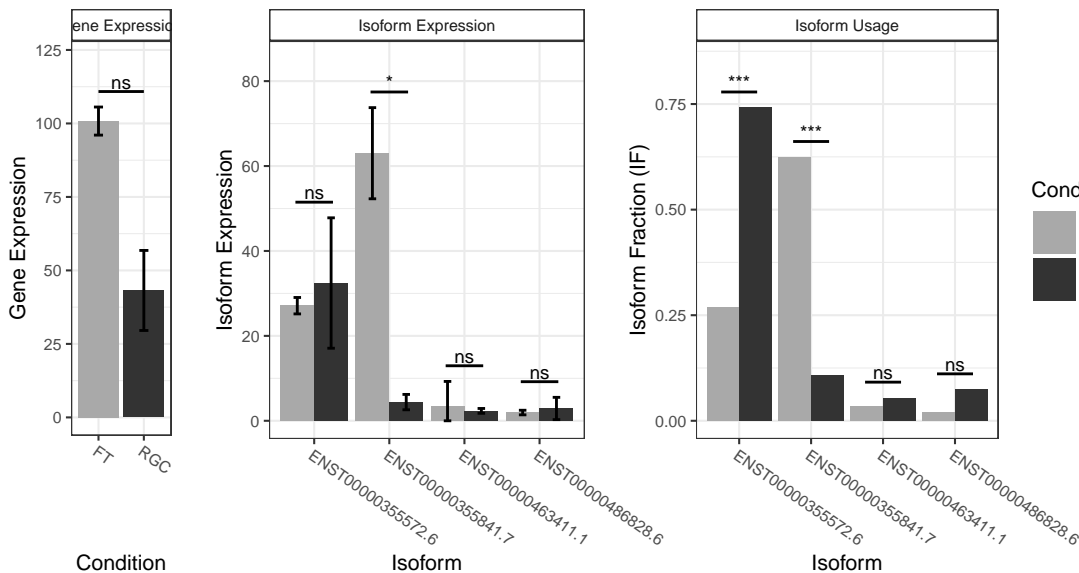
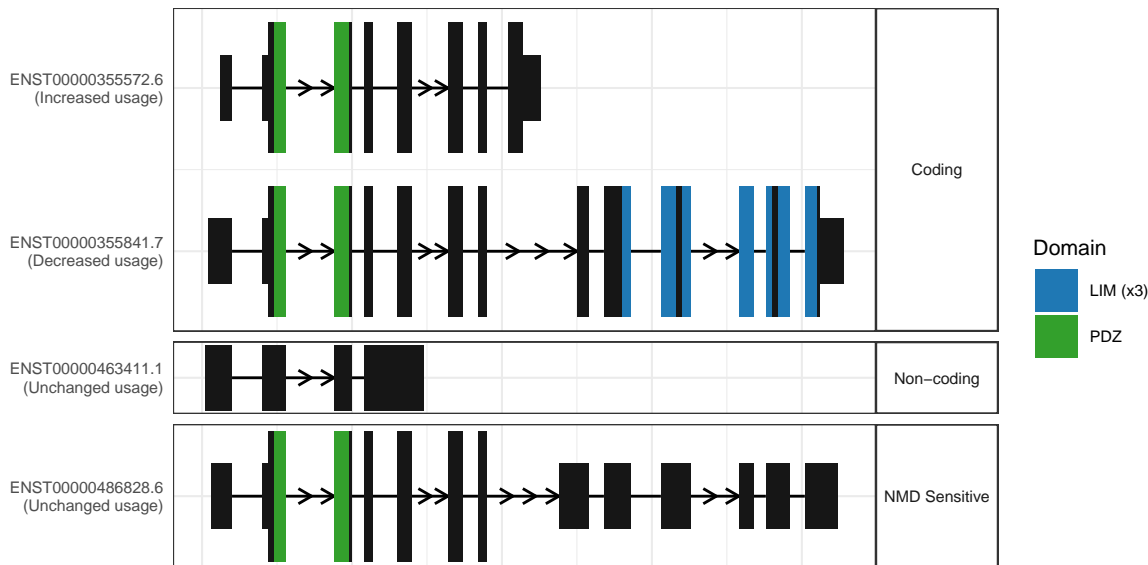


MH1



Nfil_DNAAbd_pre-N

The isoform switch in PDLIM7 (FT vs RGC)



Domain

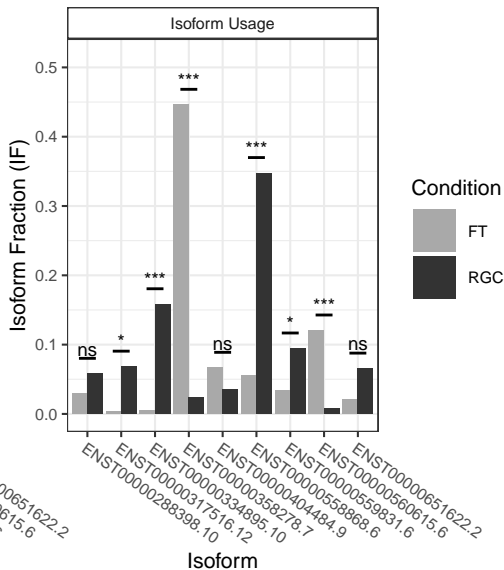
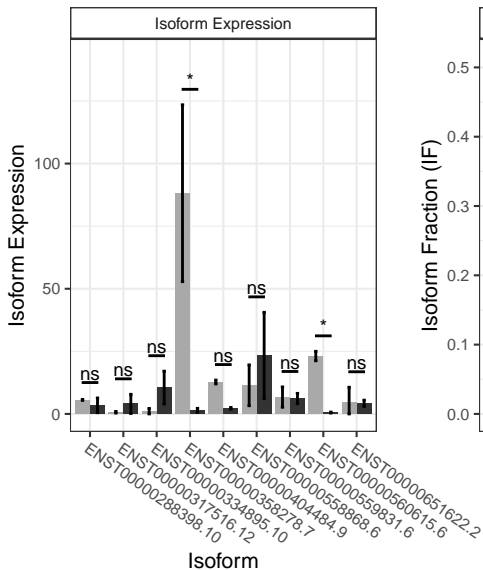
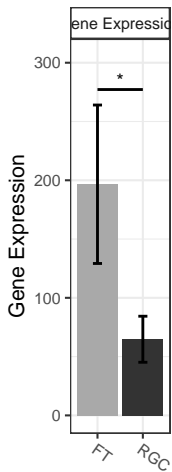
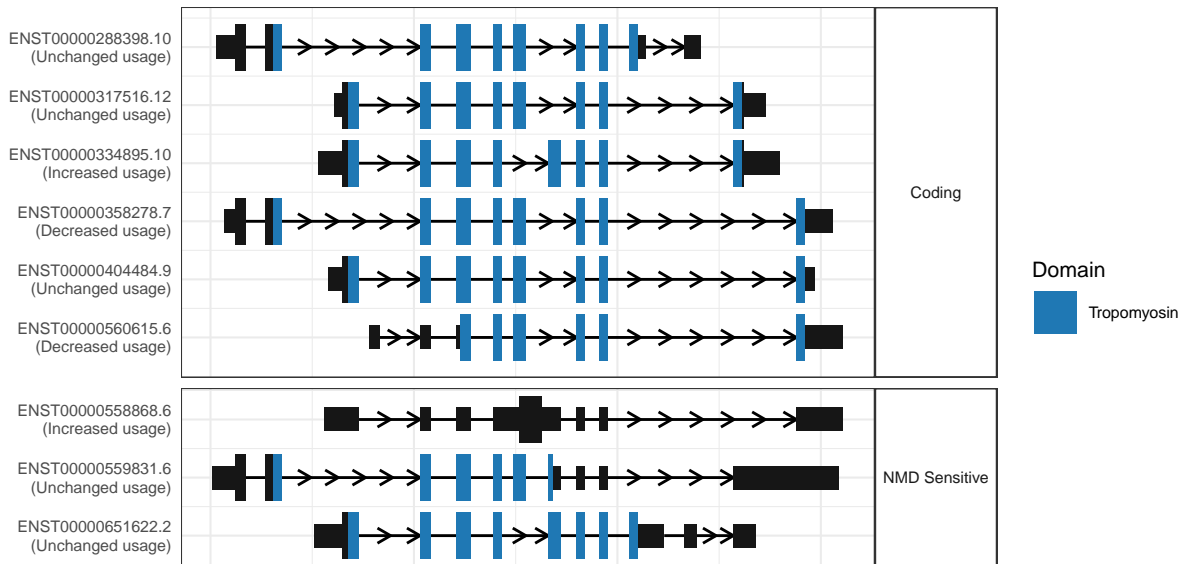


LIM (x3)



PDZ

The isoform switch in TPM1 (FT vs RGC)



Condition

Isoform

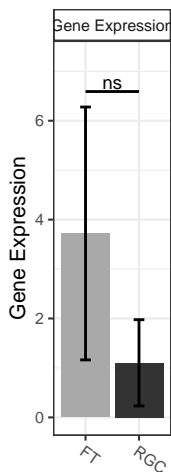
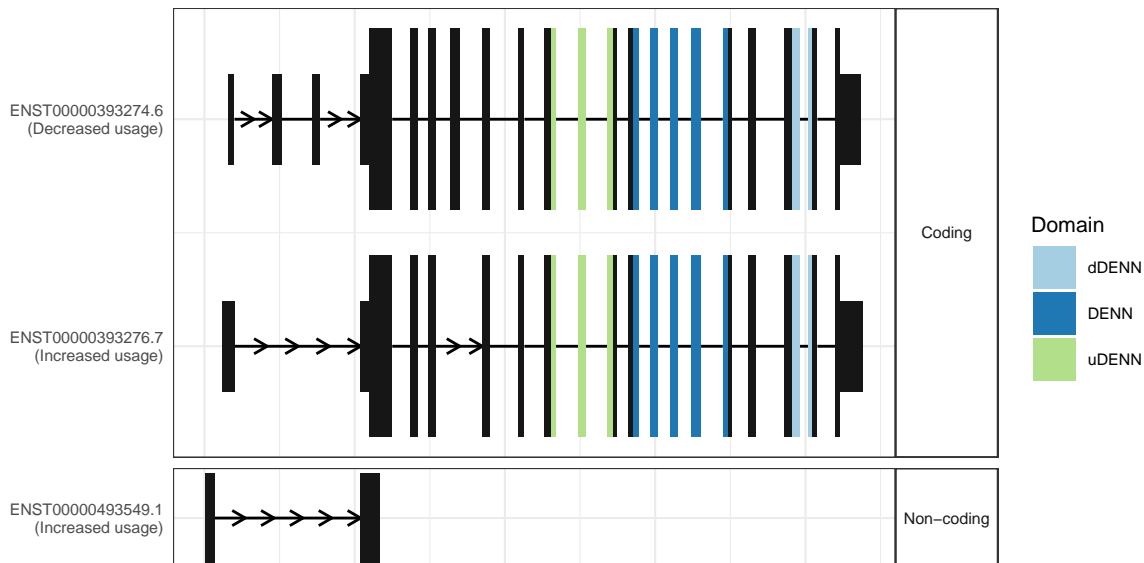
Isoform

Domain

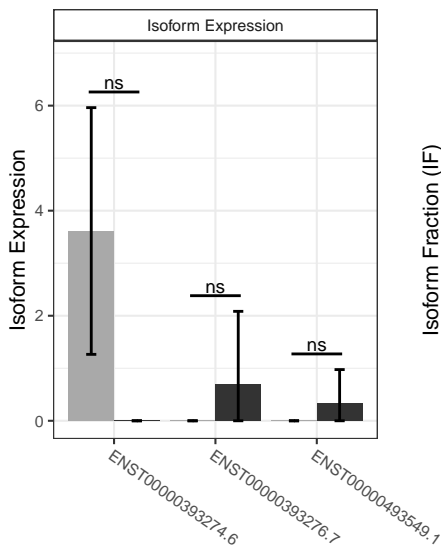


Tropomyosin

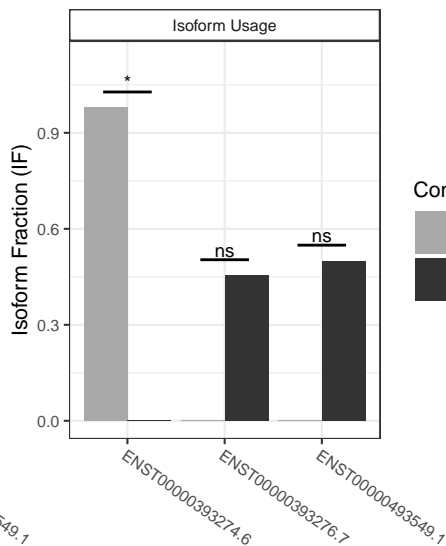
The isoform switch in DENND2C (FT vs RGC)



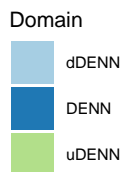
Condition



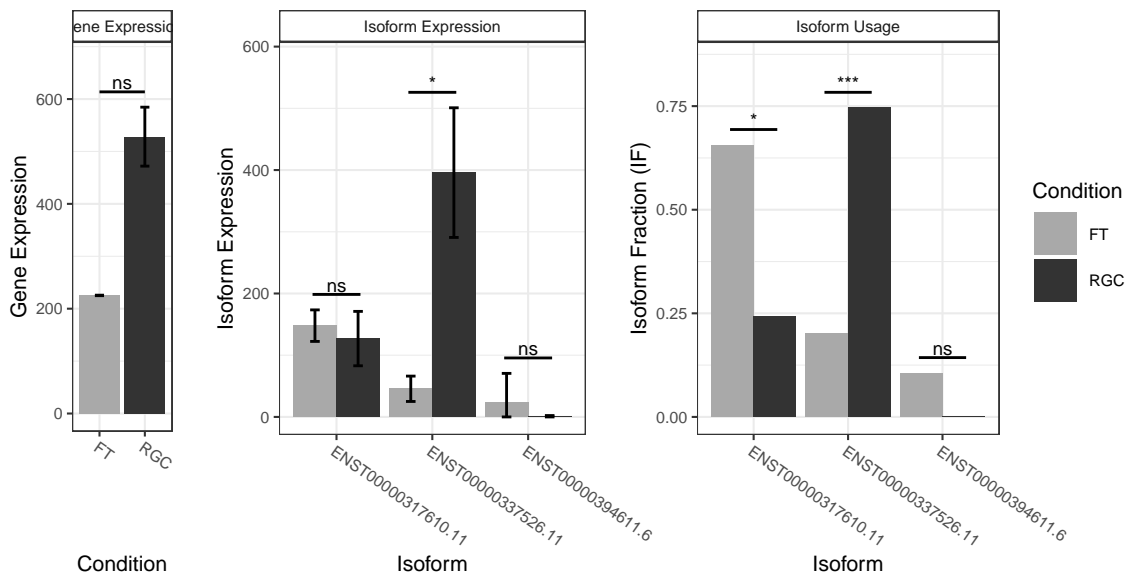
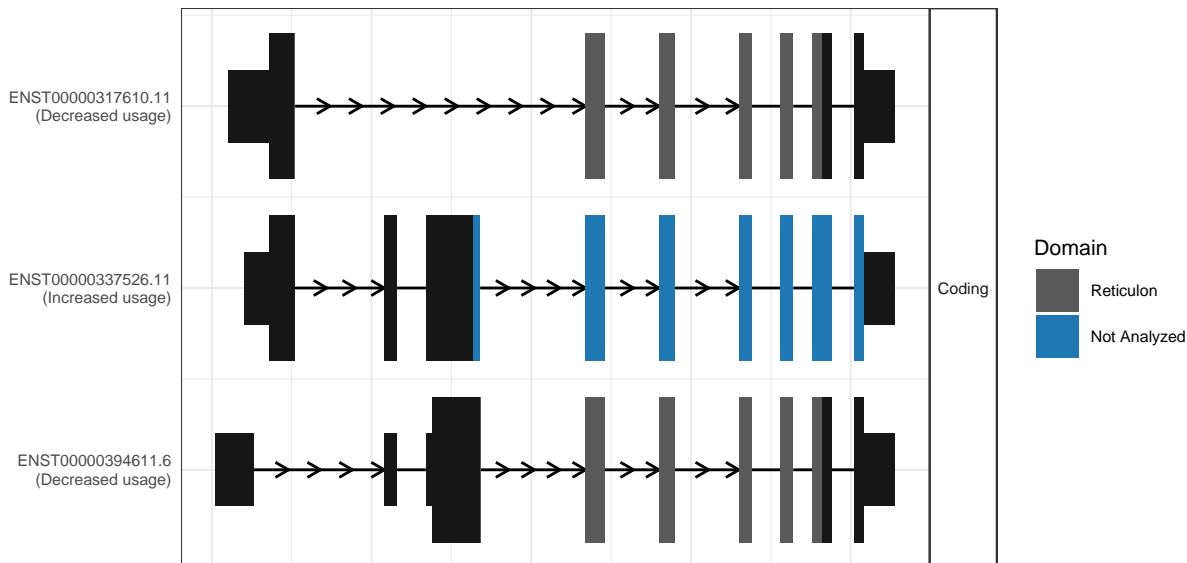
Isoform



Isoform



The isoform switch in RTN4 (FT vs RGC)



Domain

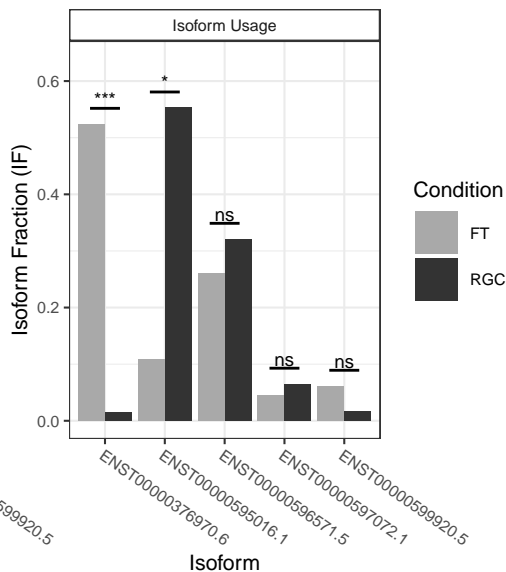
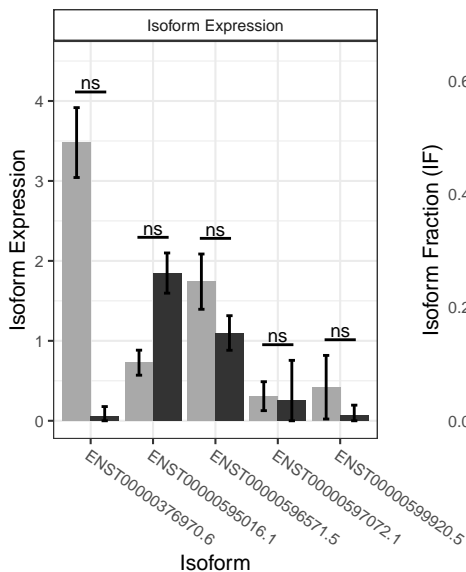
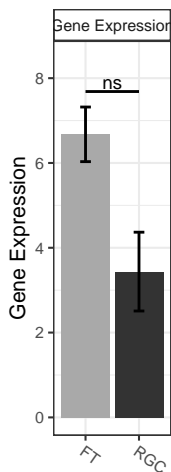
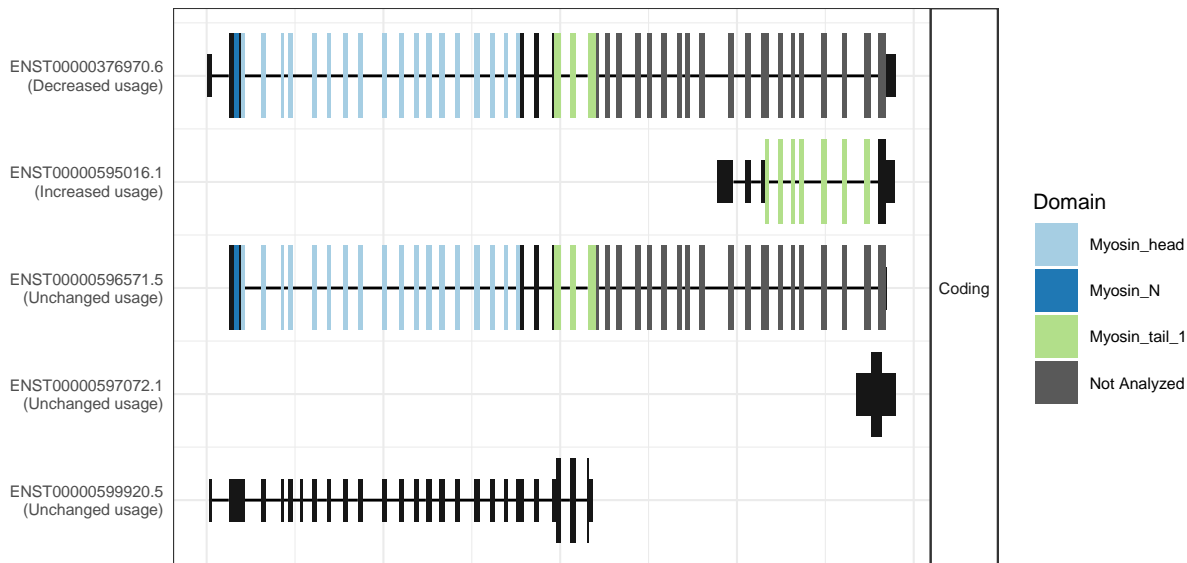


Reticulon







Not Analyzed

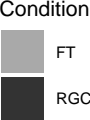
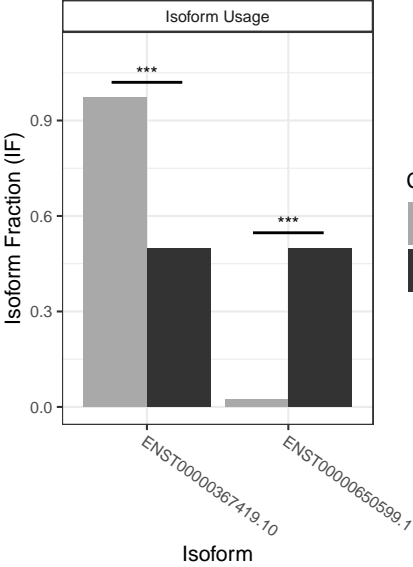
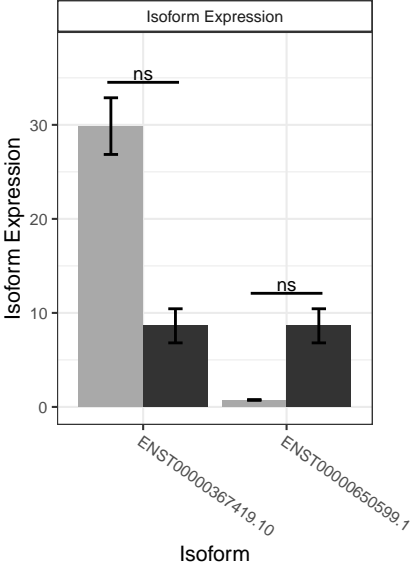
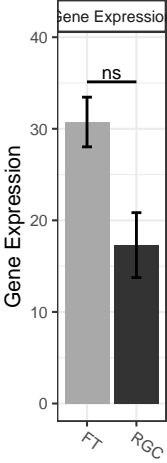
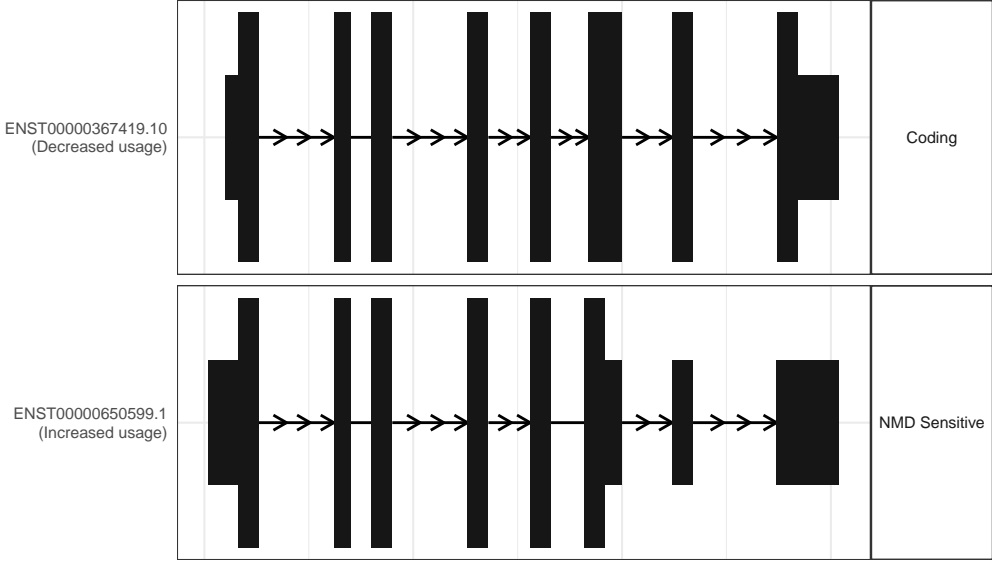
The isoform switch in MYH14 (FT vs RGC)



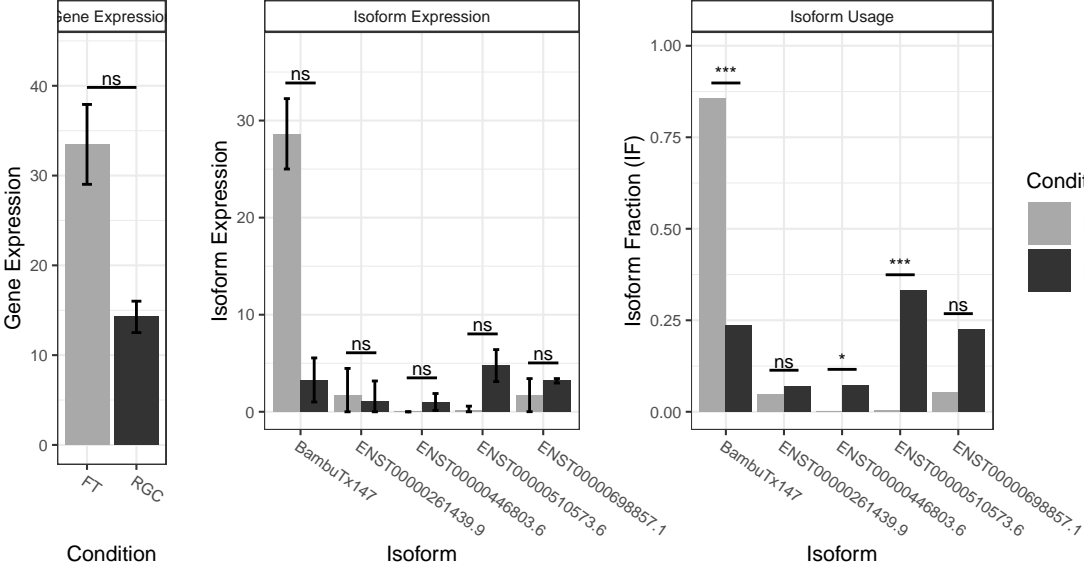
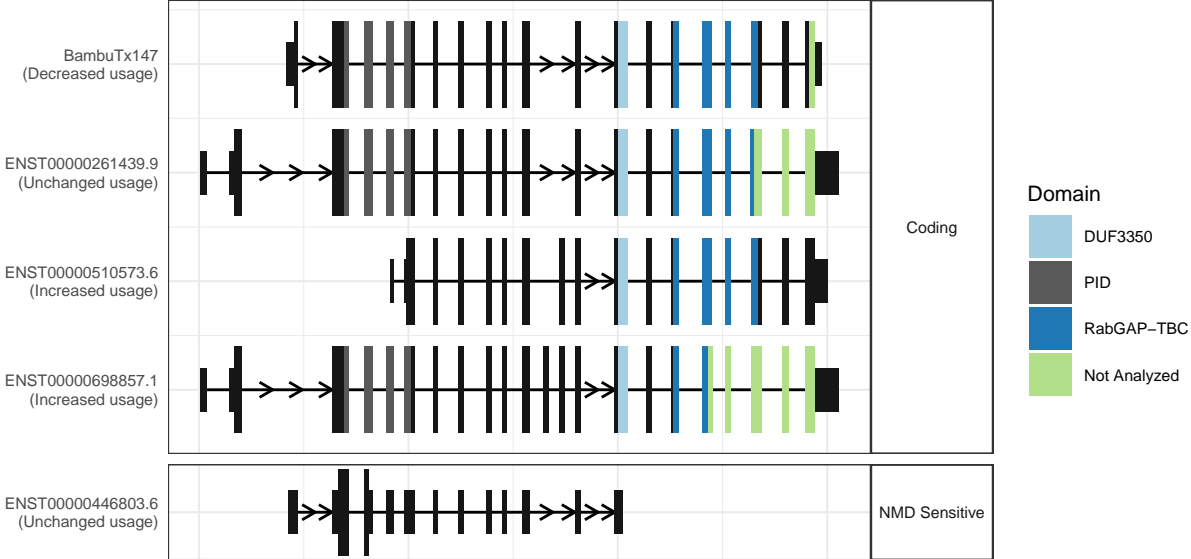
Domain

	Myosin_head
	Myosin_N
	Myosin_tail_1
	Not Analyzed





The isoform switch in GINM1 (FT vs RGC)



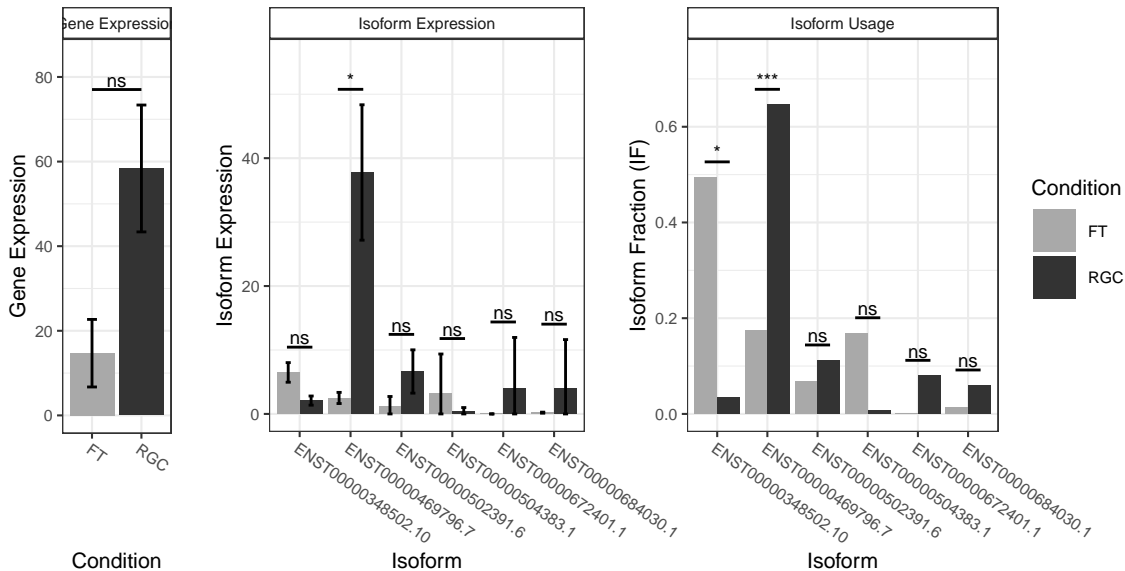
The isoform switch in TBC1D1 (FT vs RGC)



Domain

	DUF3350
	PID
	RabGAP-TBC
	Not Analyzed

	Coding
	Non-coding
	NMD Sensitive

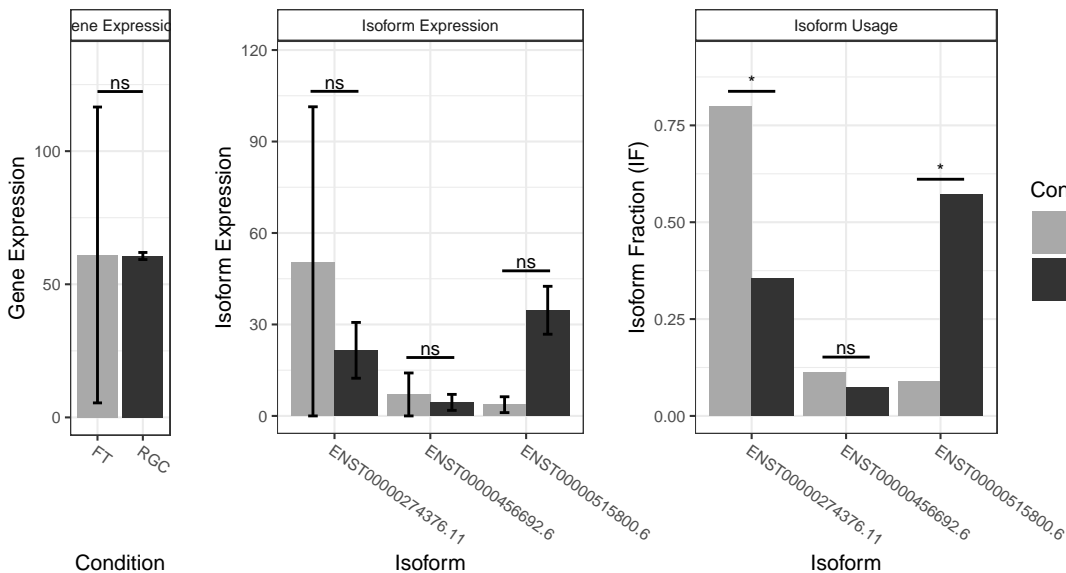
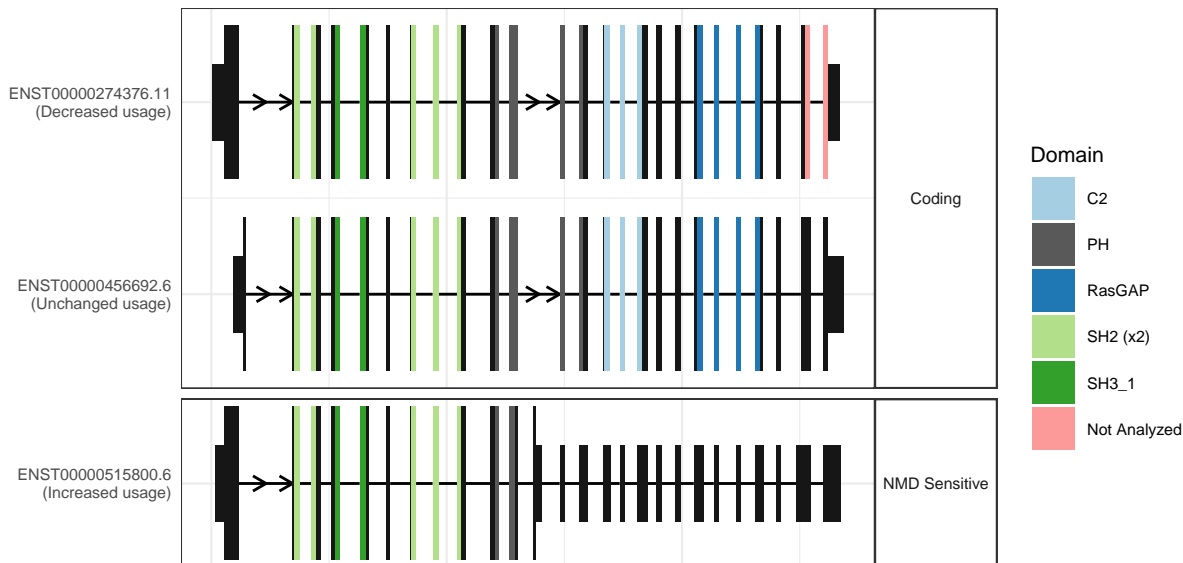


Domain









AMP-binding

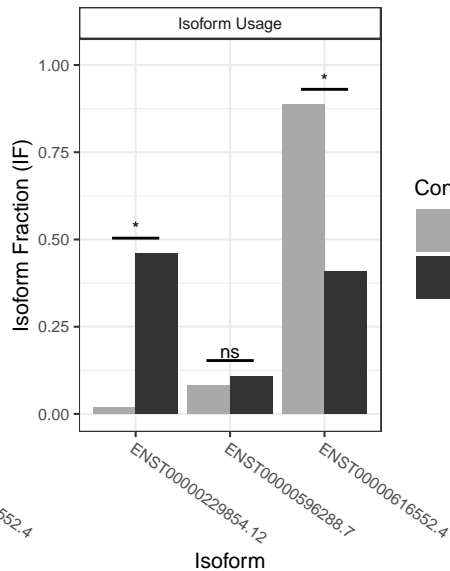
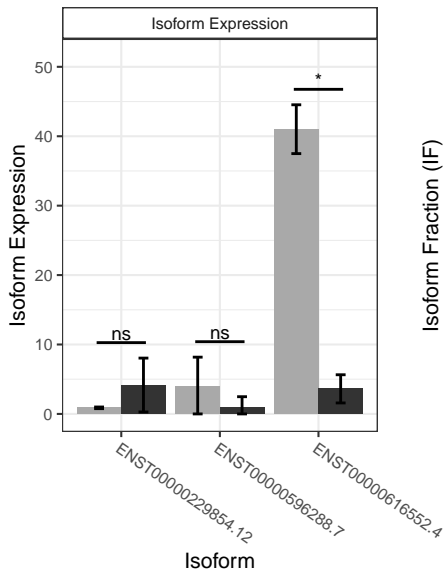
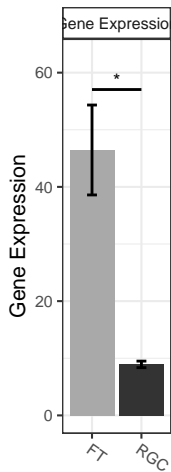
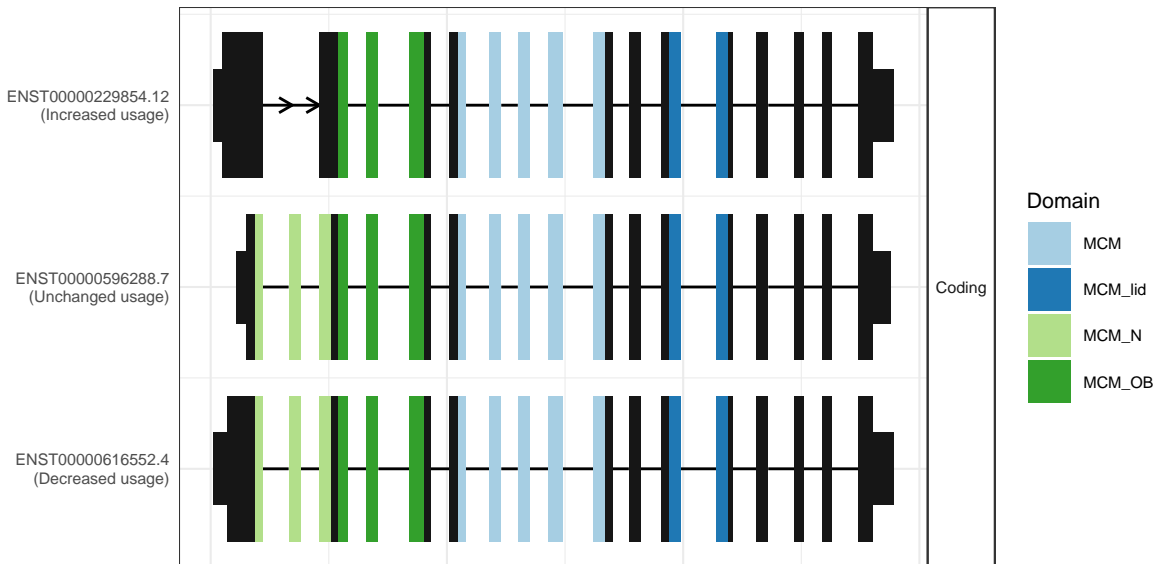
The isoform switch in RASA1 (FT vs RGC)



Domain

	C2
	PH
	RasGAP
	SH2 (x2)
	SH3_1
	Not Analyzed

The isoform switch in MCM3 (FT vs RGC)



Domain



MCM



MCM_lid

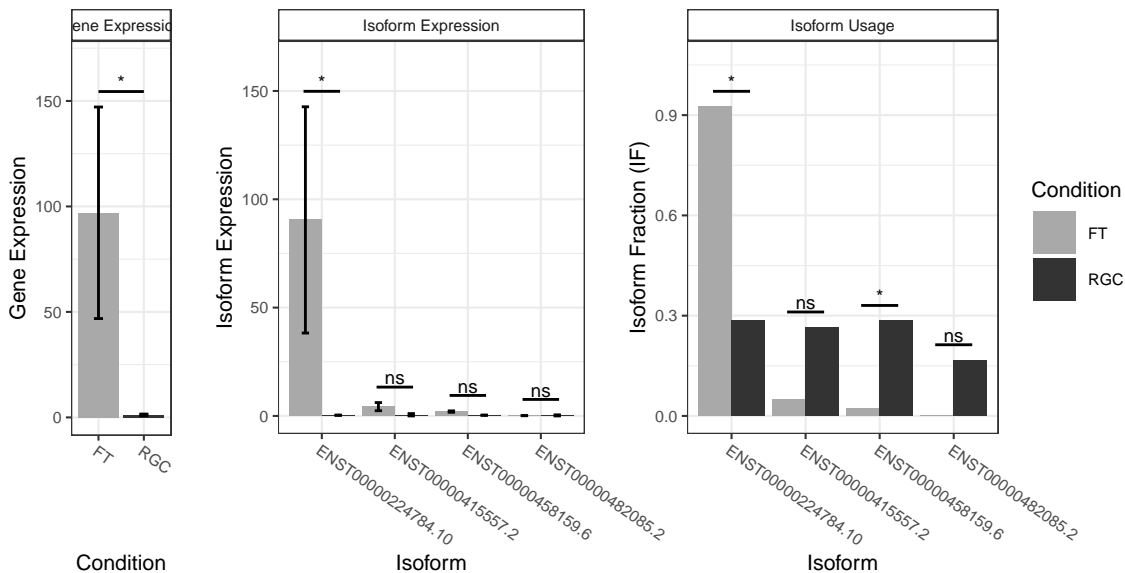


MCM_N



MCM_OB

	Coding
	Non-coding

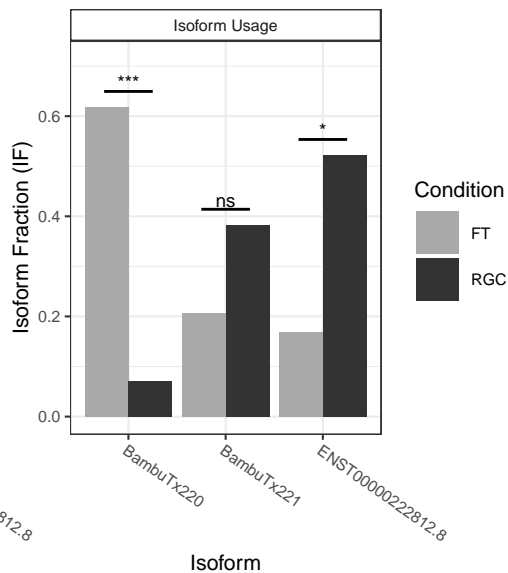
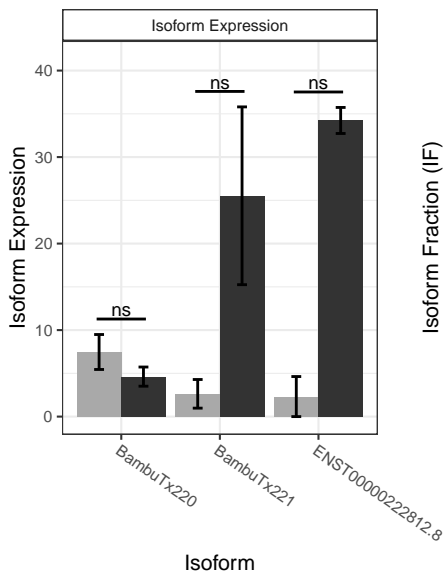
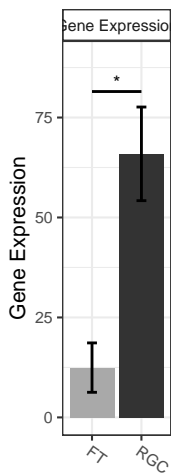
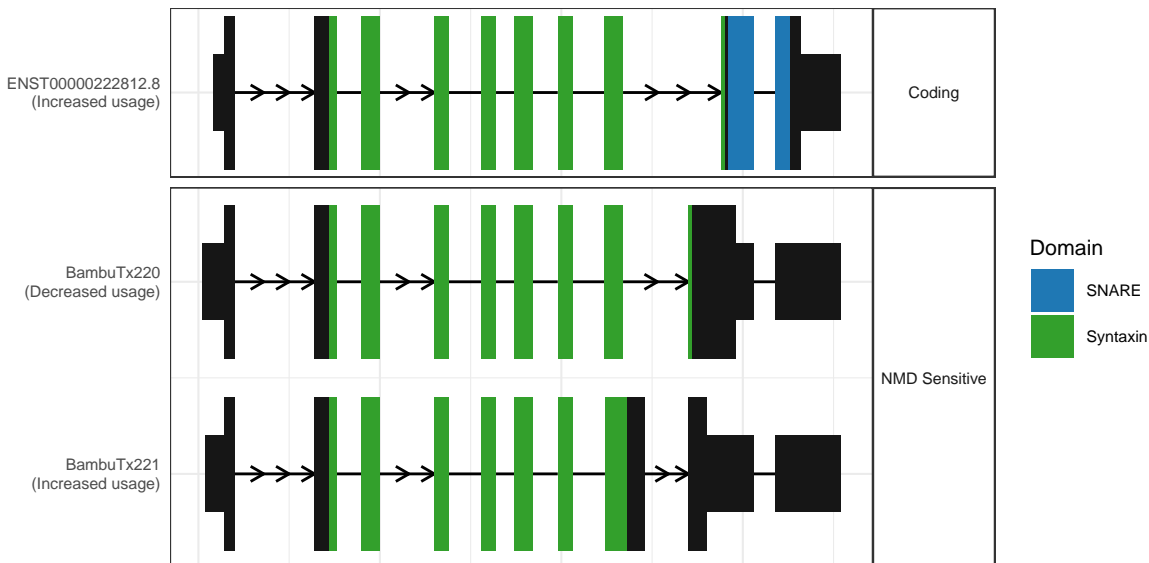


Domain



Actin

The isoform switch in STX1A (FT vs RGC)



Domain

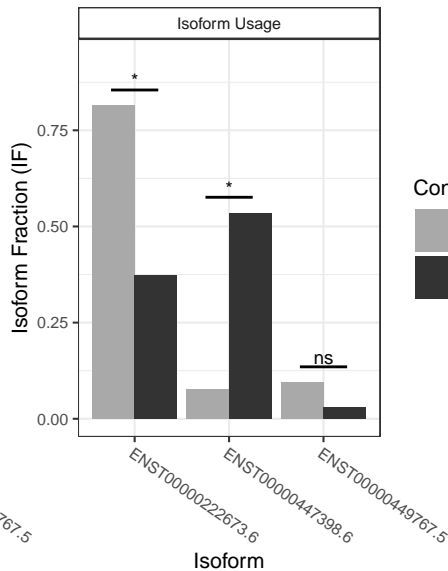
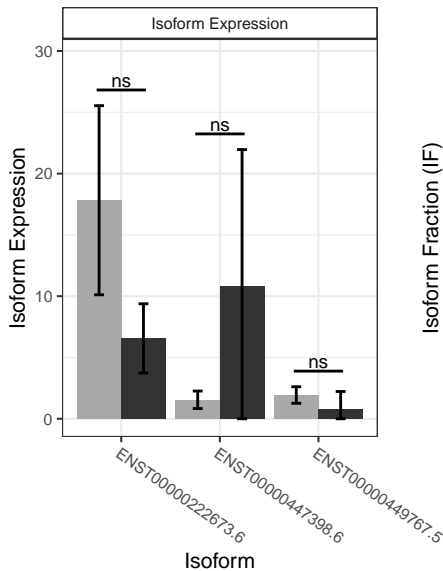
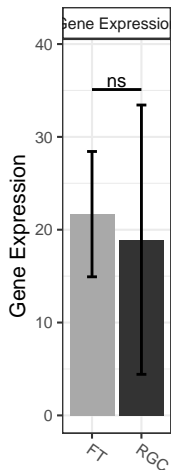
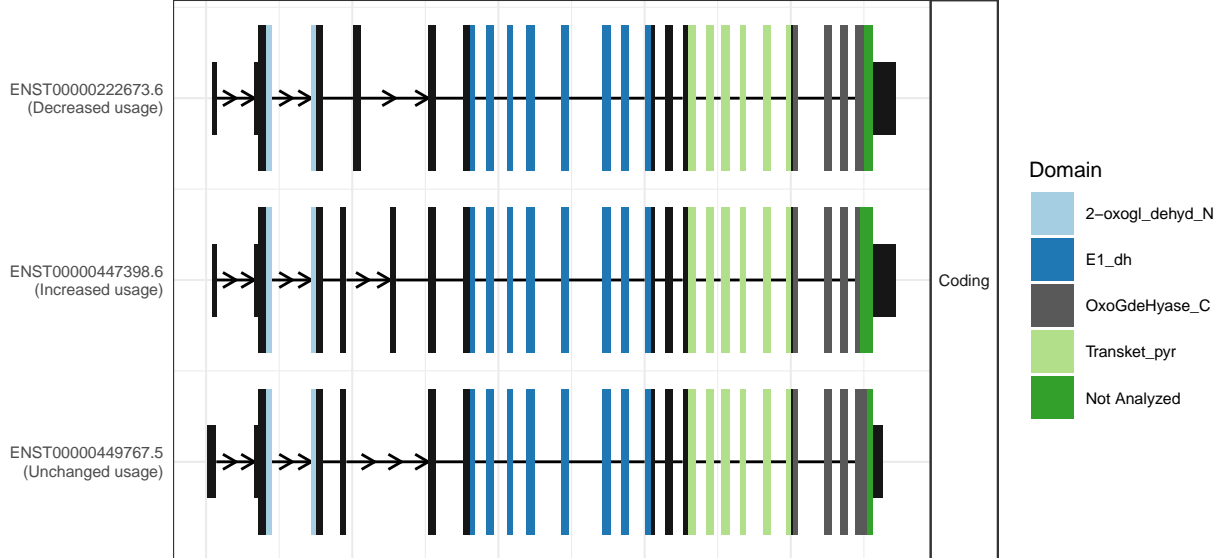


SNARE



Syntaxin

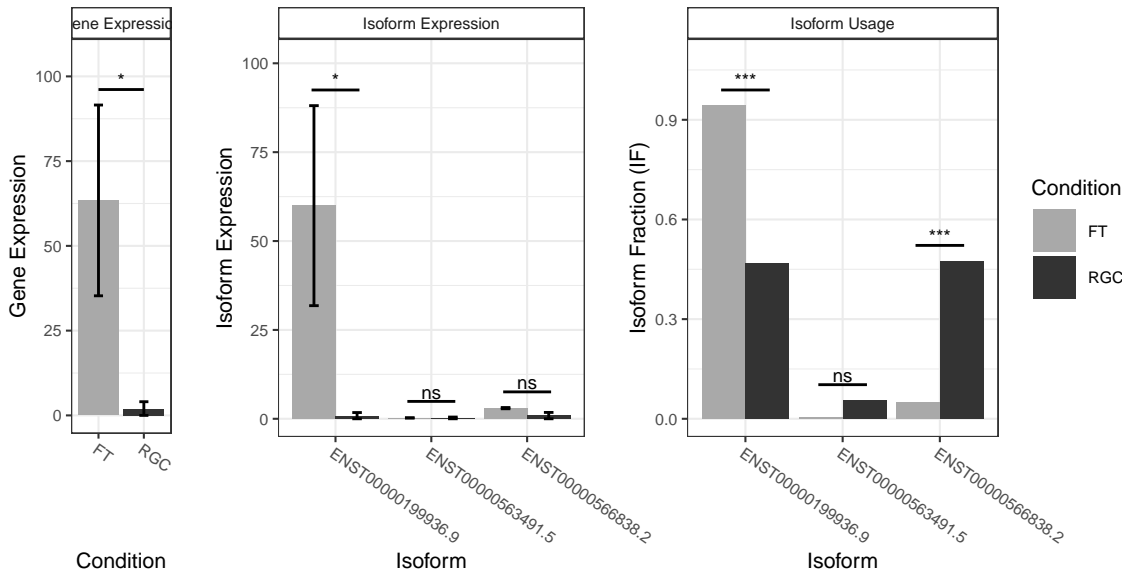
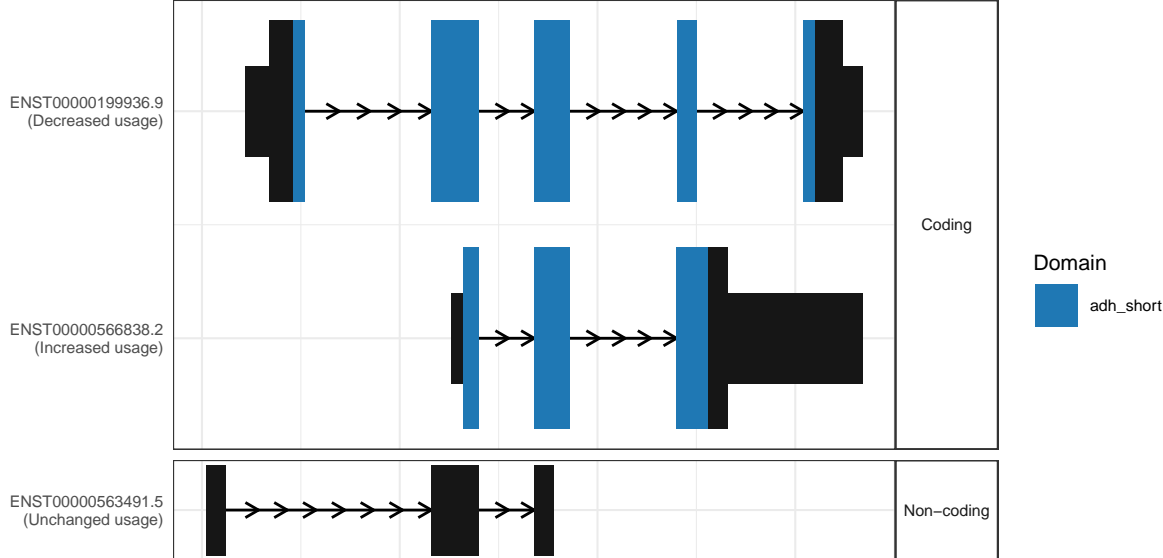
The isoform switch in OGDH (FT vs RGC)



Domain



The isoform switch in HSD17B2 (FT vs RGC)

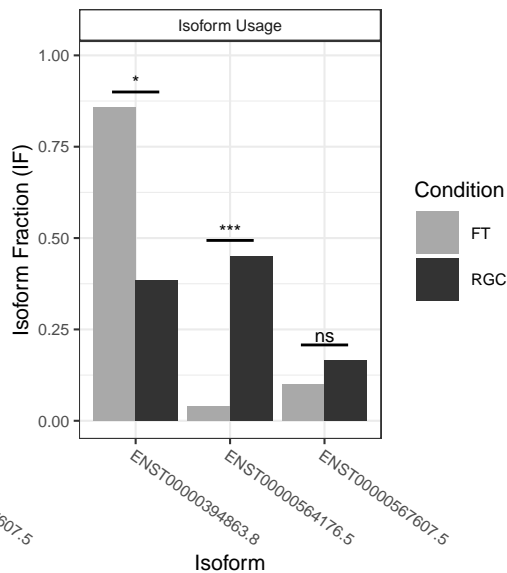
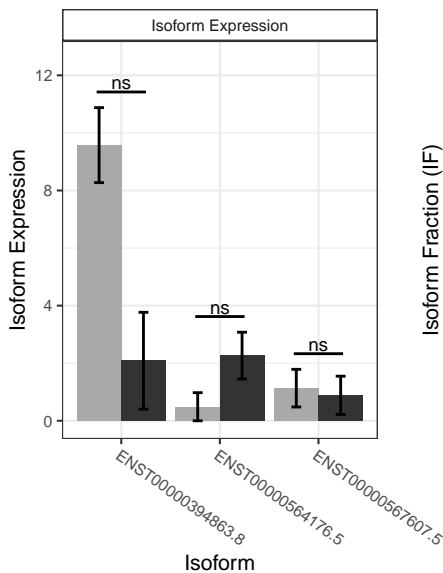
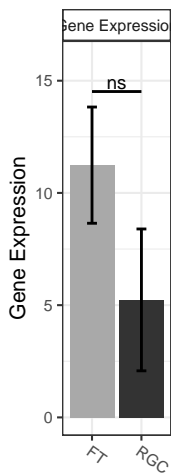
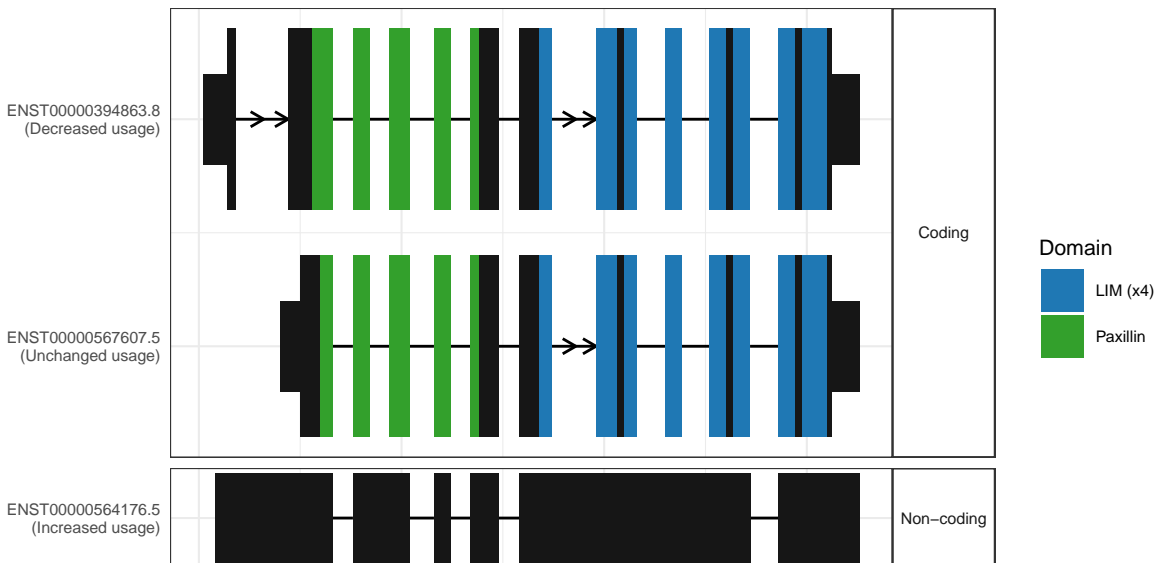


Domain



adh_short

The isoform switch in TGFB11 (FT vs RGC)



Domain

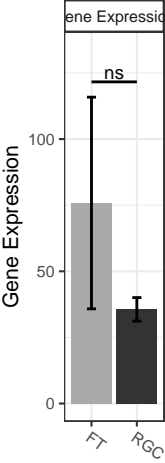
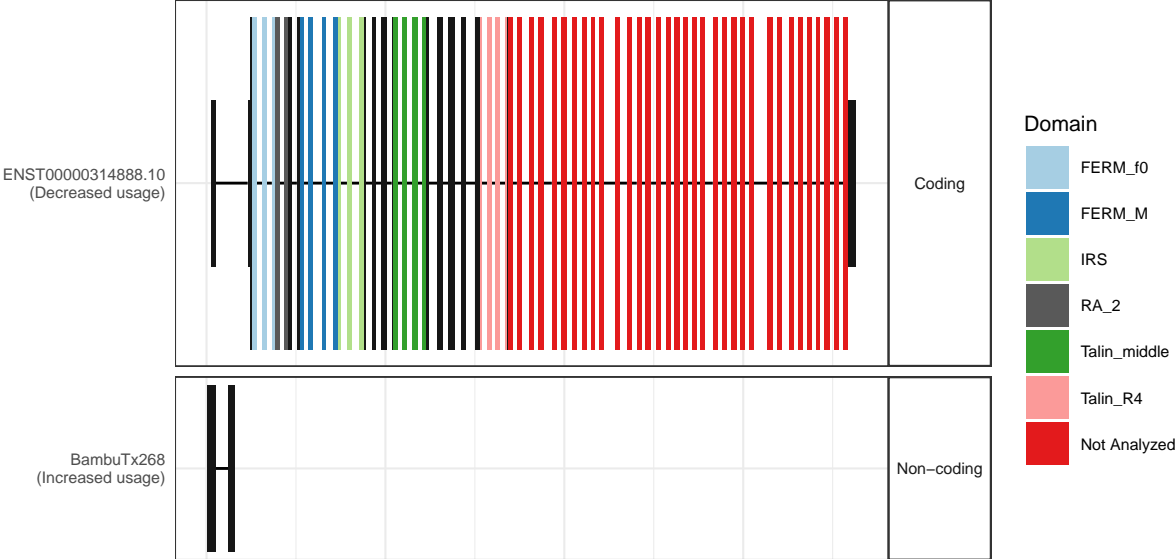


LIM (x4)

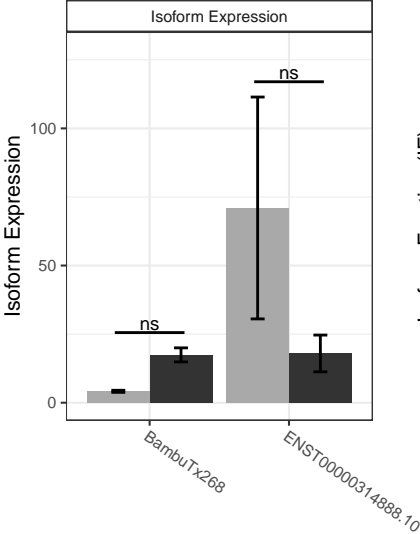


Paxillin

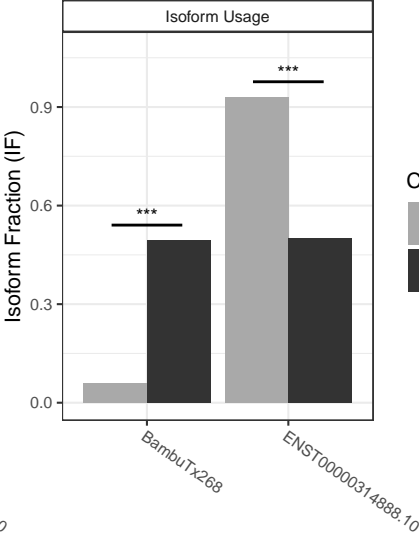
The isoform switch in TLN1 (FT vs RGC)



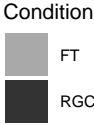
Condition










Isoform



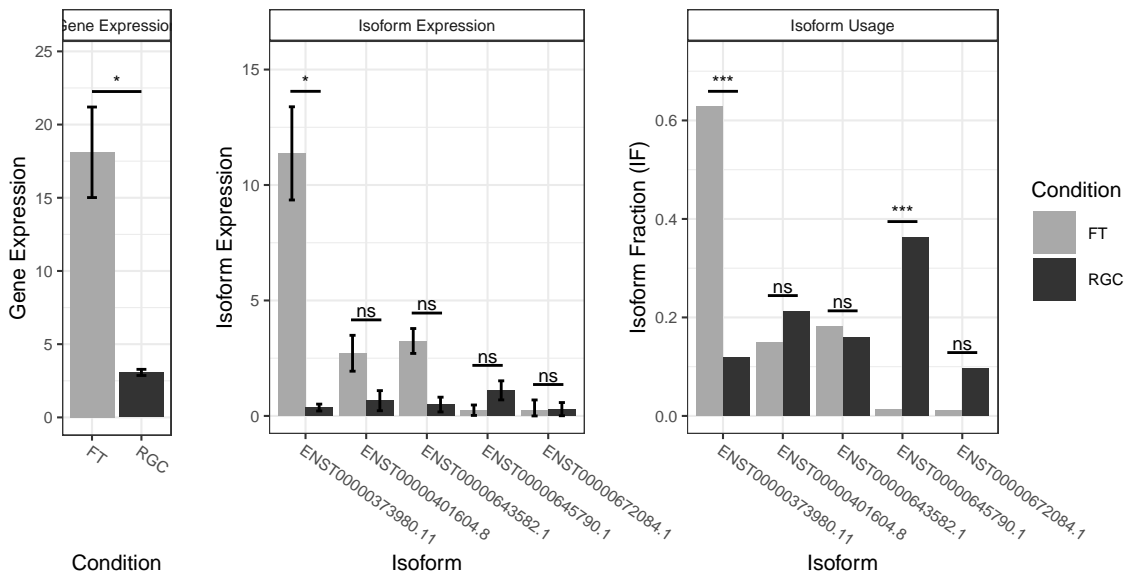
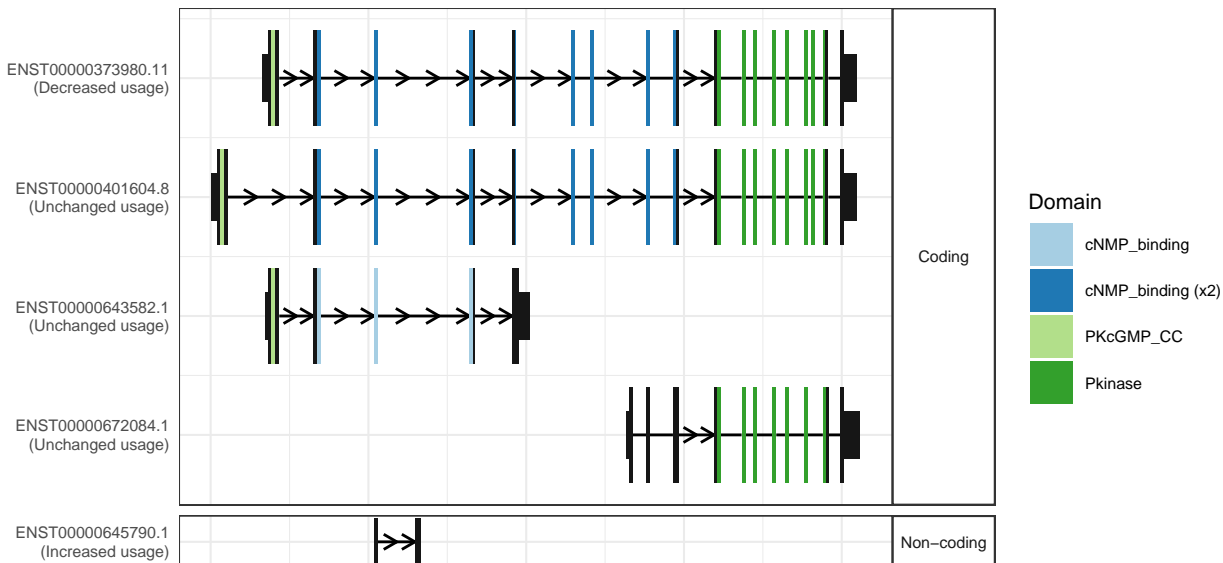
Isoform







Domain

	FERM_f0
	FERM_M
	IRS
	RA_2
	Talin_middle
	Talin_R4
	Not Analyzed

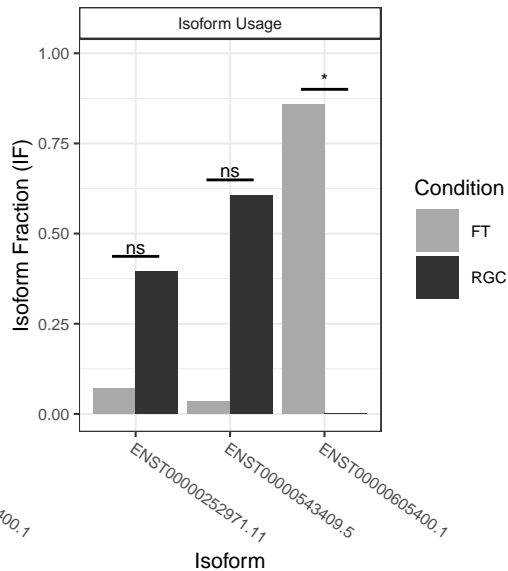
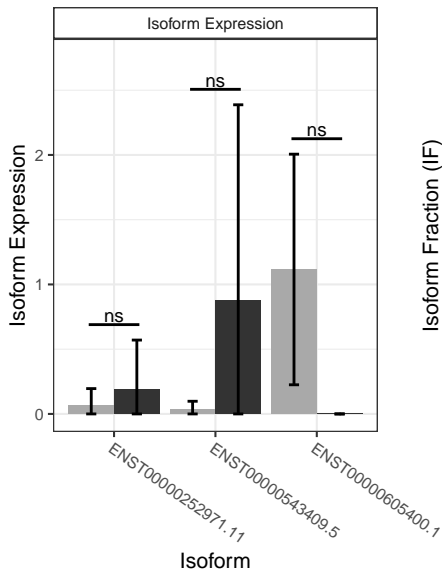
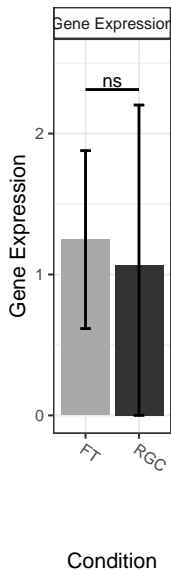
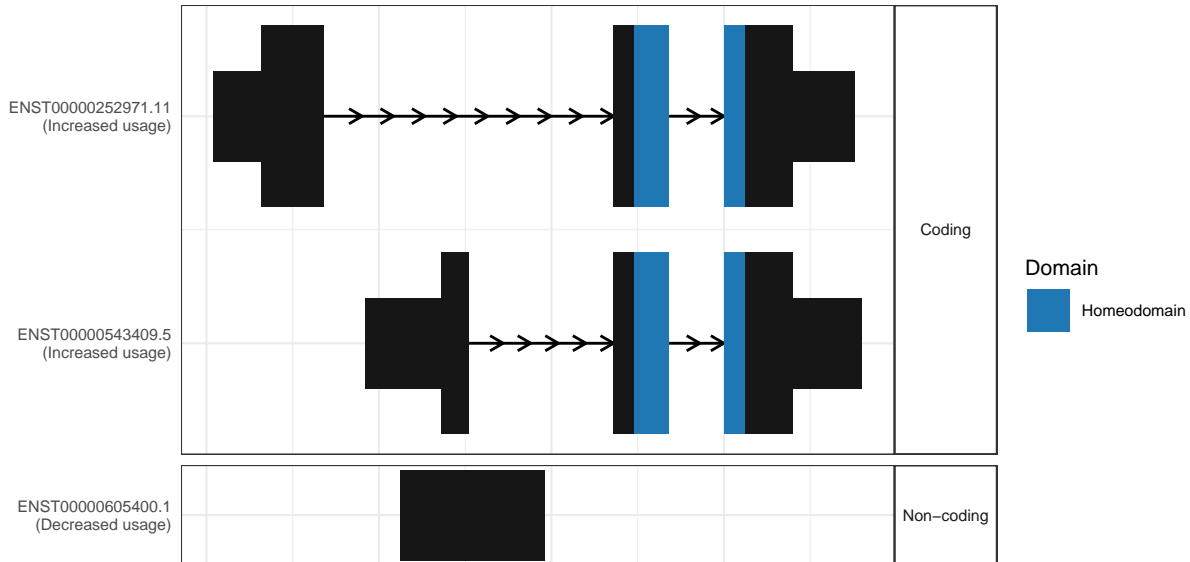
The isoform switch in PRKG1 (FT vs RGC)



Domain

	cNMP_binding
	cNMP_binding (x2)
	PKcGMP_CC
	Pkinase

The isoform switch in MNX1 (FT vs RGC)

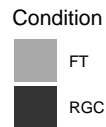
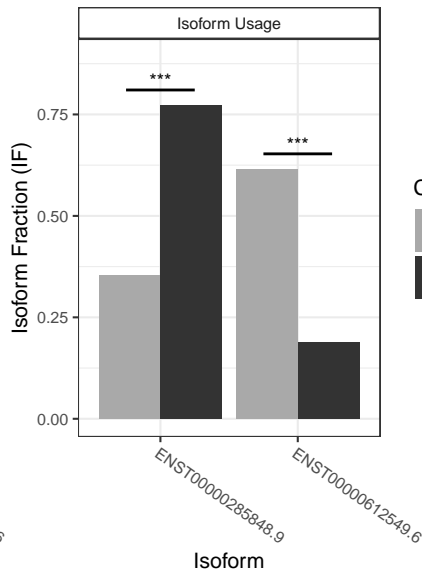
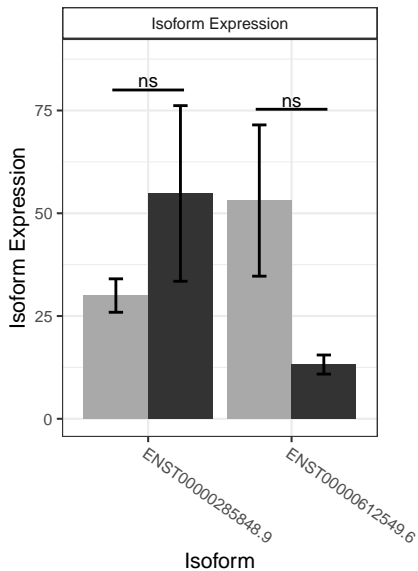
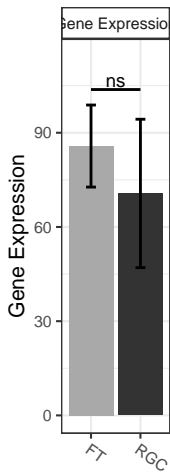
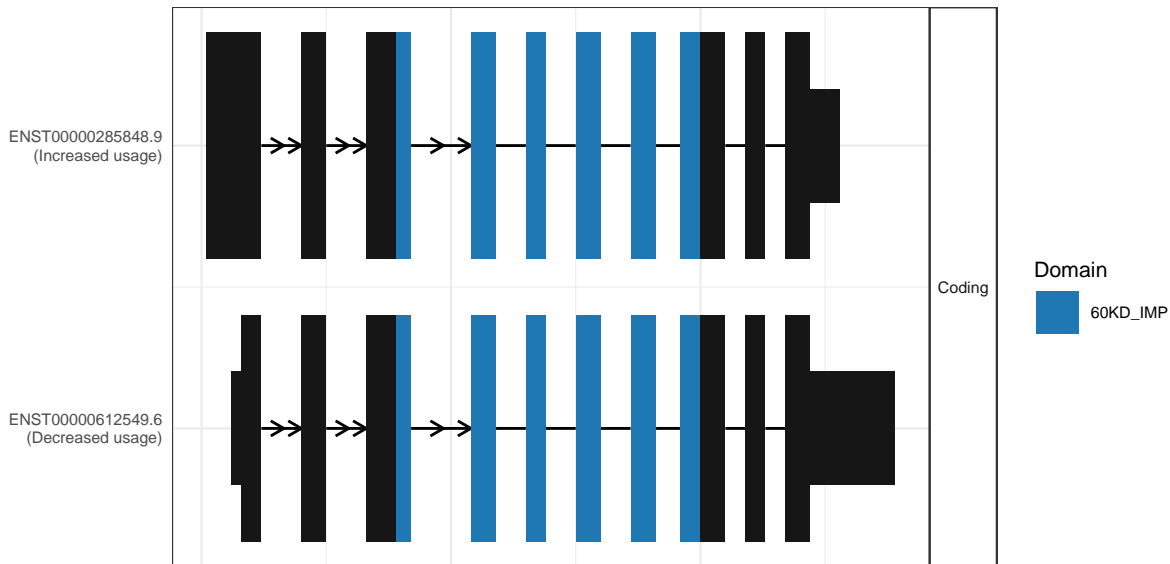


Domain



Homeodomain

The isoform switch in OXA1L (FT vs RGC)

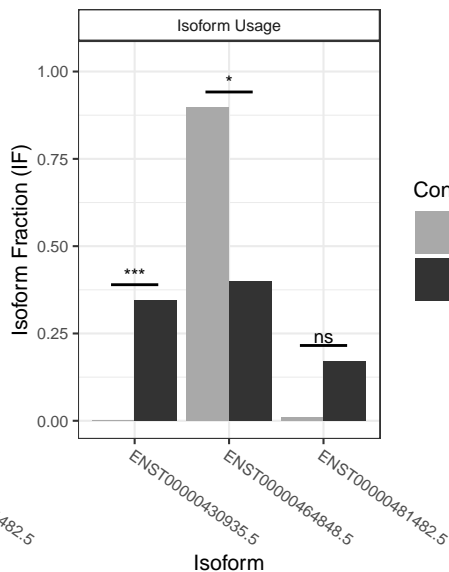
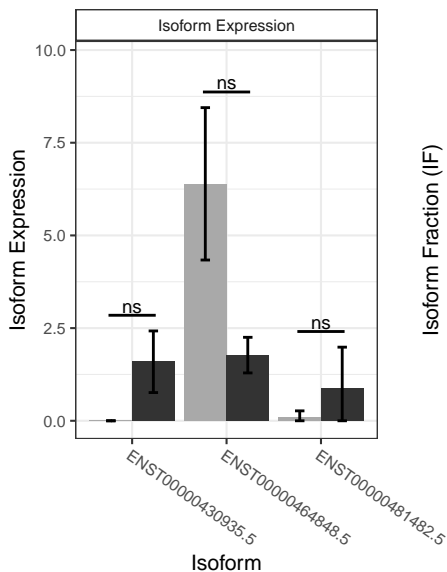
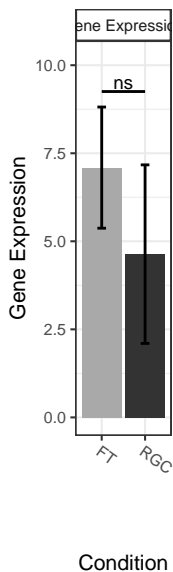
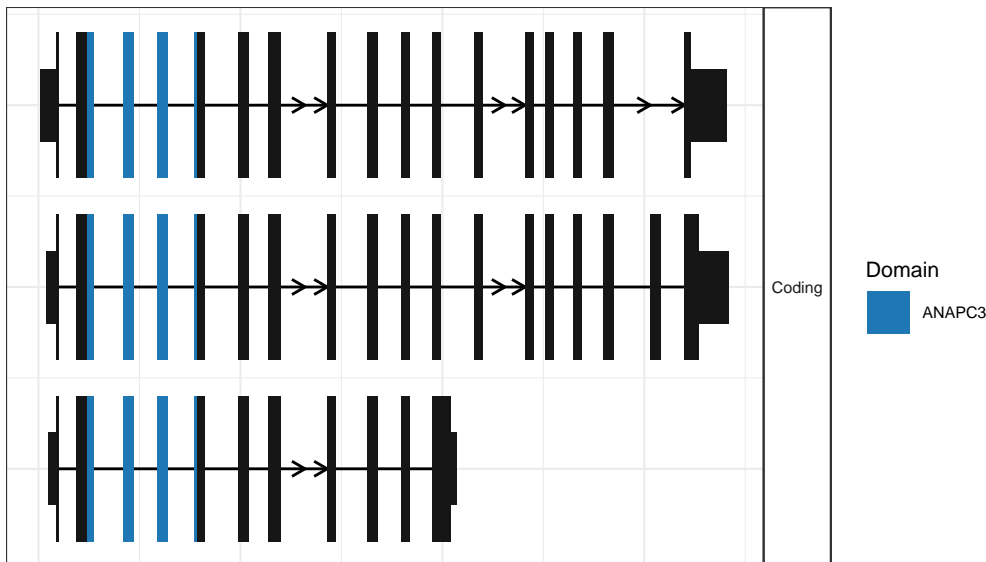


Domain



60KD_IMP

The isoform switch in IFT56 (FT vs RGC)

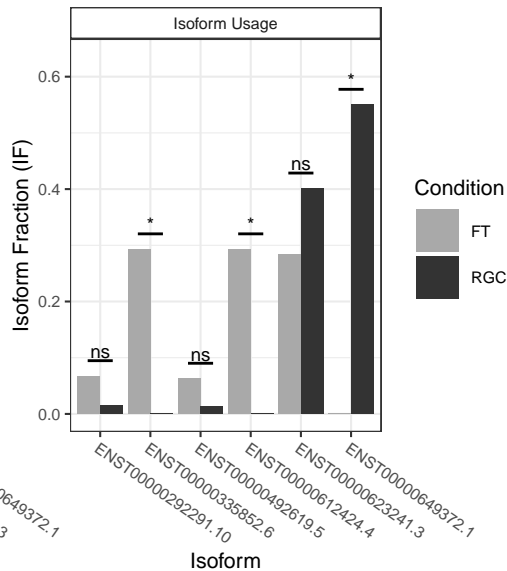
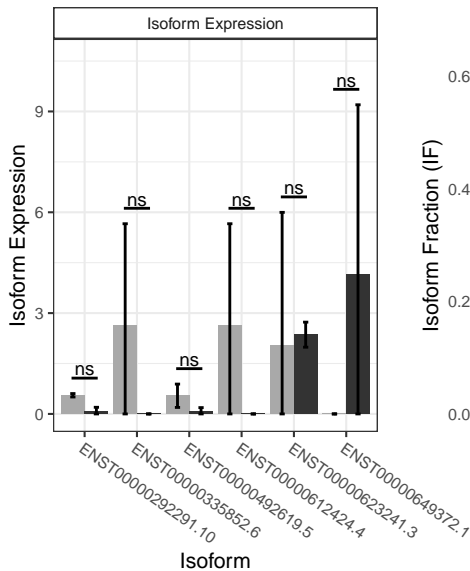
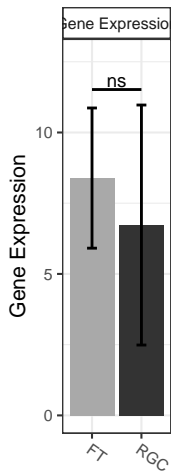
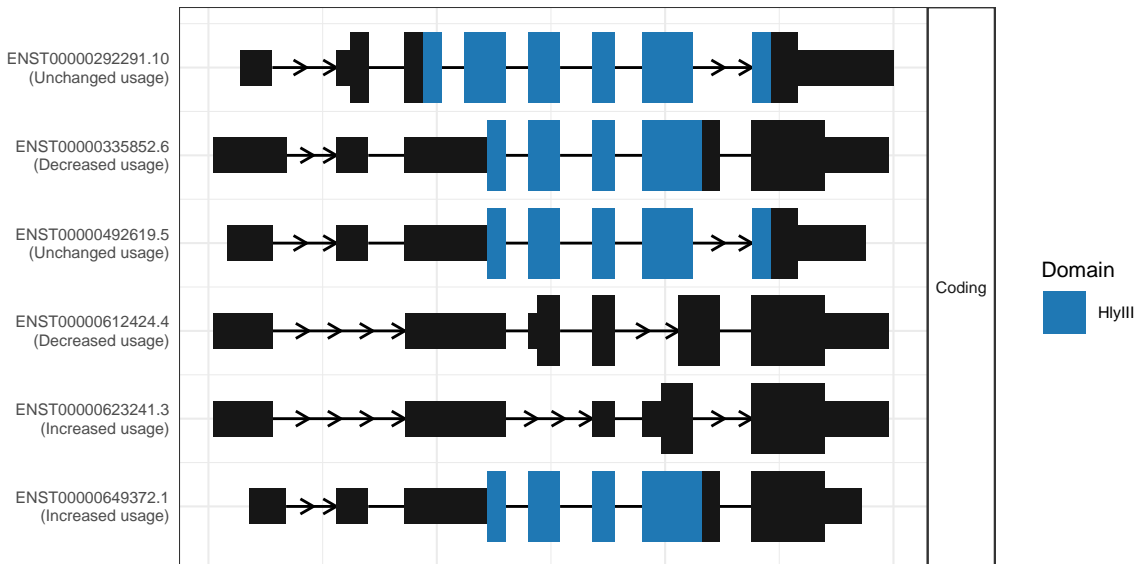


Domain



ANAPC3

The isoform switch in PAQR6 (FT vs RGC)

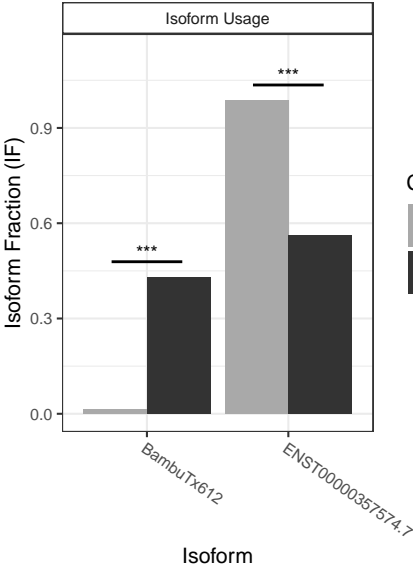
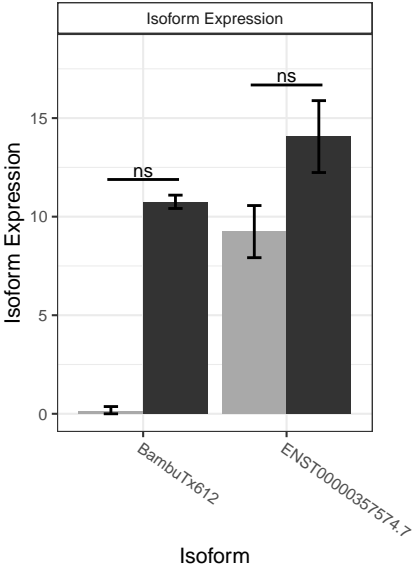
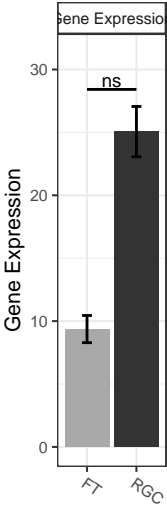
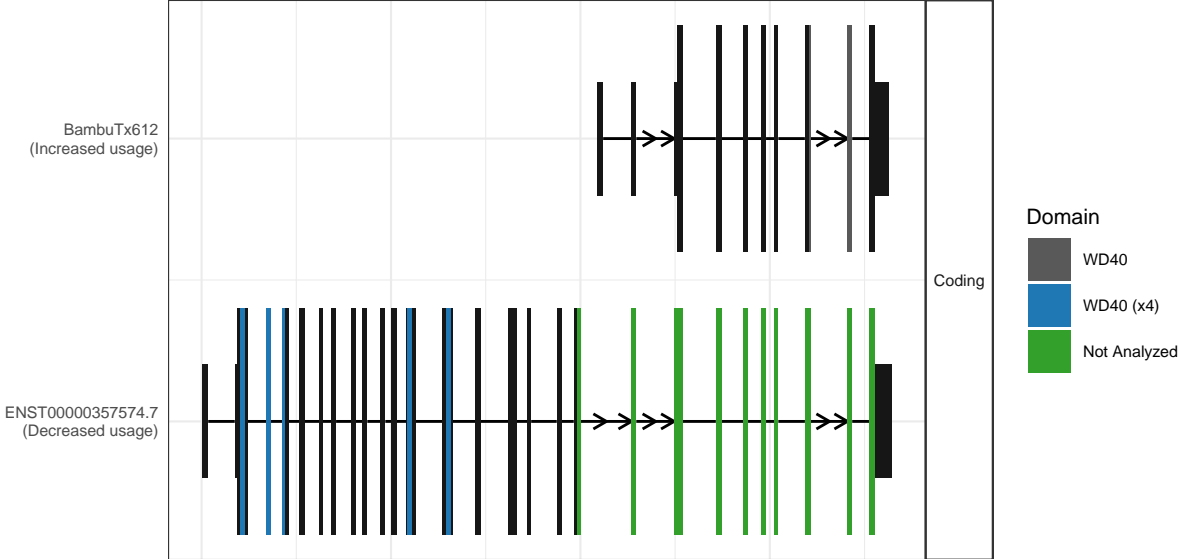


Domain



HlyIII

The isoform switch in WDR7 (FT vs RGC)



Domain



WD40

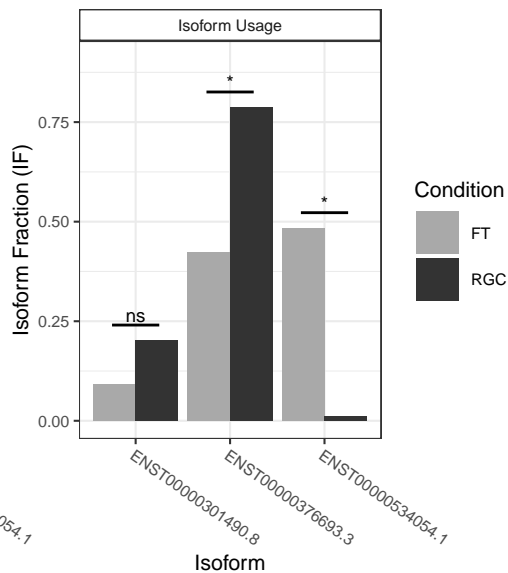
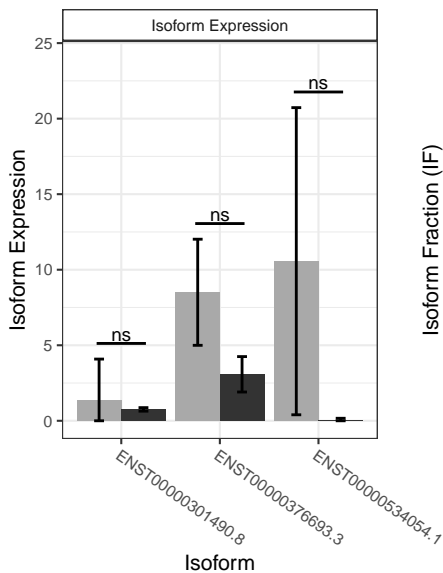
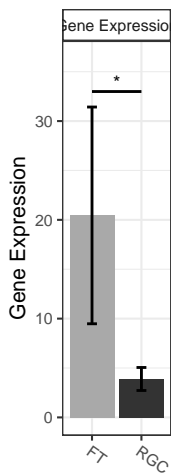
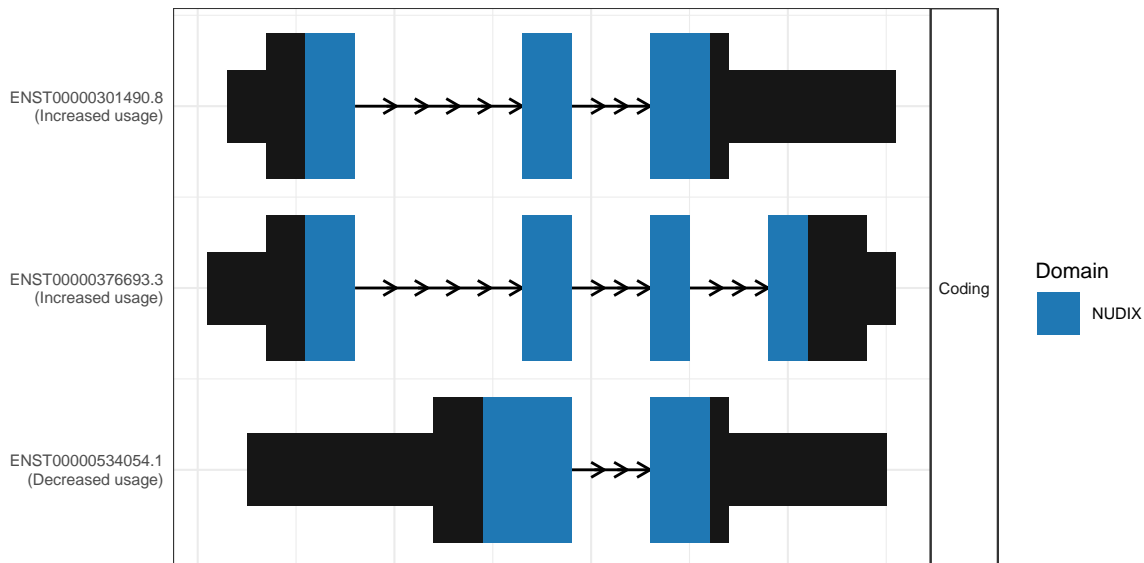


WD40 (x4)



Not Analyzed

The isoform switch in NUDT8 (FT vs RGC)

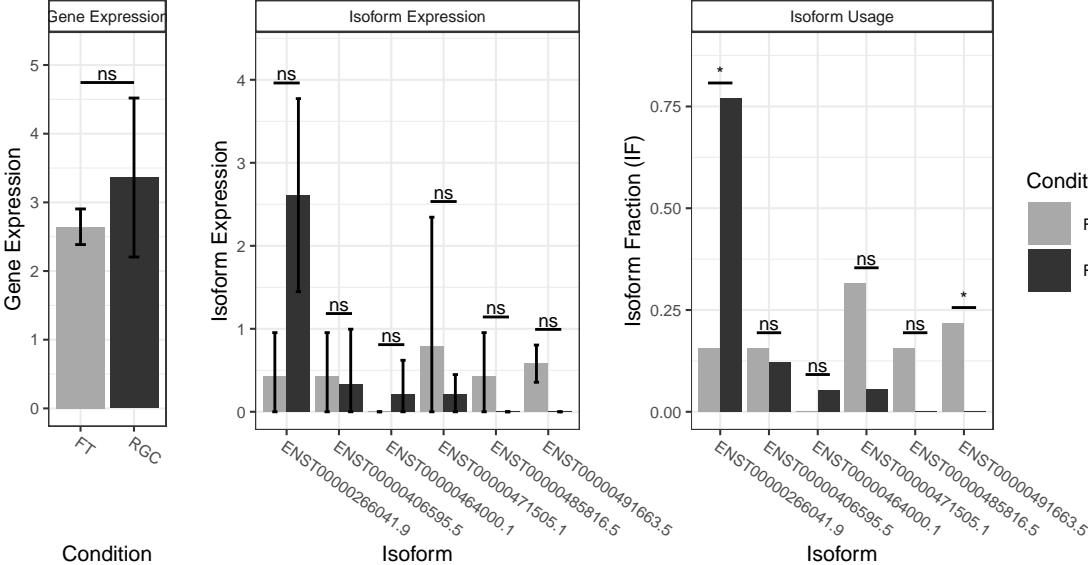
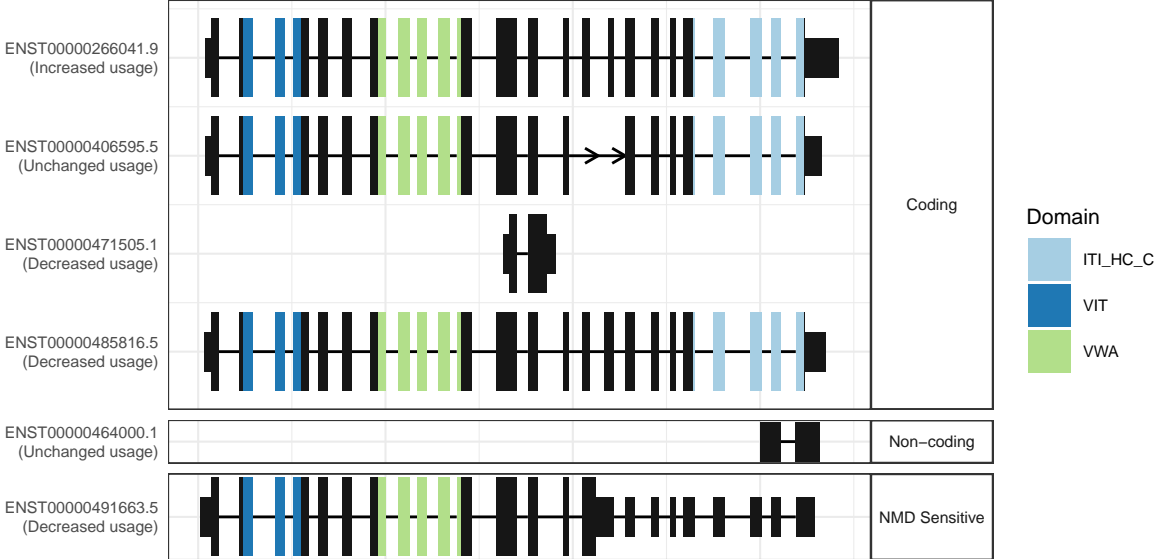


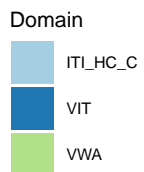
Domain



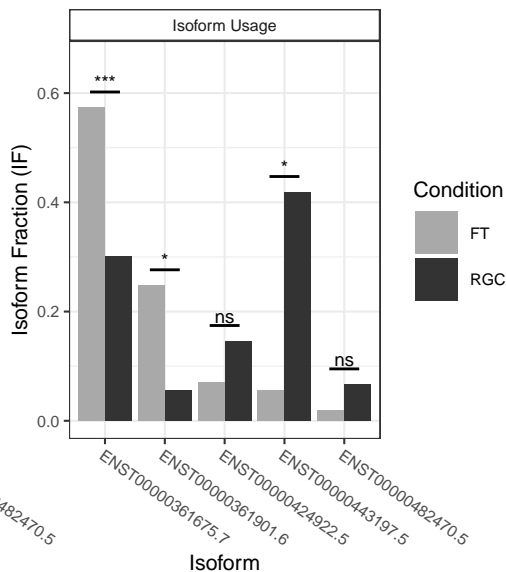
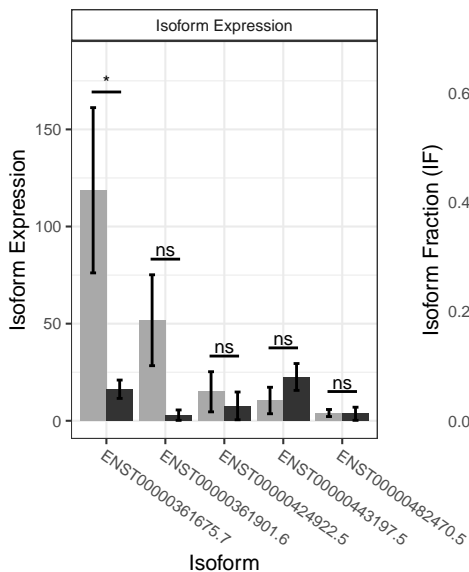
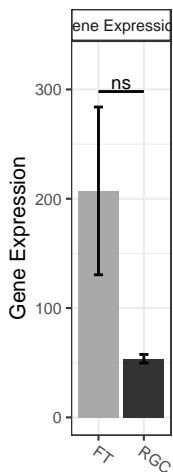
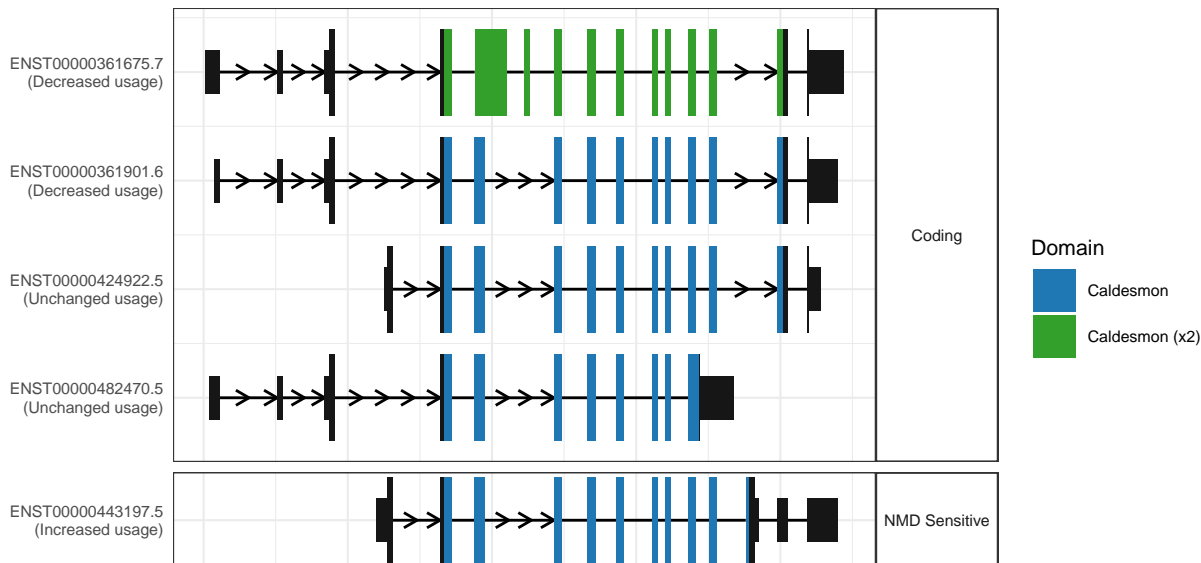
NUDIX

The isoform switch in ITIH4 (FT vs RGC)





The isoform switch in CALD1 (FT vs RGC)



Domain

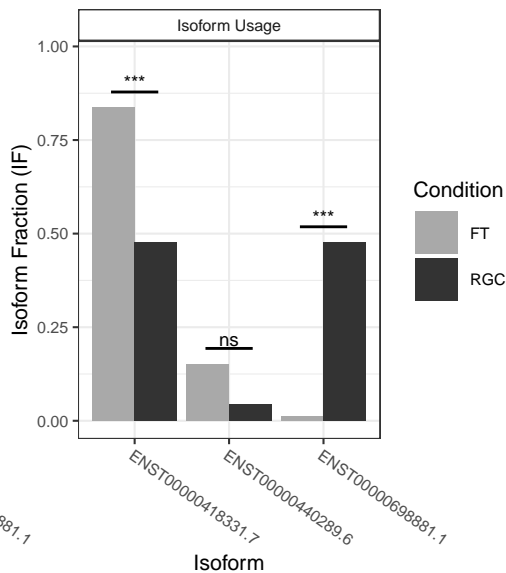
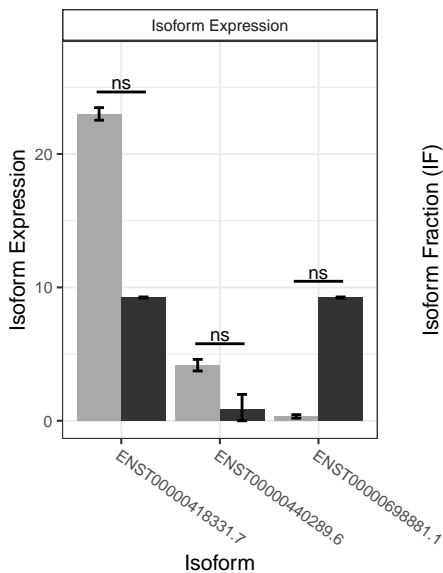
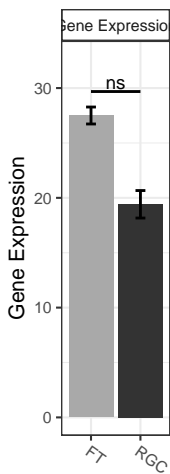
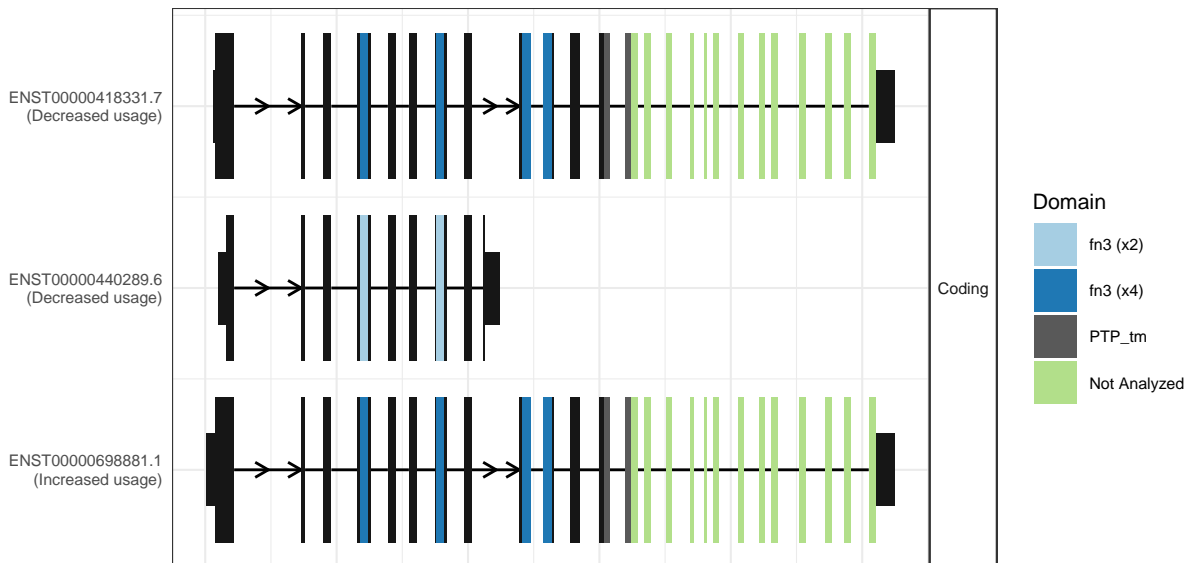


Caldesmon

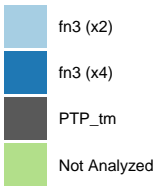


Caldesmon (x2)

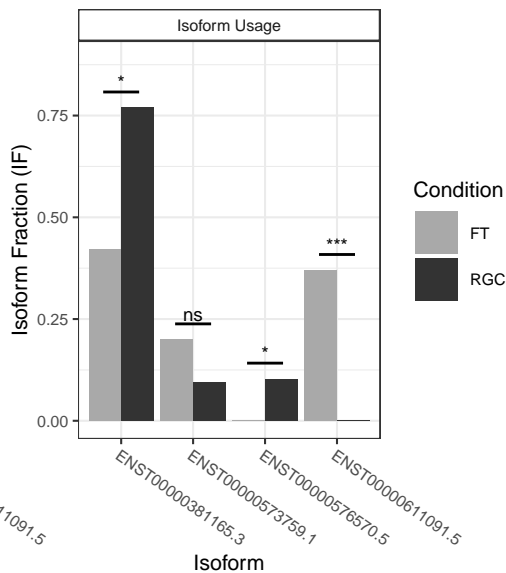
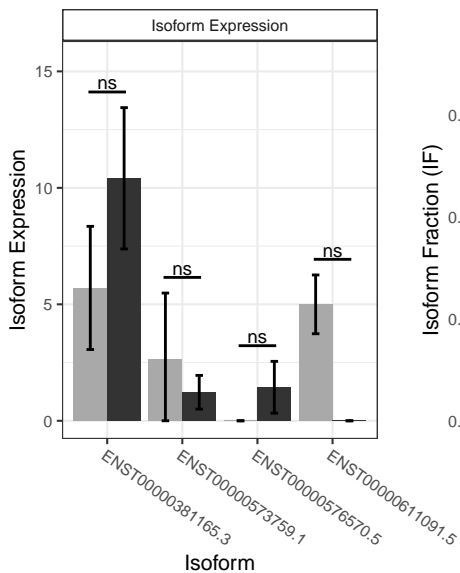
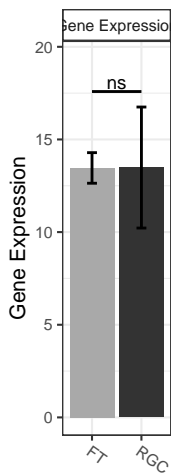
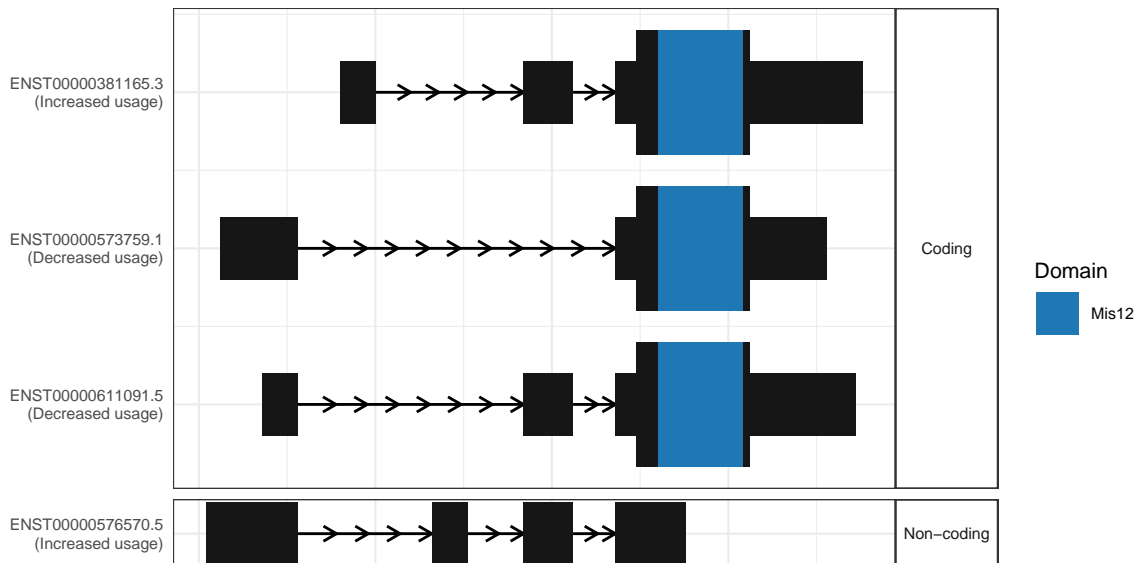
The isoform switch in PTPRJ (FT vs RGC)



Domain



The isoform switch in MIS12 (FT vs RGC)

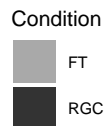
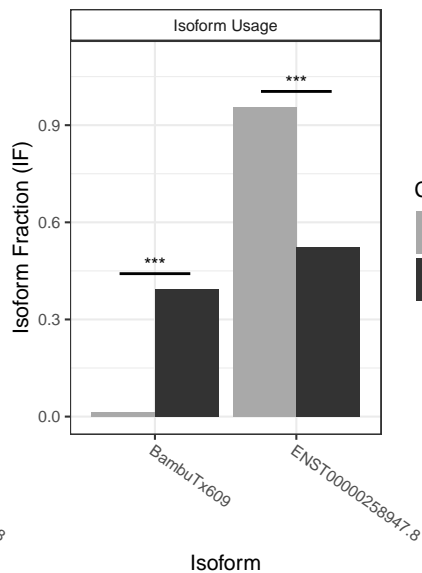
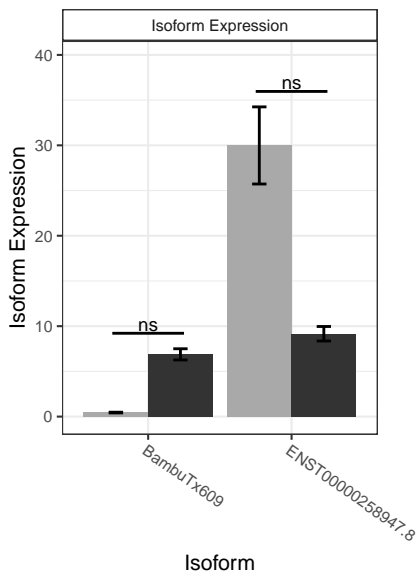
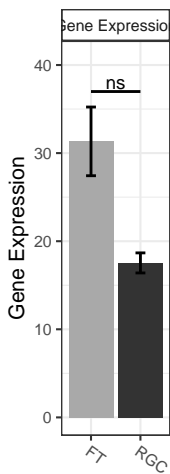
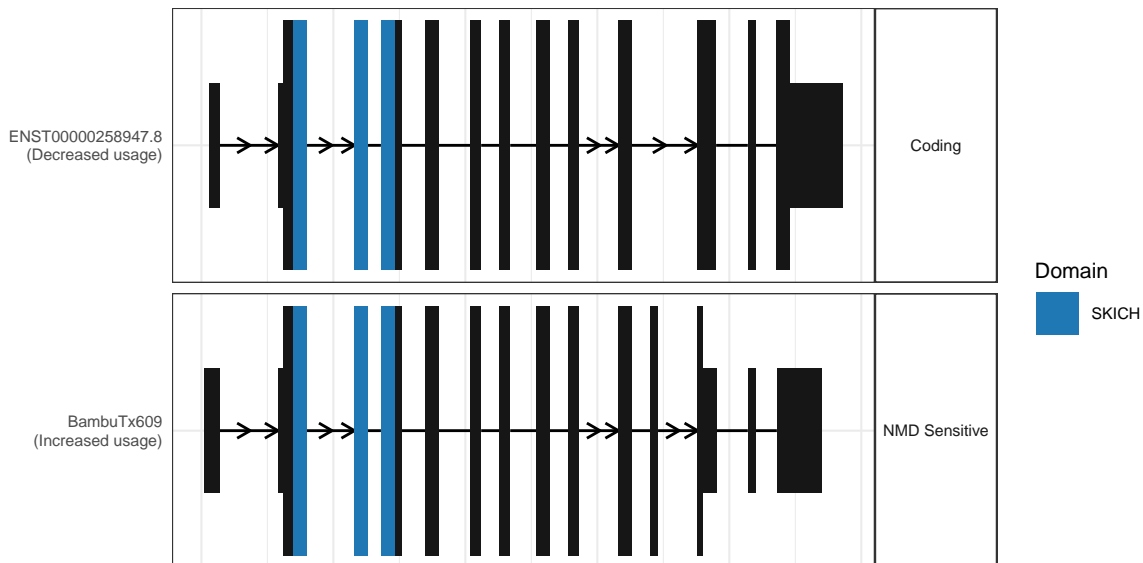


Domain



Mis12

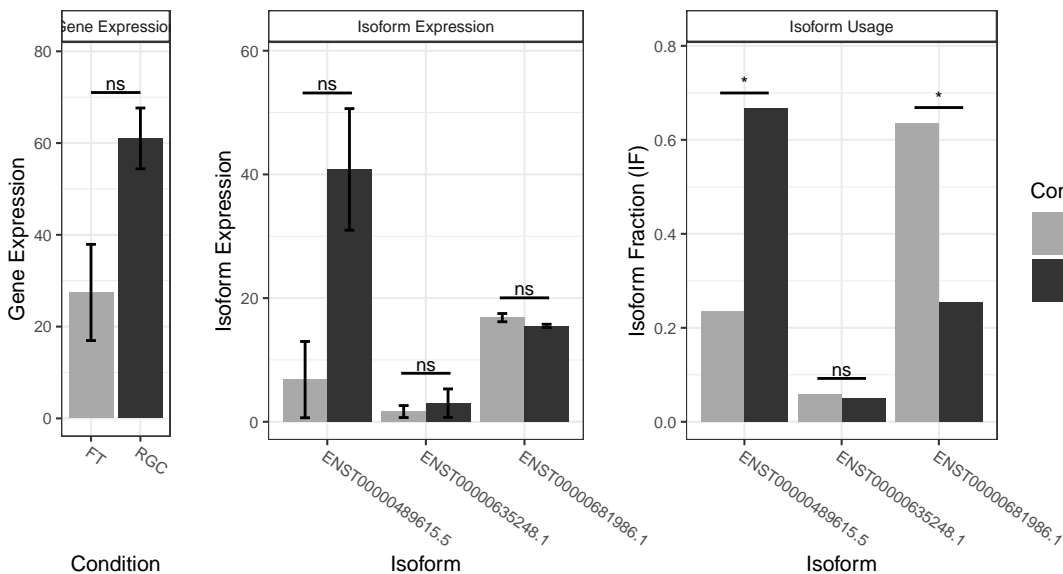
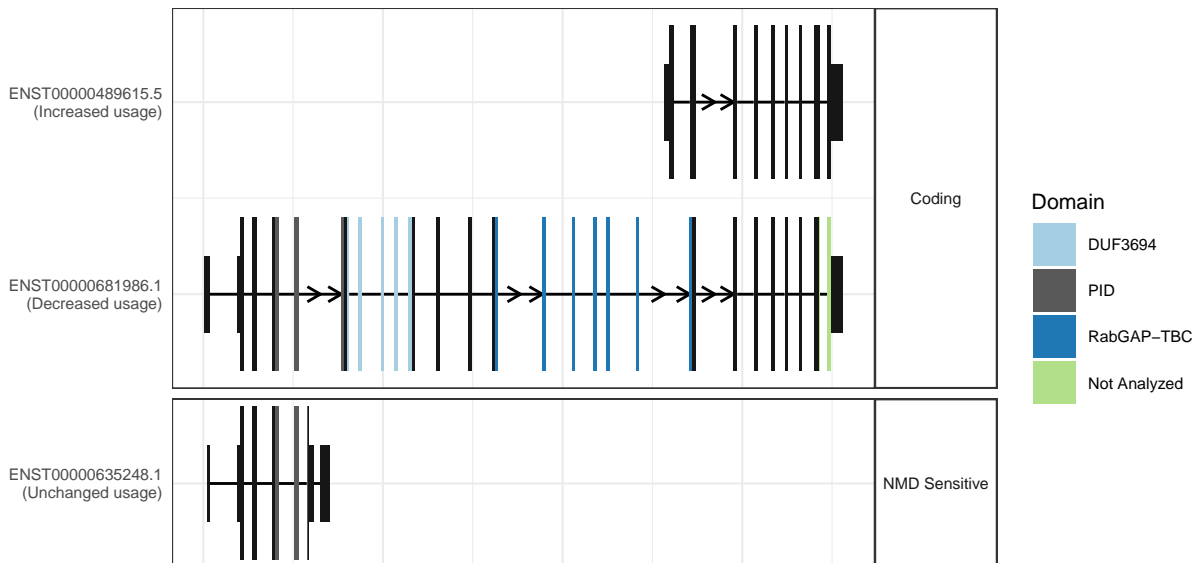
The isoform switch in CALCOCO2 (FT vs RGC)



Domain

 SKICH

The isoform switch in RABGAP1L (FT vs RGC)



Domain



DUF3694



PID

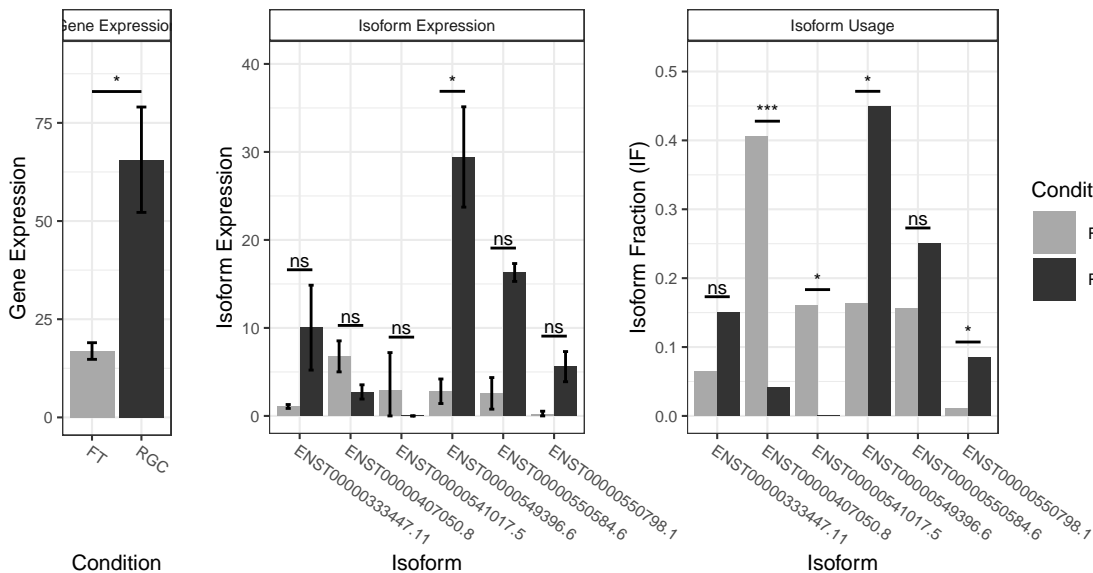
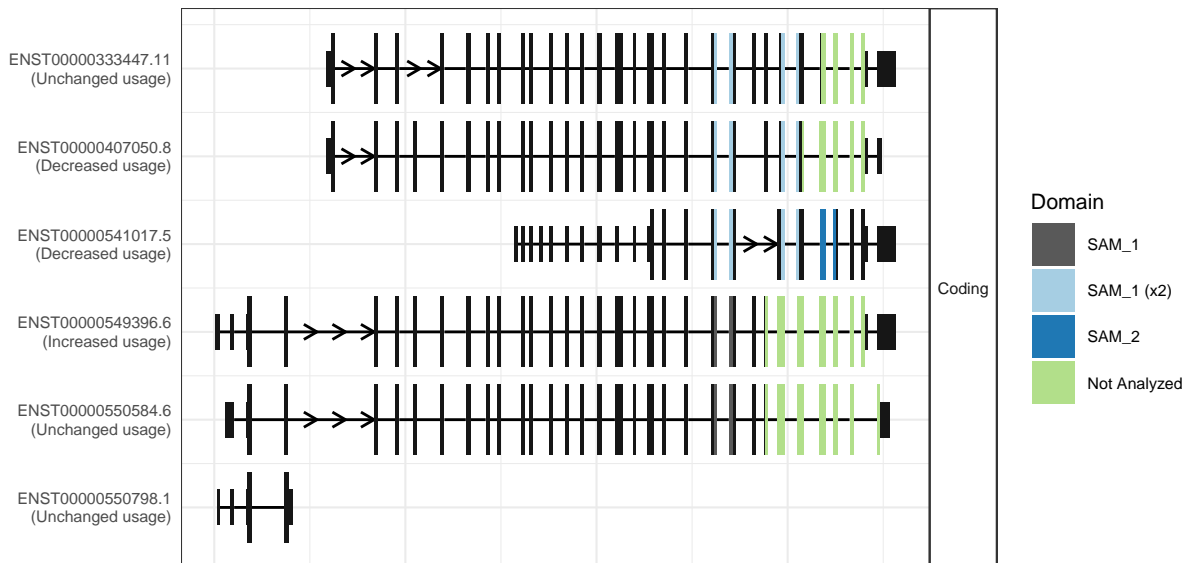


RabGAP-TBC



Not Analyzed

The isoform switch in PPFIA2 (FT vs RGC)



Domain



SAM_1



SAM_1 (x2)

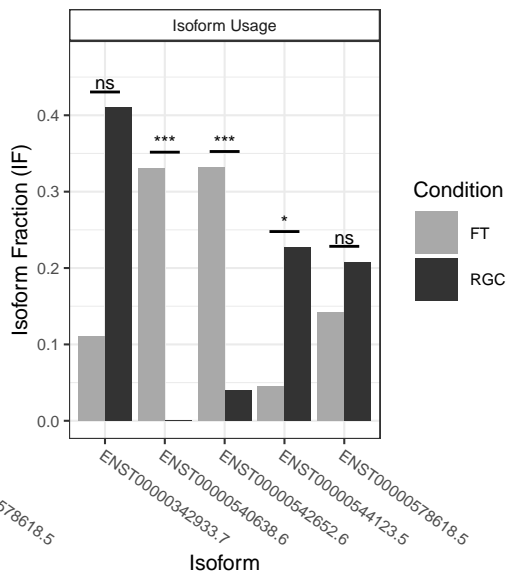
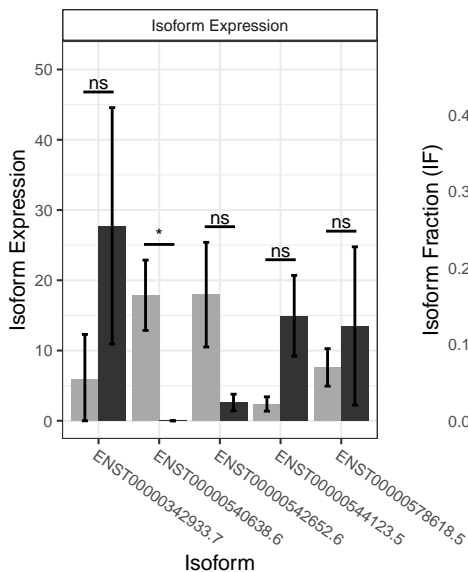
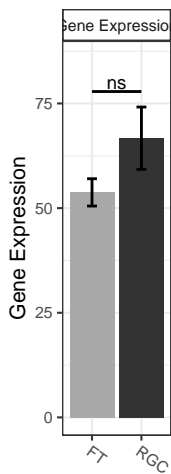
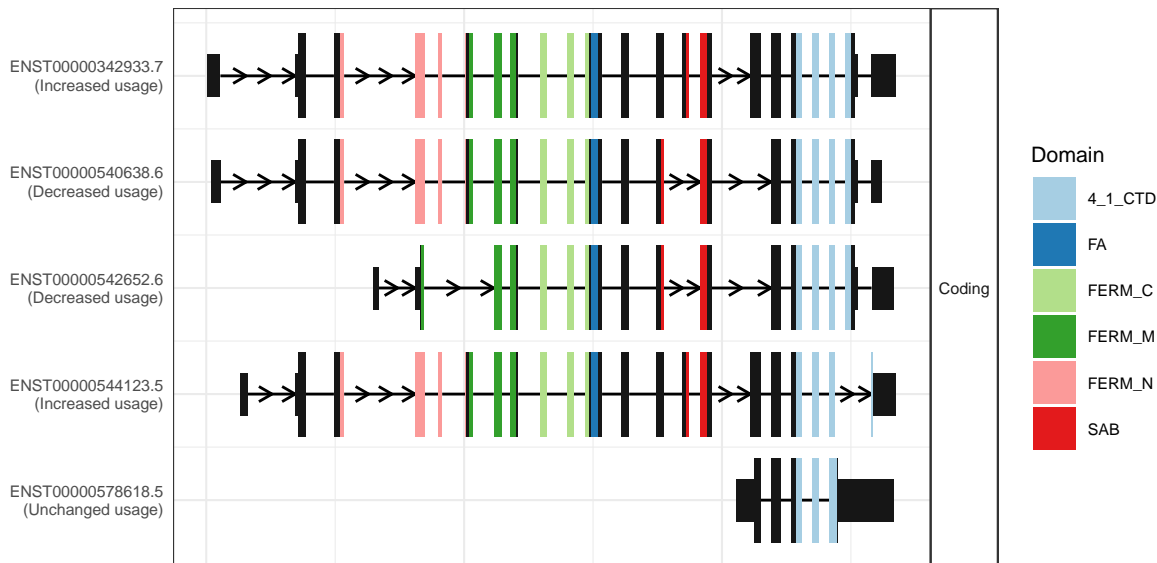


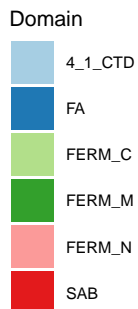
SAM_2



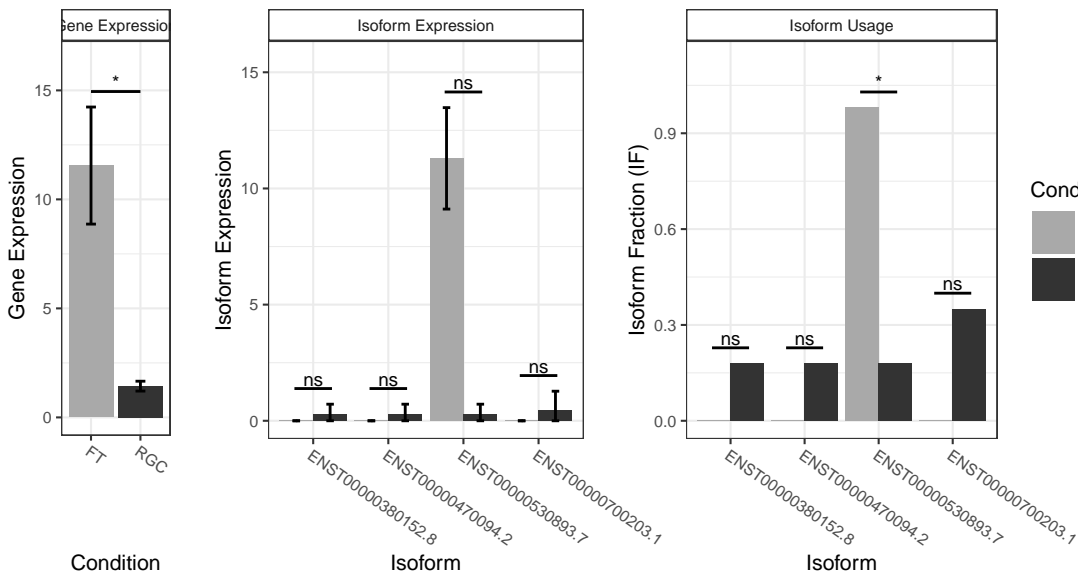
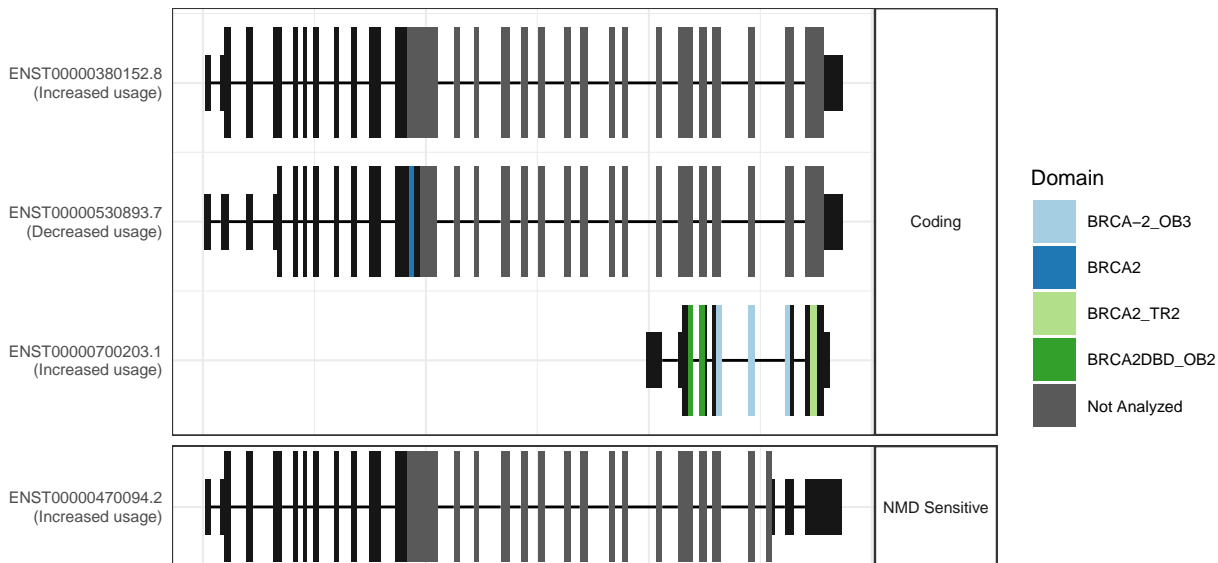
Not Analyzed

The isoform switch in EPB41L3 (FT vs RGC)

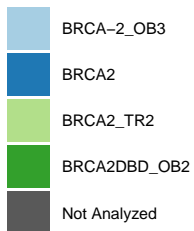




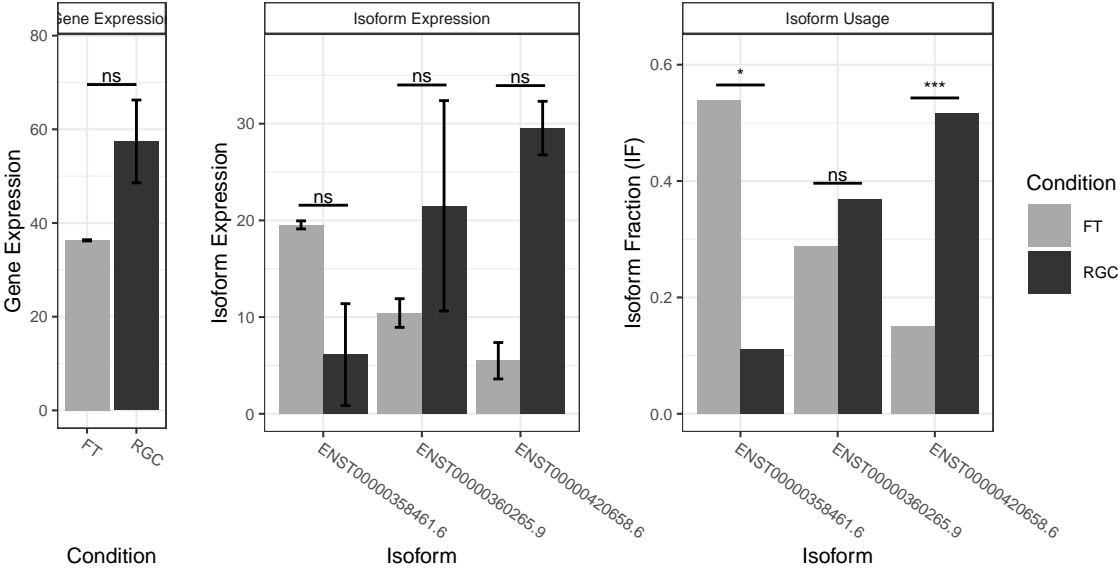
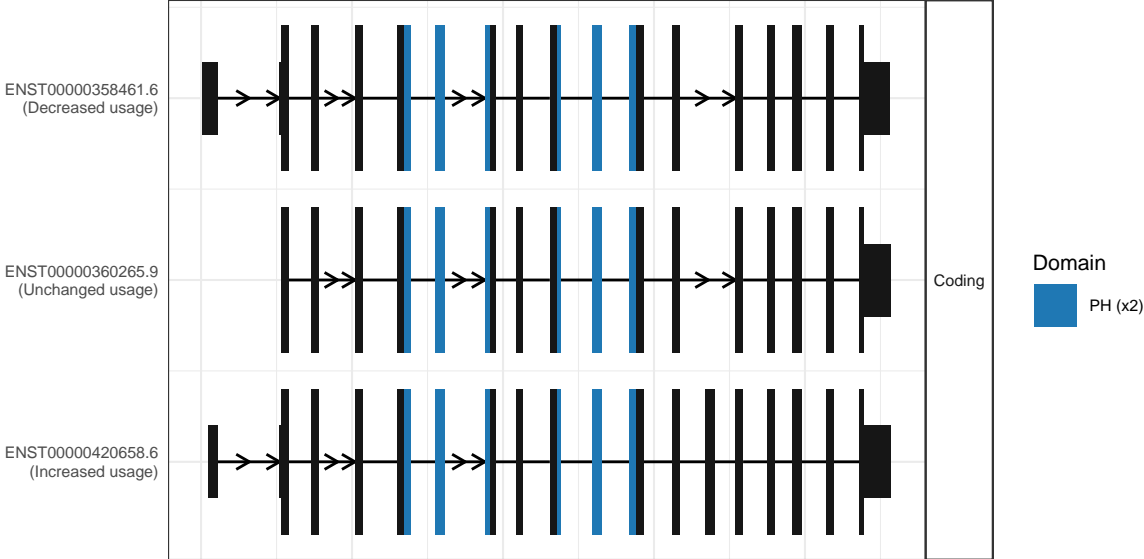
The isoform switch in BRCA2 (FT vs RGC)



Domain



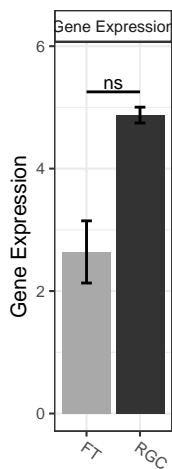
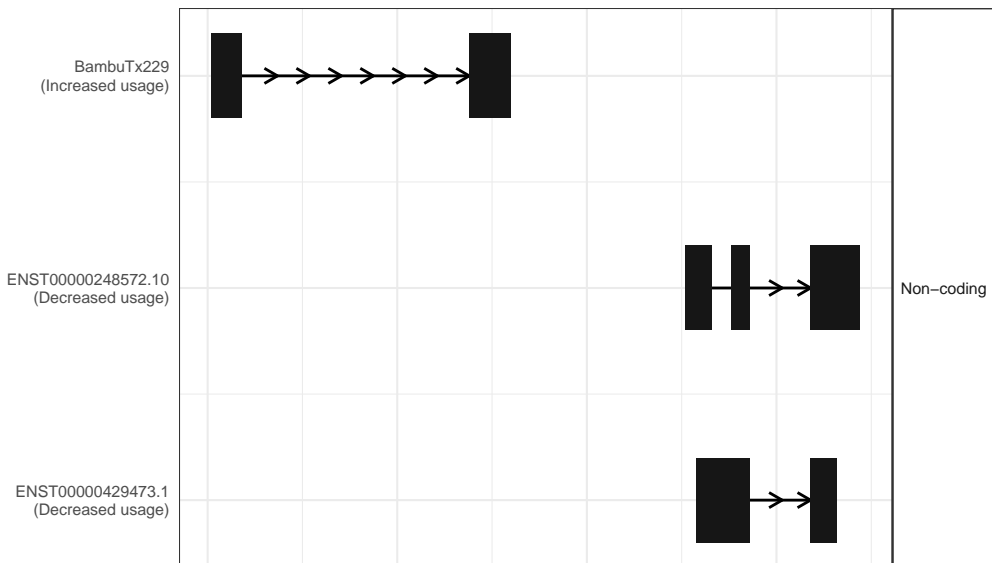
The isoform switch in AFAP1 (FT vs RGC)



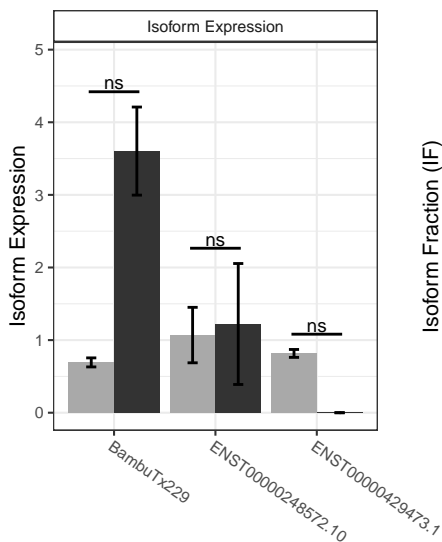
Domain

 PH (x2)

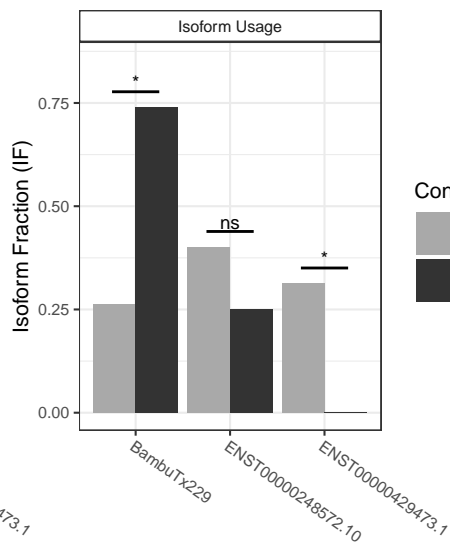
The isoform switch in GNGT1 (FT vs RGC)



Condition



Isoform

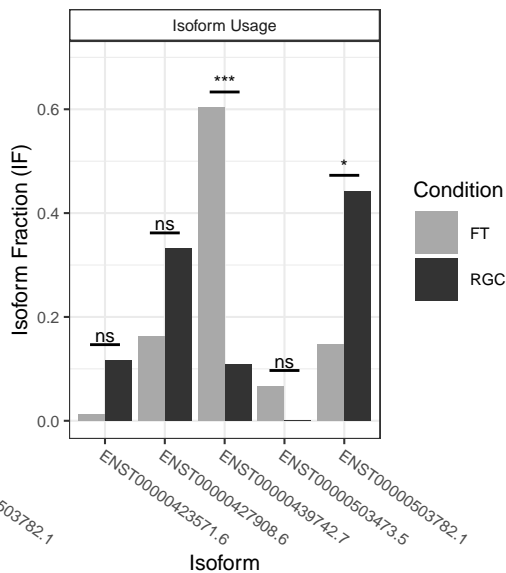
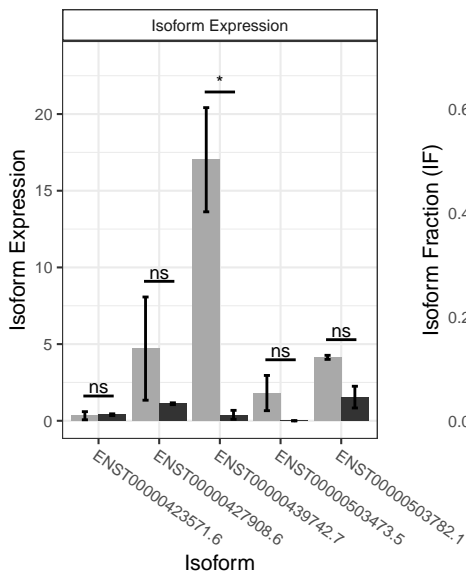
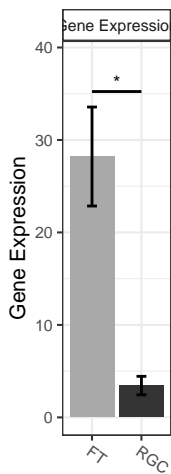
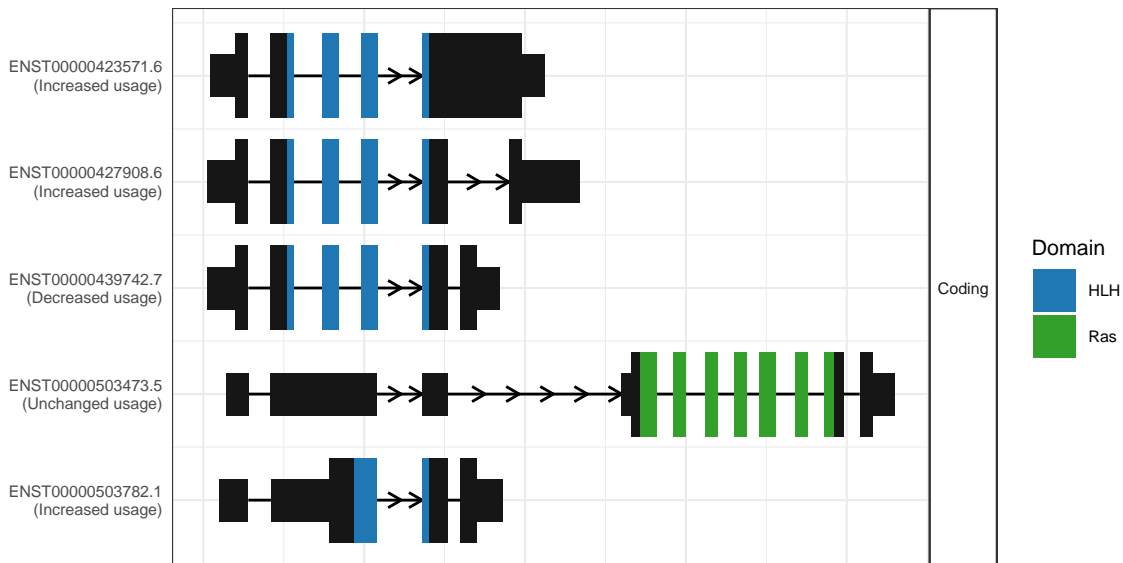


Isoform

Condition



The isoform switch in MXD3 (FT vs RGC)



Domain

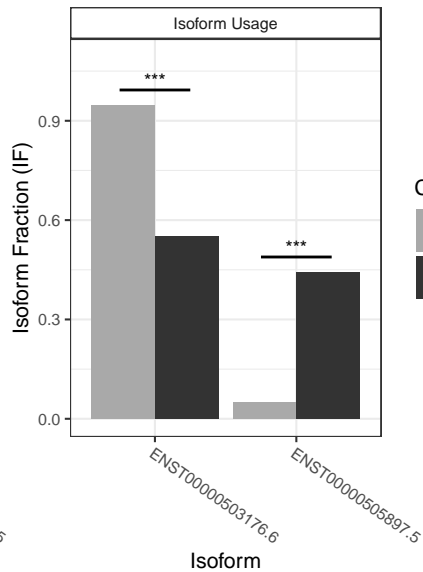
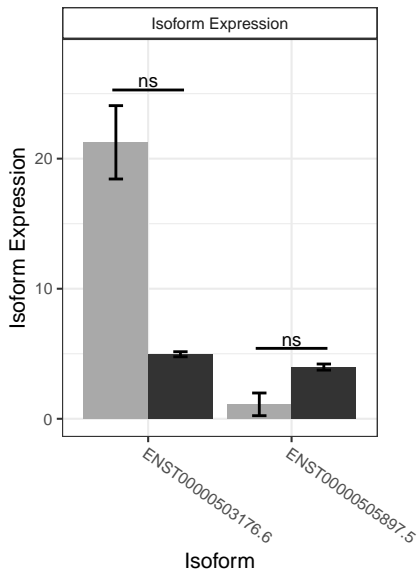
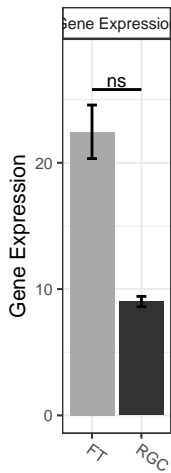
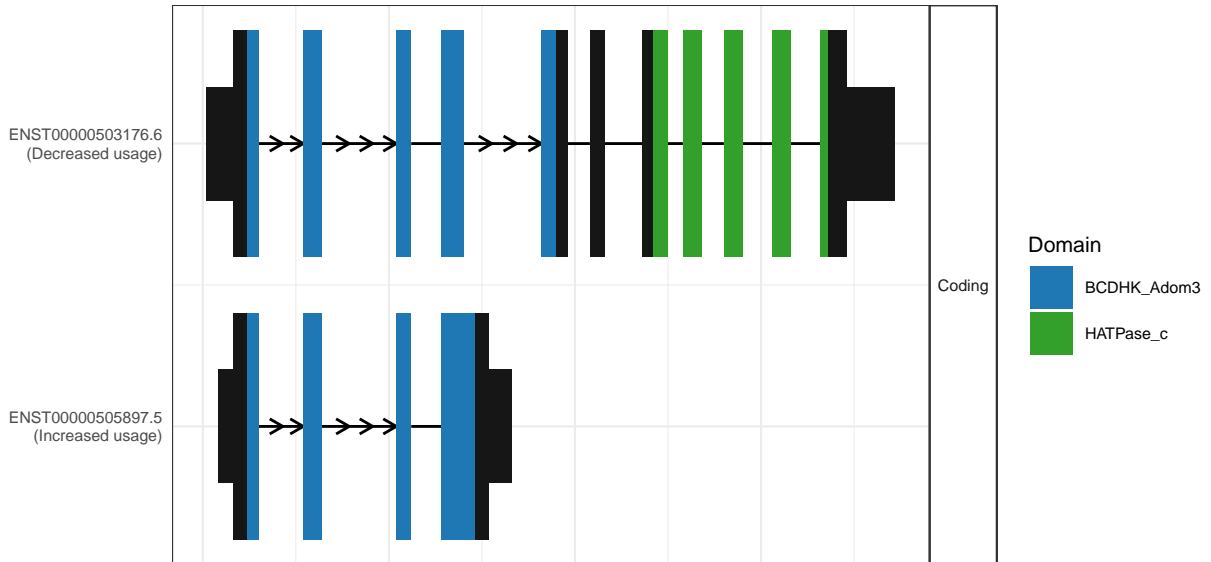


HLH

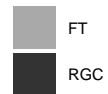


Ras

The isoform switch in PDK2 (FT vs RGC)



Condition



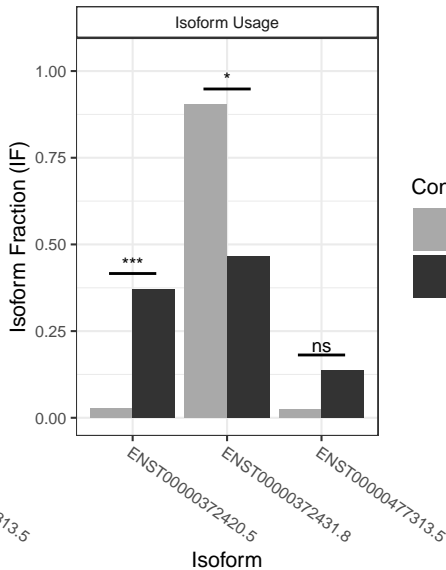
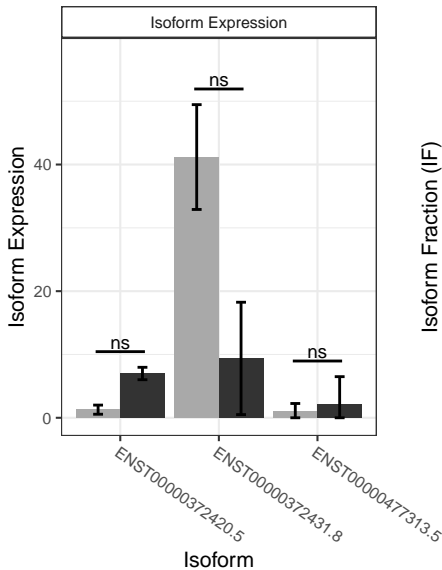
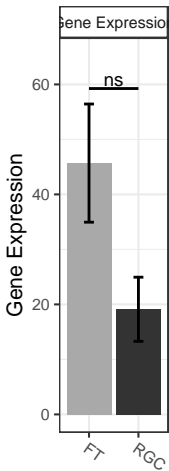
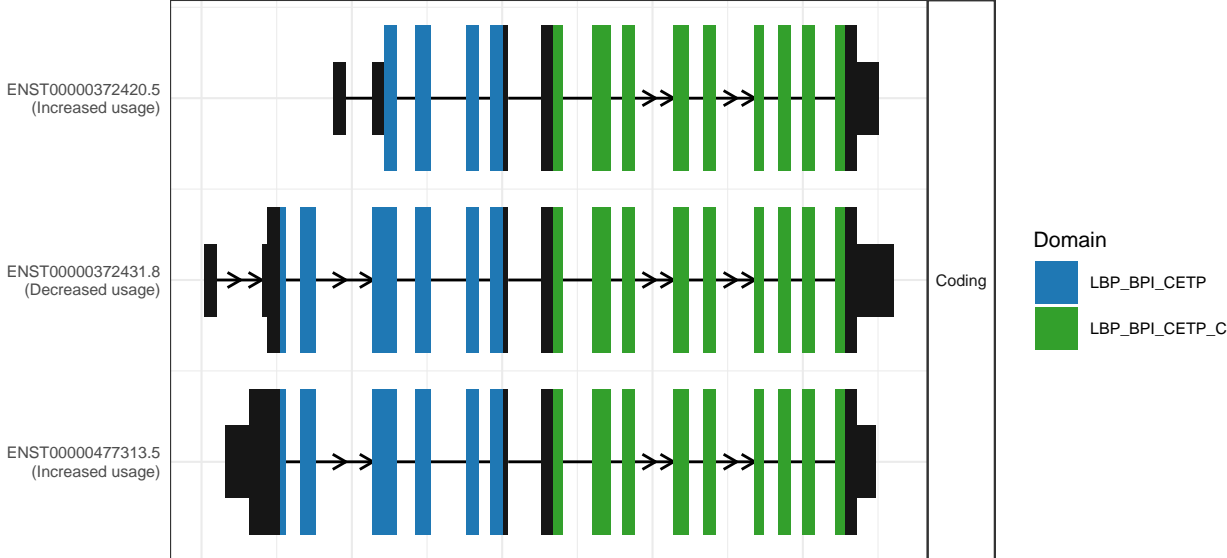
Domain



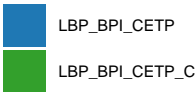
BCDHK_Adom3

HATPase_c

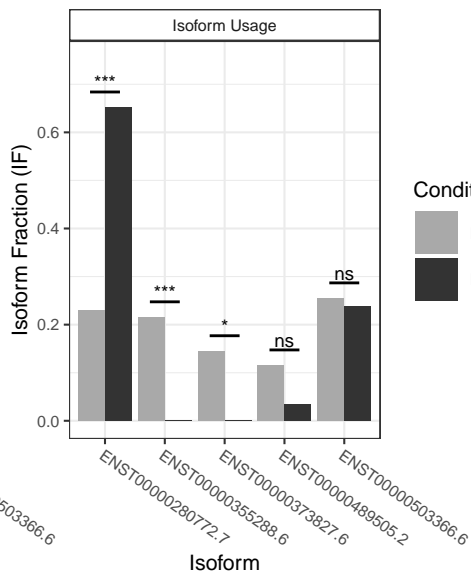
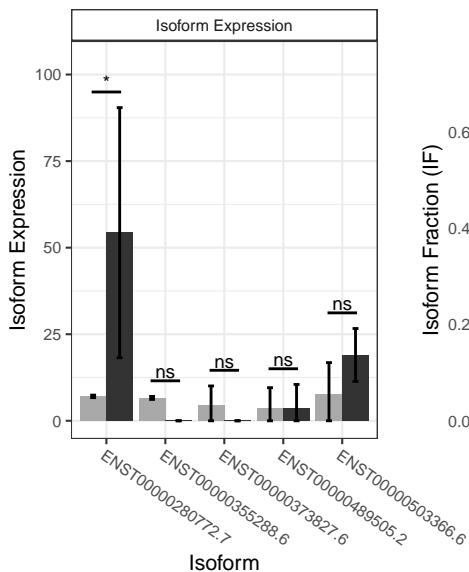
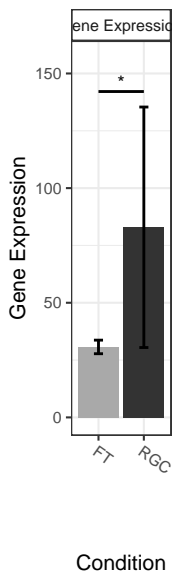
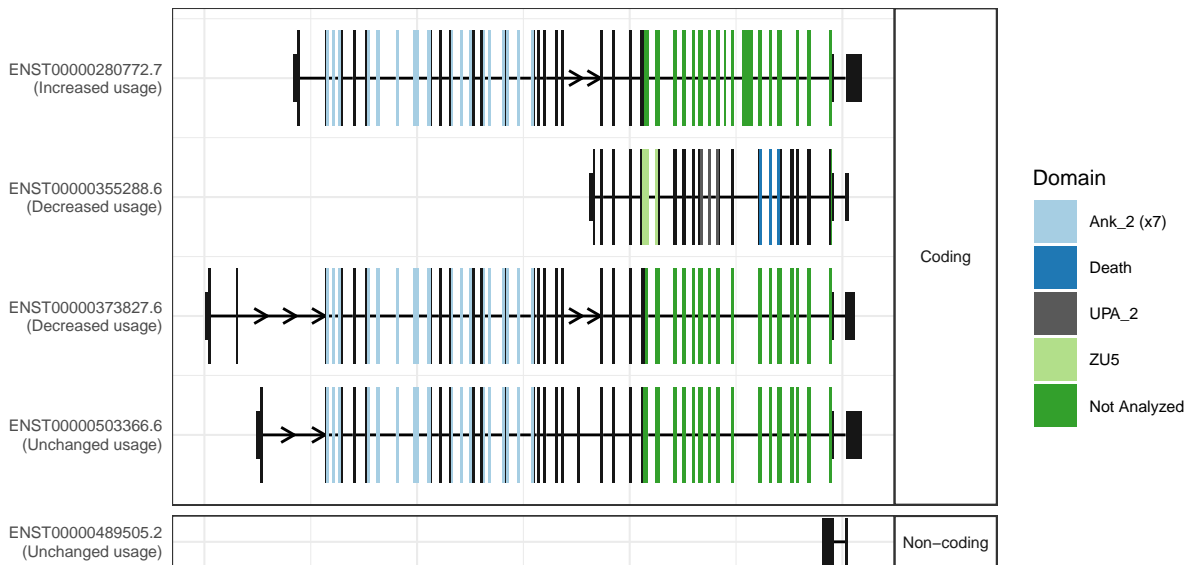
The isoform switch in PLTP (FT vs RGC)



Domain



The isoform switch in ANK3 (FT vs RGC)



Domain



Ank_2 (x7)



Death



UPA_2

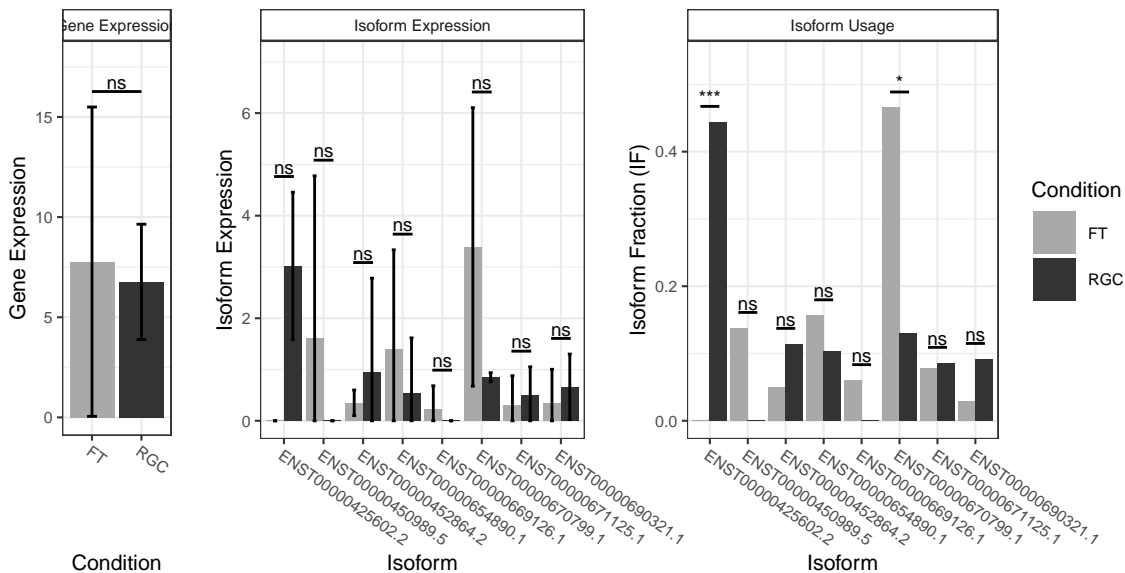
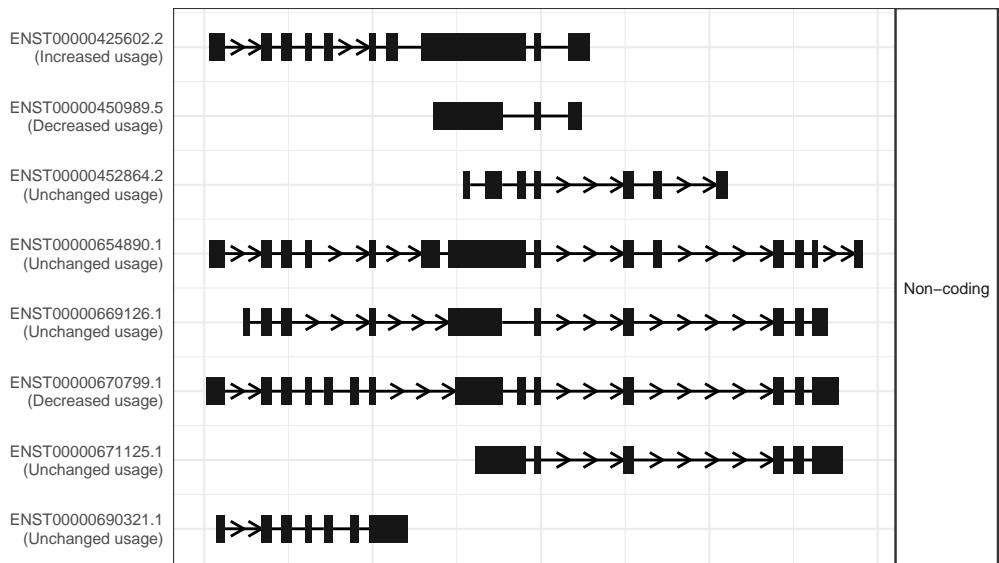


ZU5

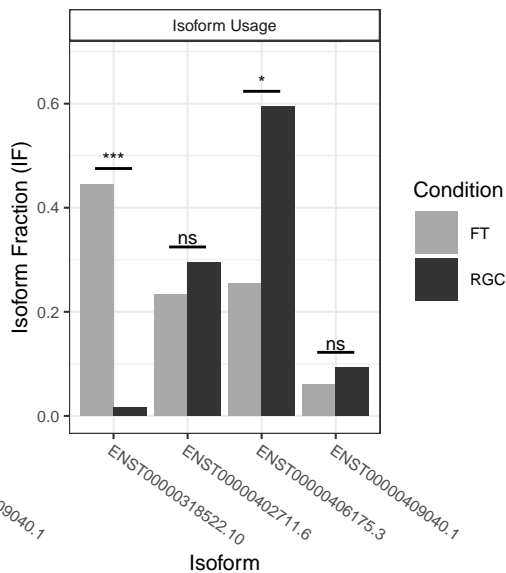
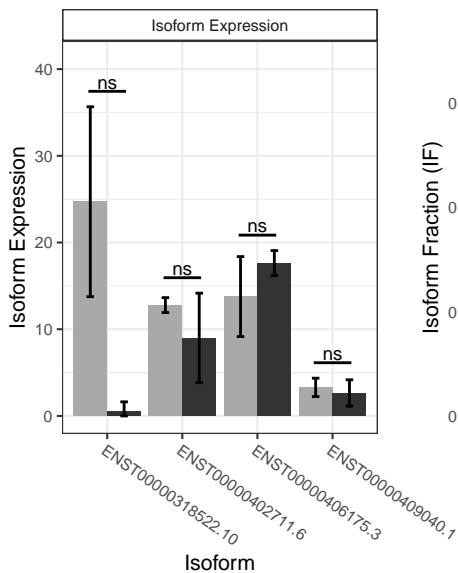
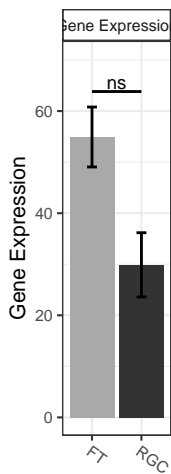
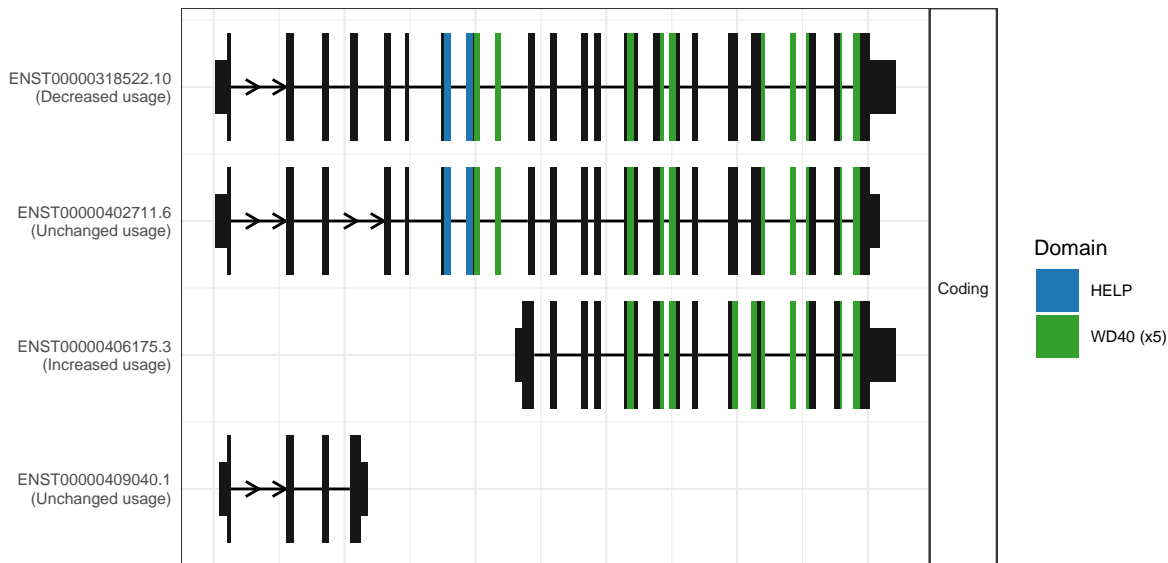


Not Analyzed

The isoform switch in EOLA2-DT (FT vs RGC)



The isoform switch in EML4 (FT vs RGC)



Domain

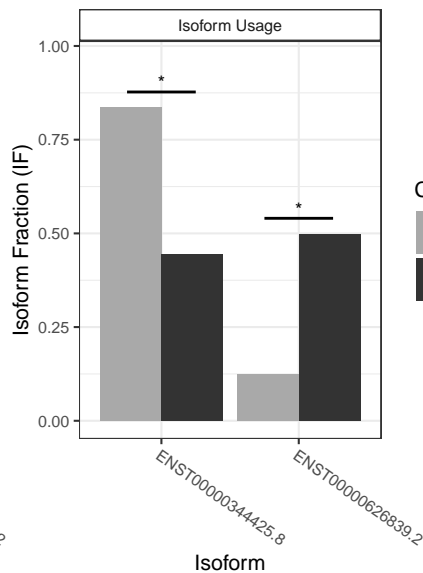
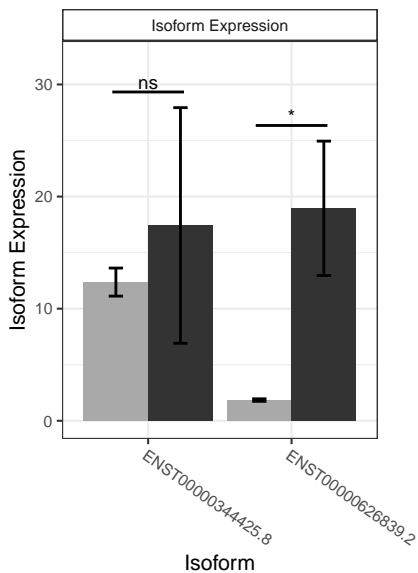
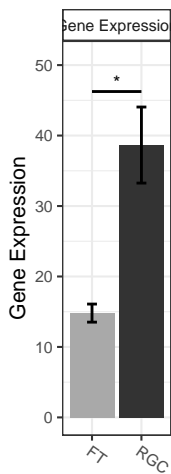
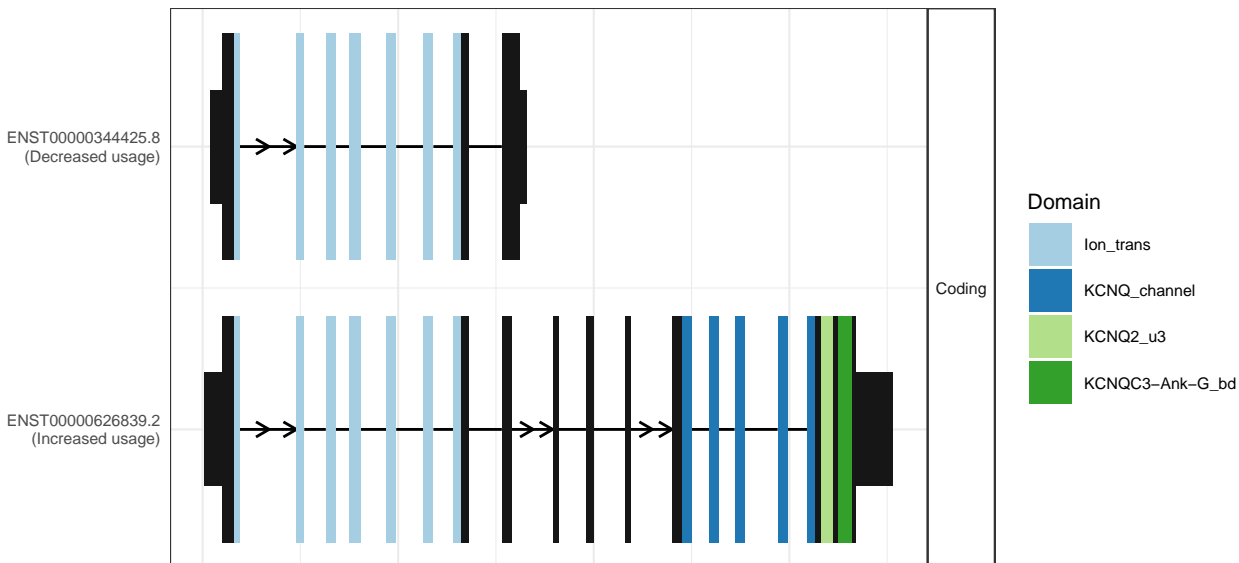


HELP







WD40 (x5)

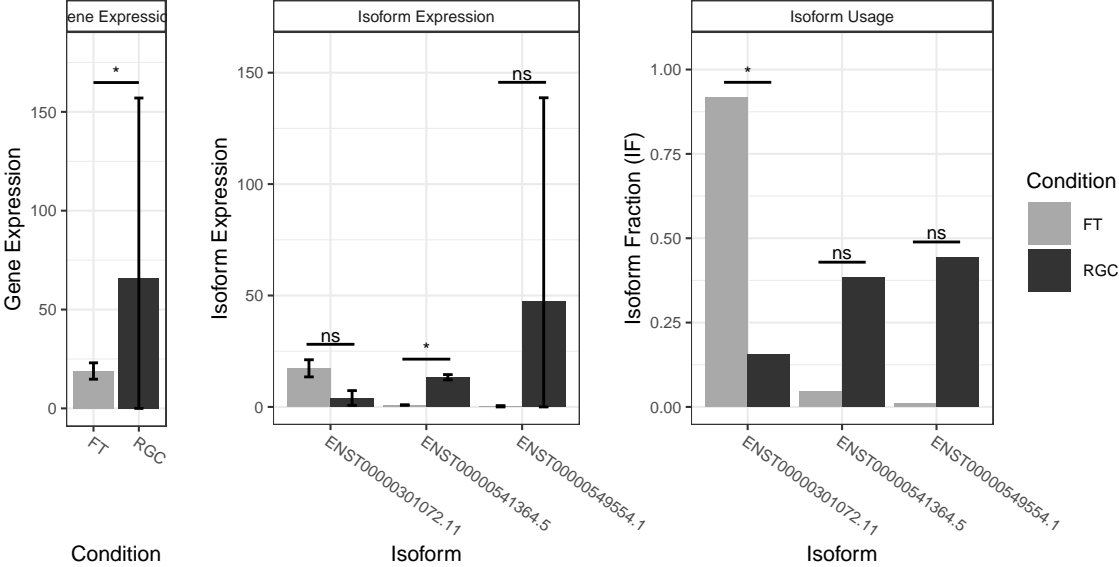
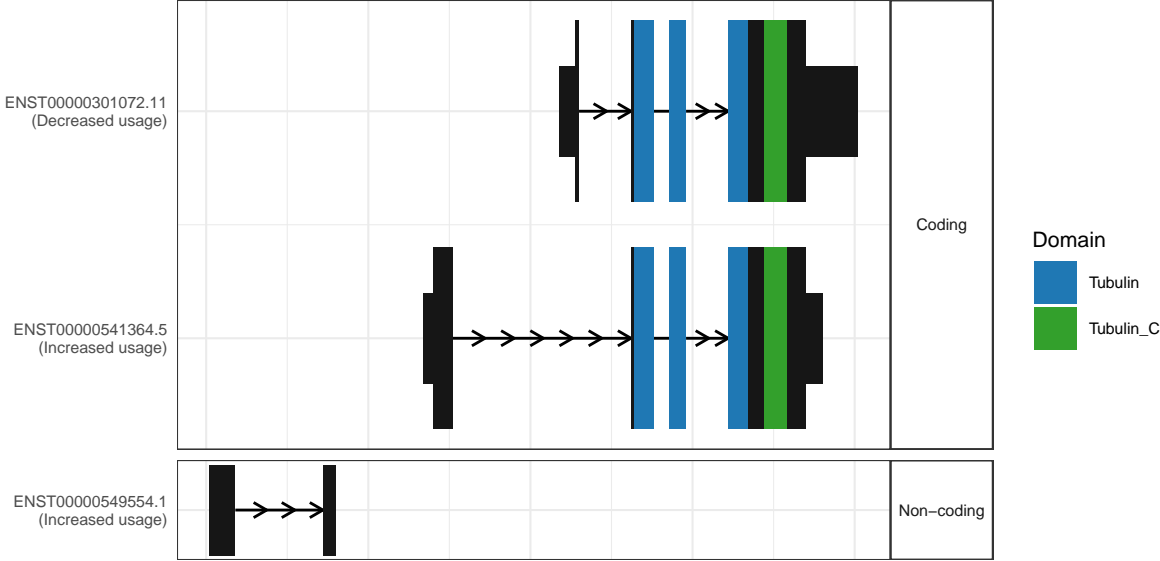
The isoform switch in KCNQ2 (FT vs RGC)



Domain

	Ion_trans
	KCNQ_channel
	KCNQ2_u3
	KCNQC3-Ank-G_bd

The isoform switch in TUBA1C (FT vs RGC)



Domain

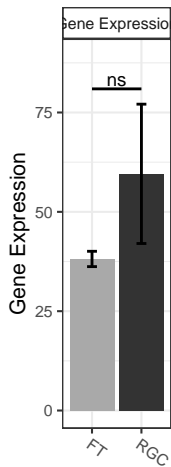
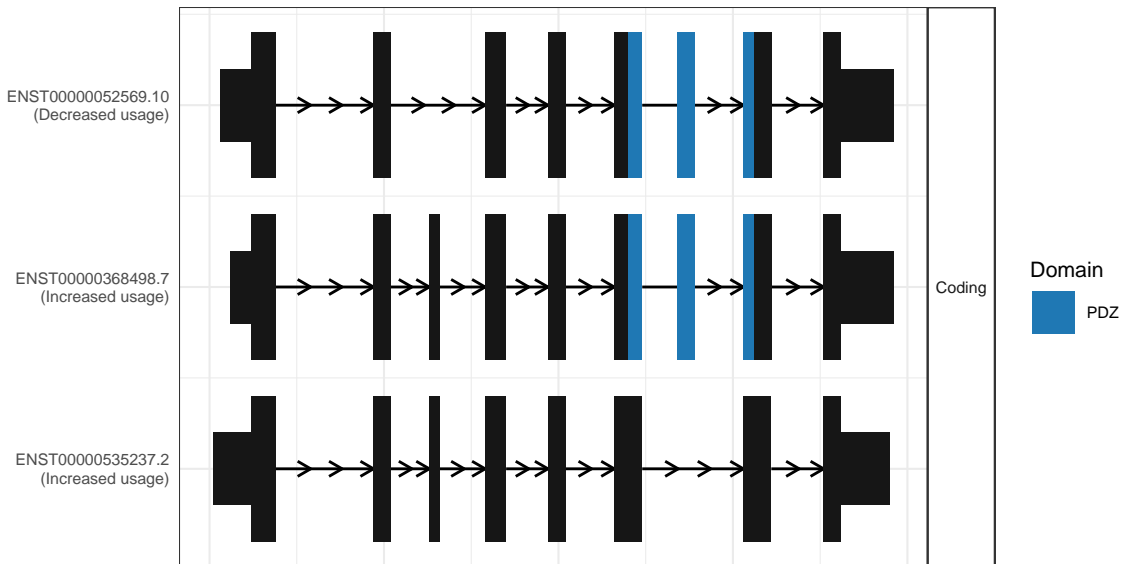


Tubulin

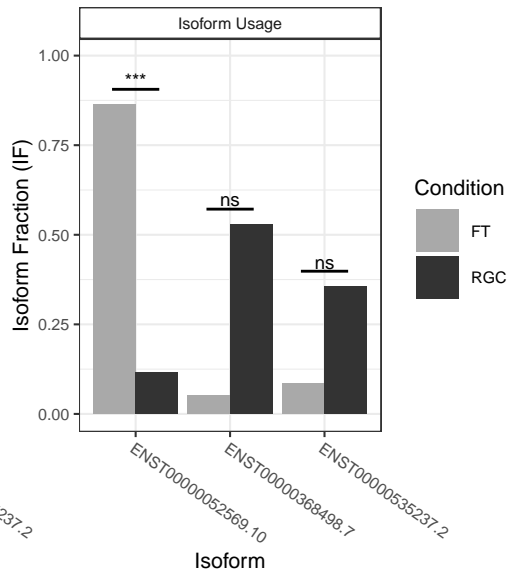
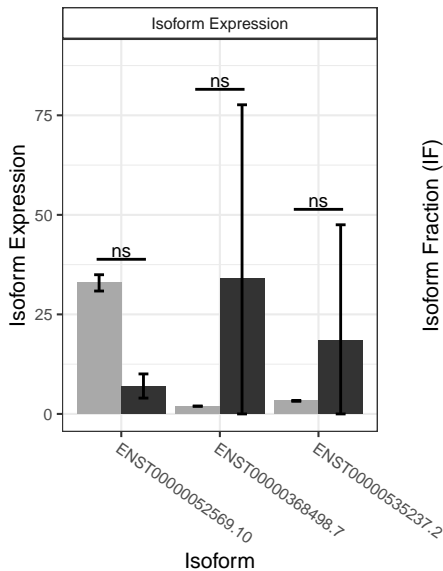


Tubulin_C

The isoform switch in GOPC (FT vs RGC)



Condition

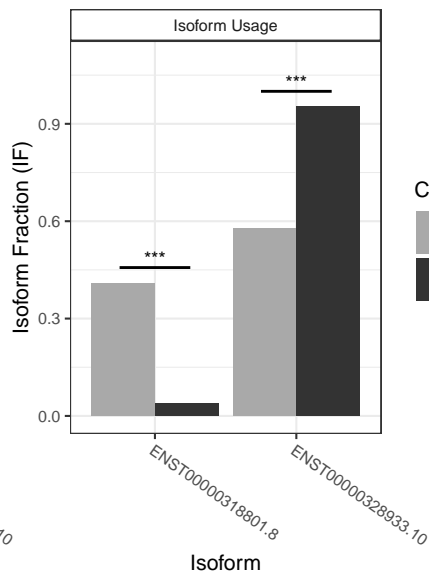
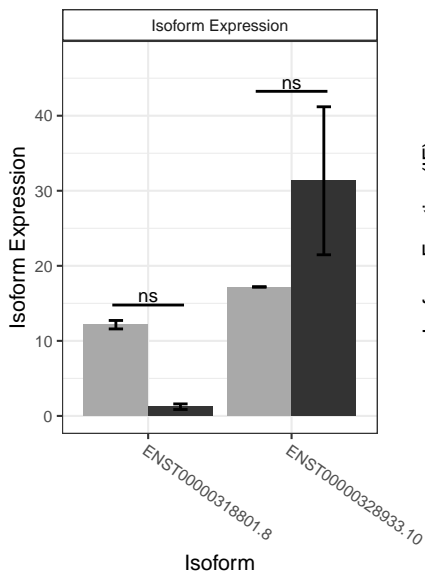
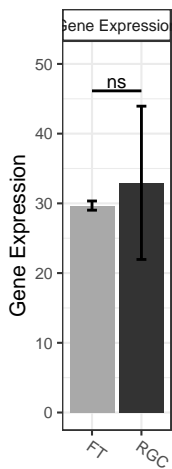
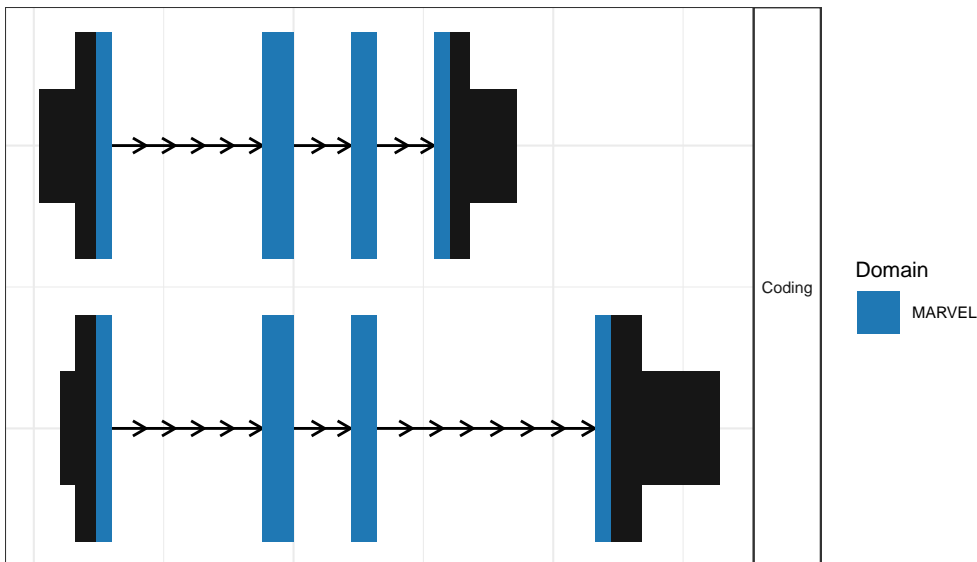


Domain



PDZ

The isoform switch in SYNGR1 (FT vs RGC)

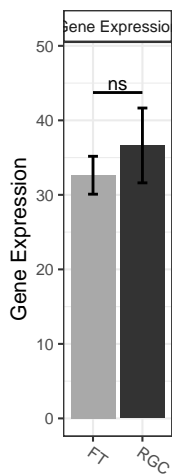
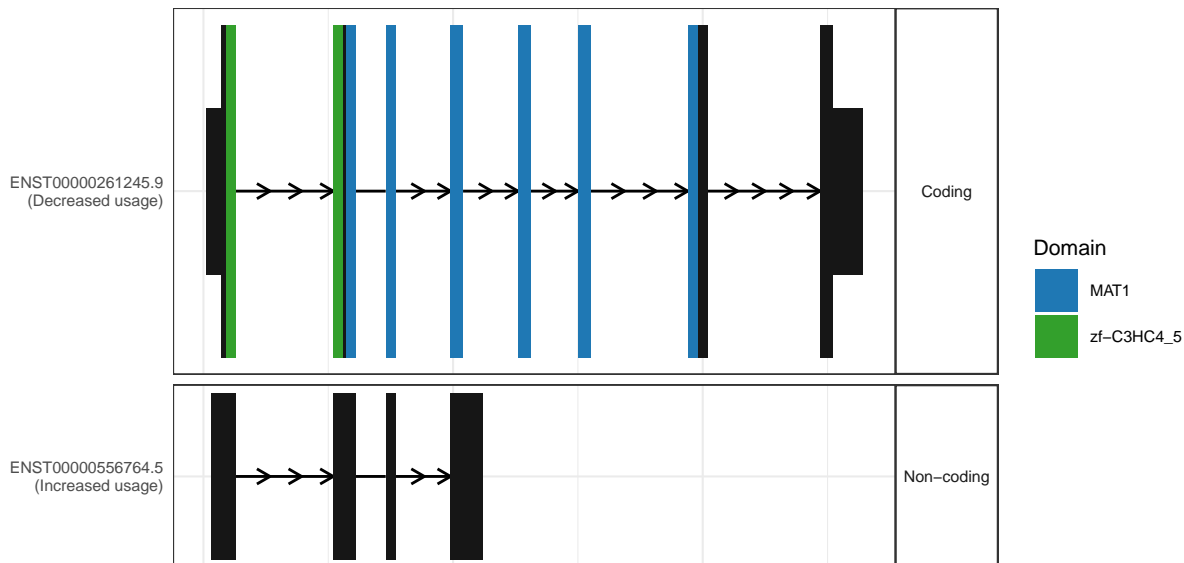


Domain

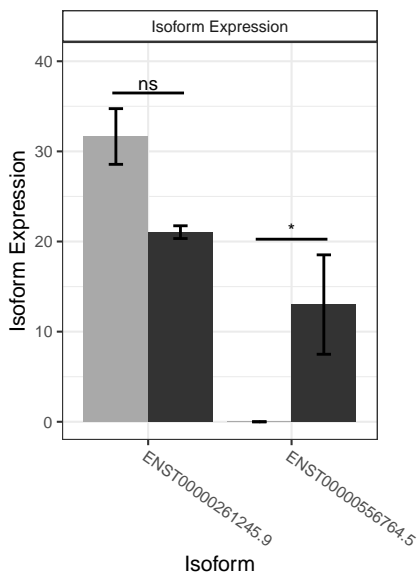


MARVEL

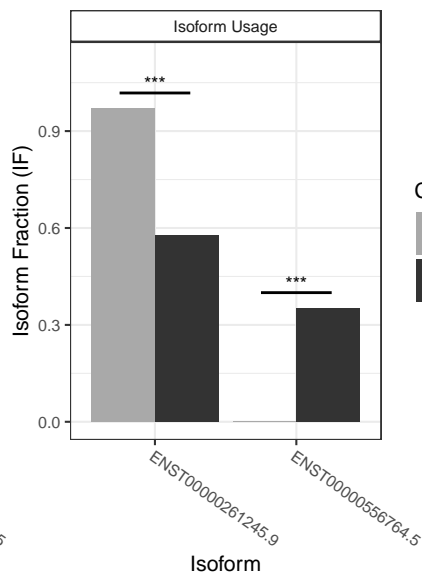
The isoform switch in MNAT1 (FT vs RGC)



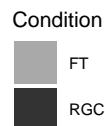
Condition



Isoform



Isoform



Domain

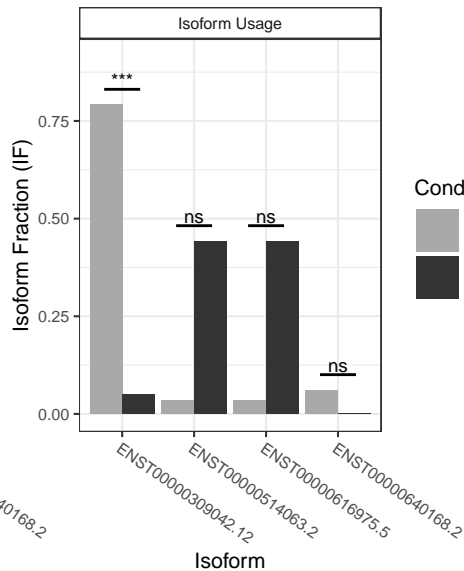
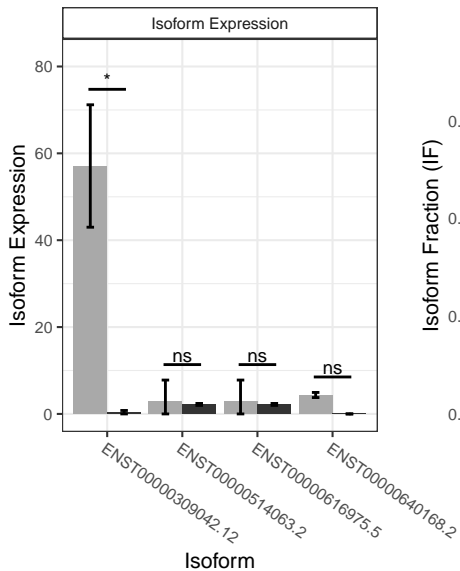
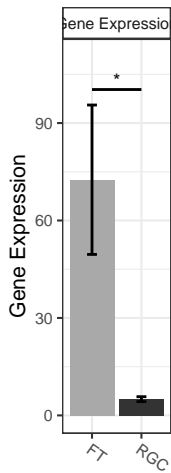
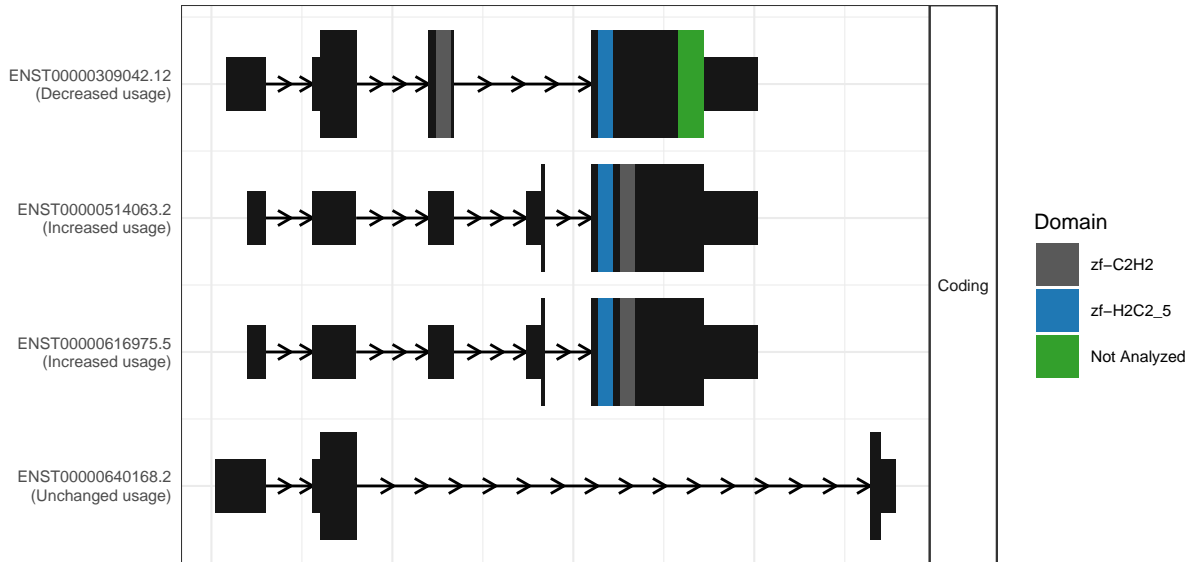


MAT1



zf-C3HC4_5

The isoform switch in REST (FT vs RGC)



Domain



zf-C2H2

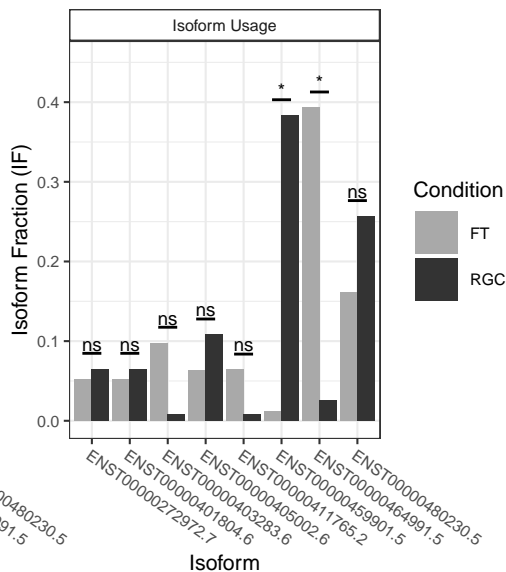
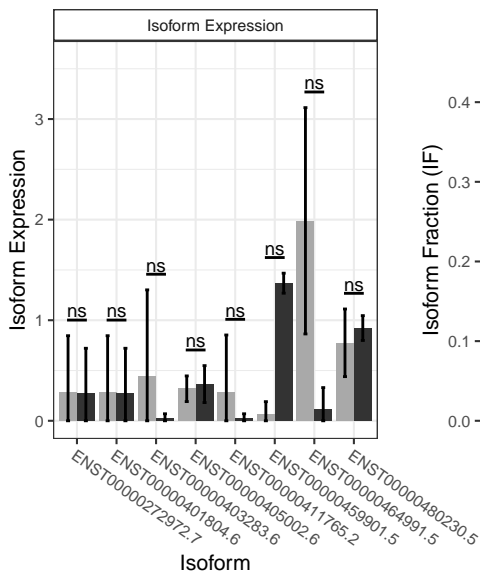
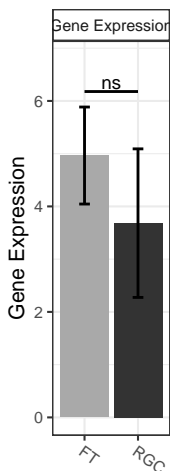
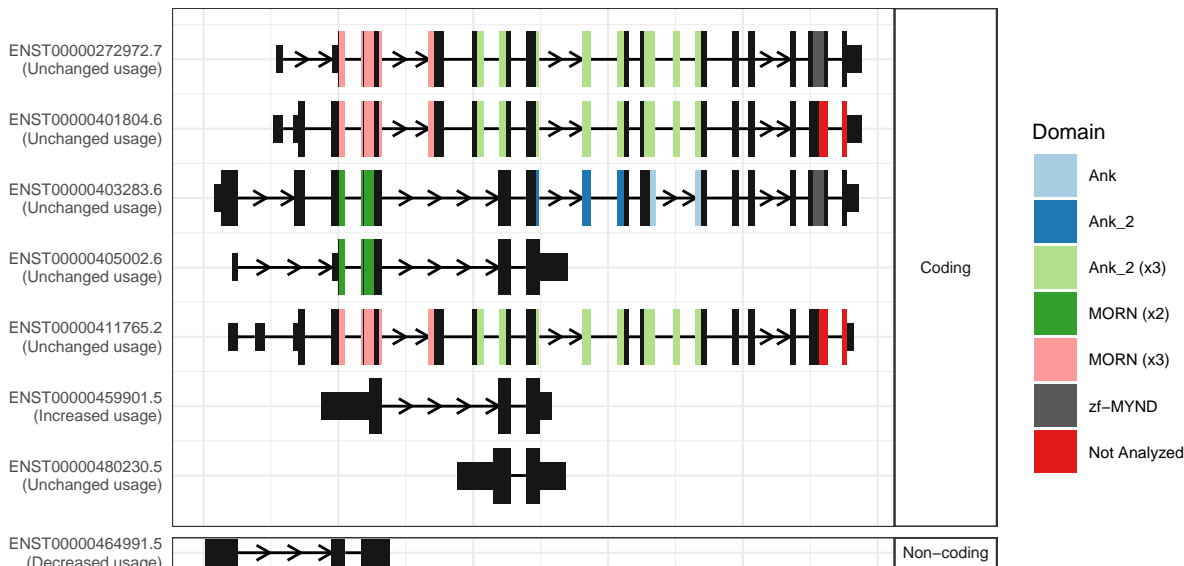


zf-H2C2_5



Not Analyzed

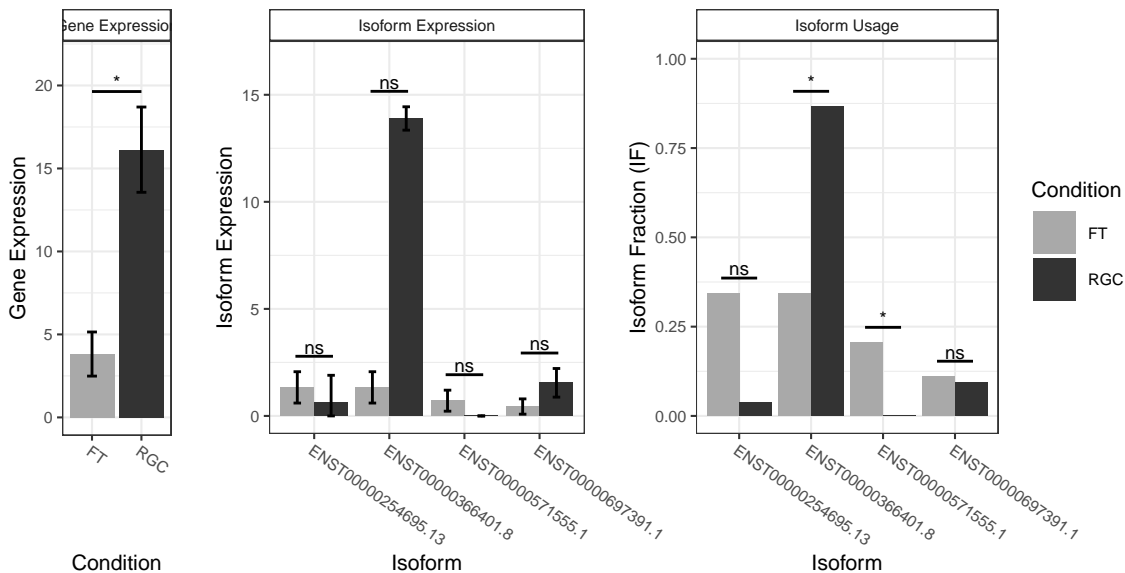
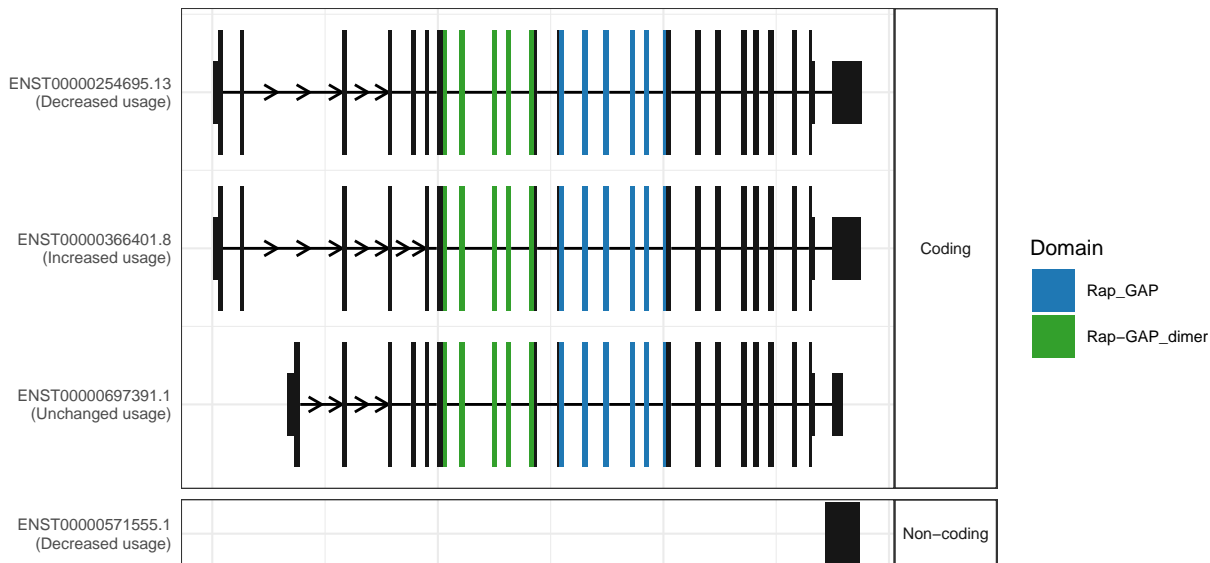
The isoform switch in ANKMYY1 (FT vs RGC)



Domain

Ank	Ank
Ank_2	Ank_2
Ank_2 (x3)	Ank_2 (x3)
MORN (x2)	MORN (x2)
MORN (x3)	MORN (x3)
zf-MYND	zf-MYND
Not Analyzed	Not Analyzed

The isoform switch in RAP1GAP2 (FT vs RGC)



Domain

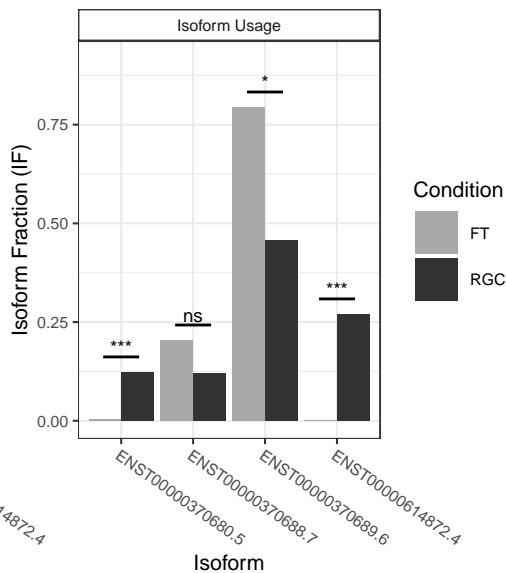
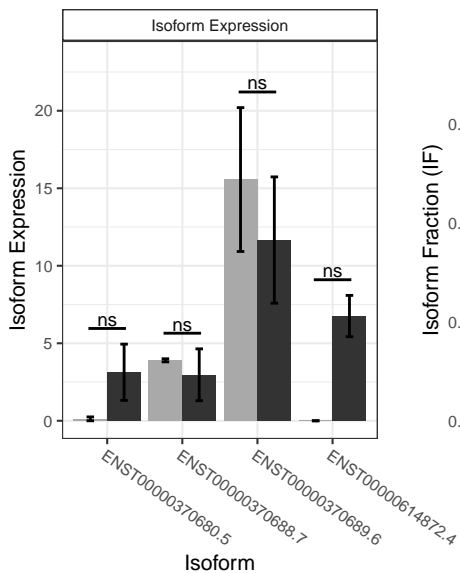
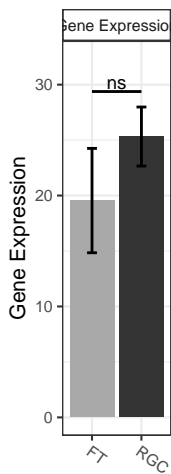
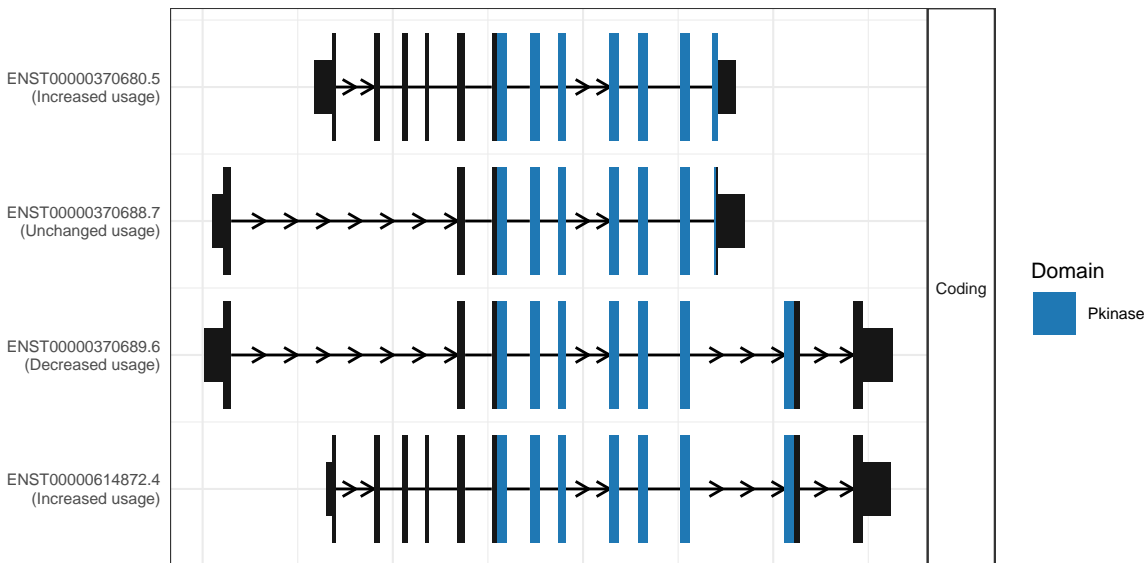


Rap_GAP



Rap-GAP_dimer

The isoform switch in PRKACB (FT vs RGC)

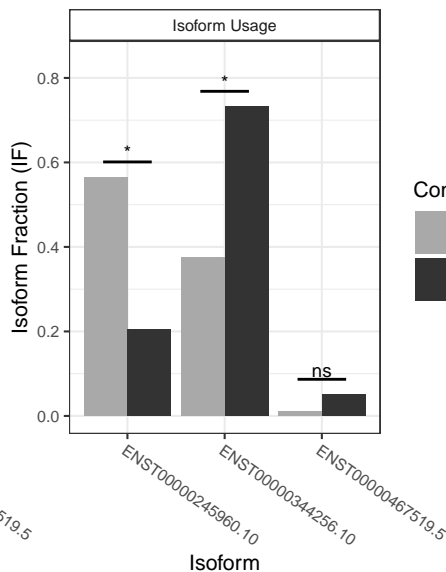
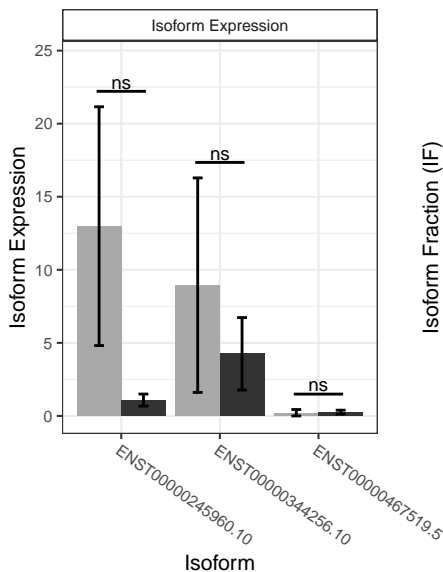
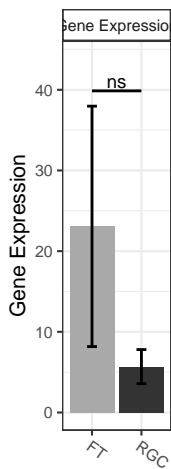
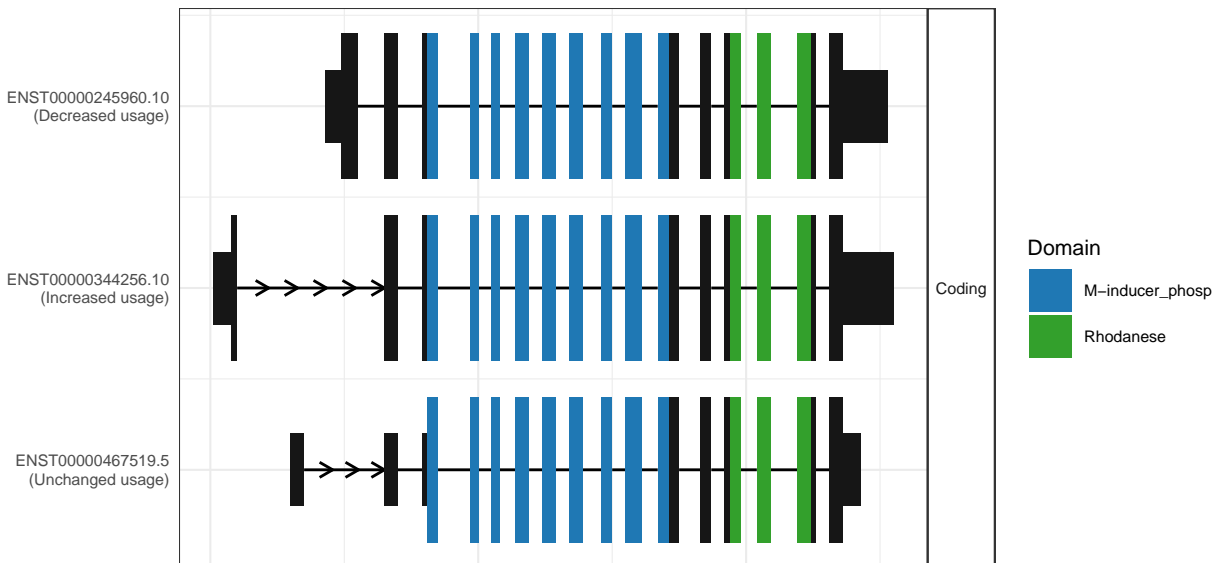


Domain



Pkinase

The isoform switch in CDC25B (FT vs RGC)



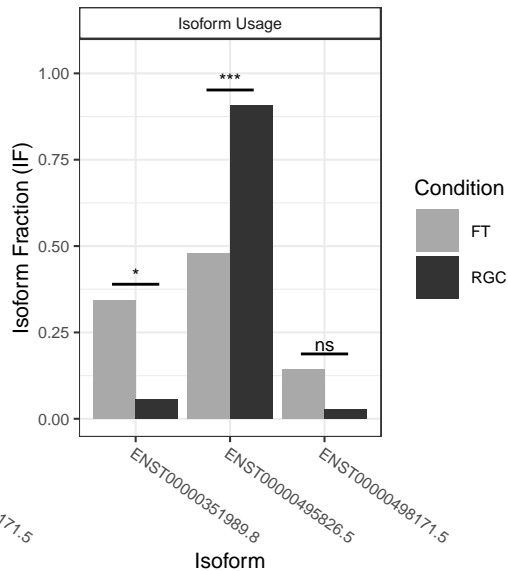
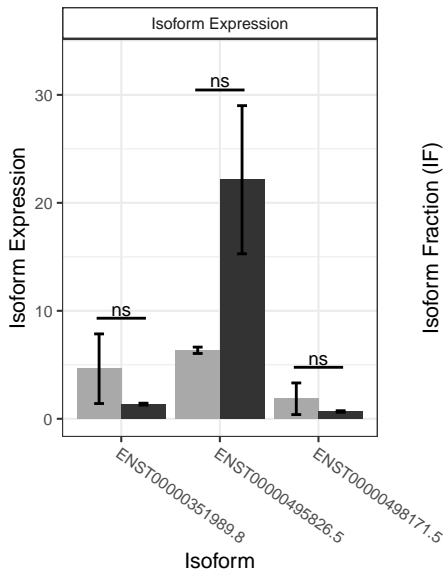
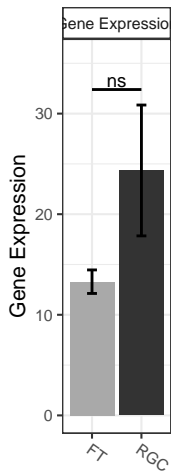
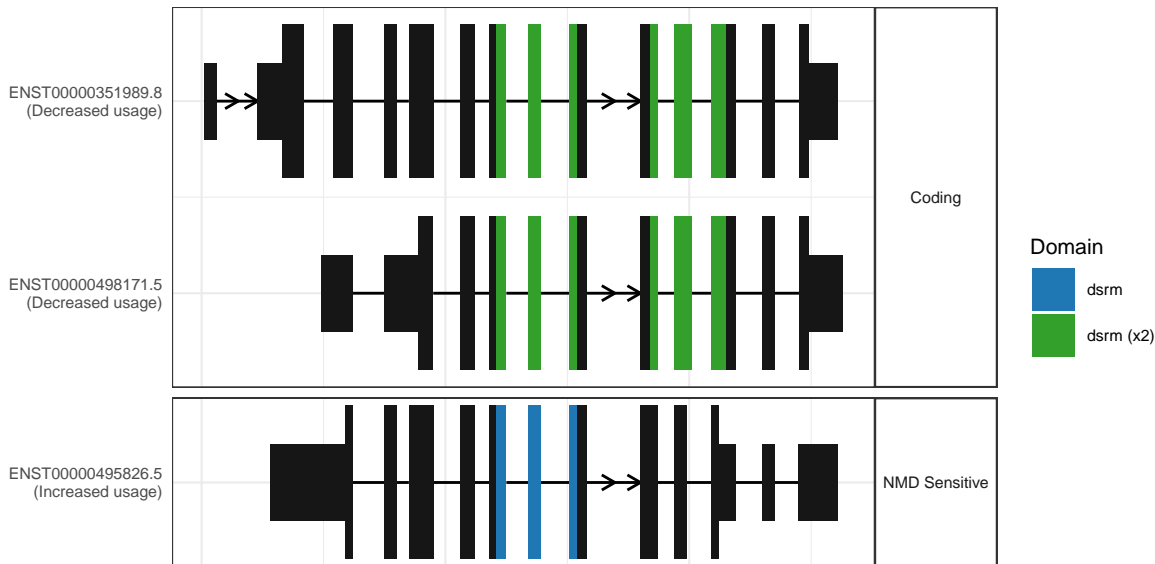
Domain



M-inducer_phosp

Rhodanese

The isoform switch in DGCR8 (FT vs RGC)



Domain

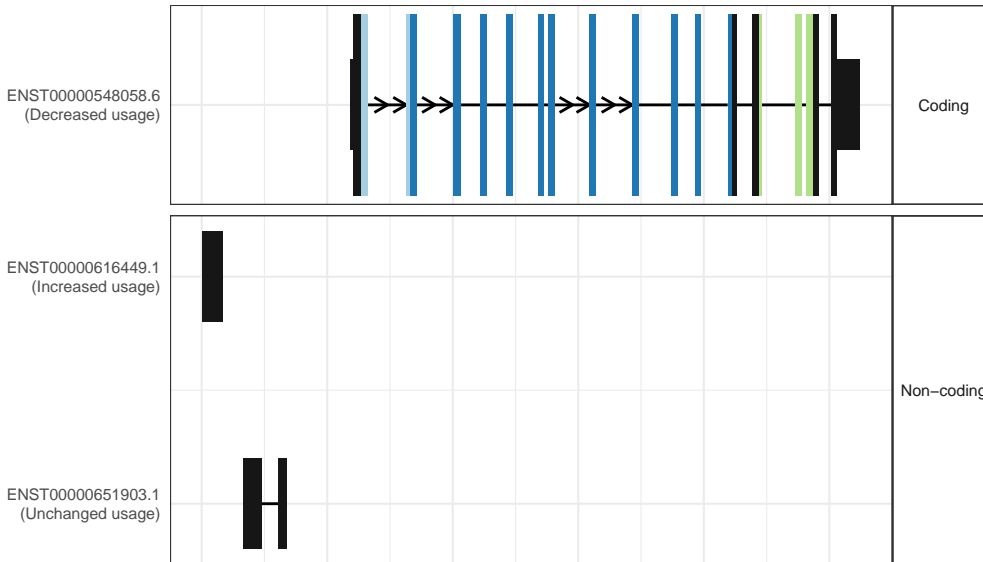


dsrm

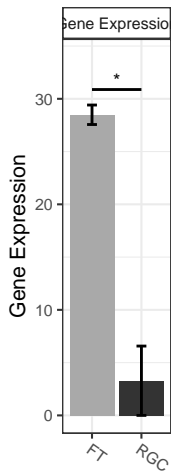
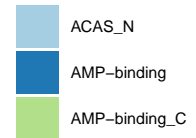


dsrm (x2)

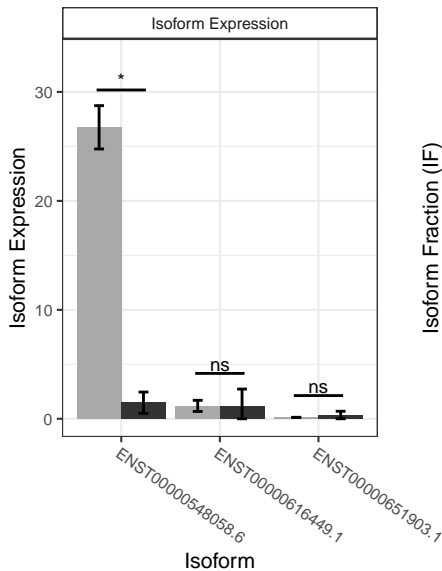
The isoform switch in ACSS3 (FT vs RGC)



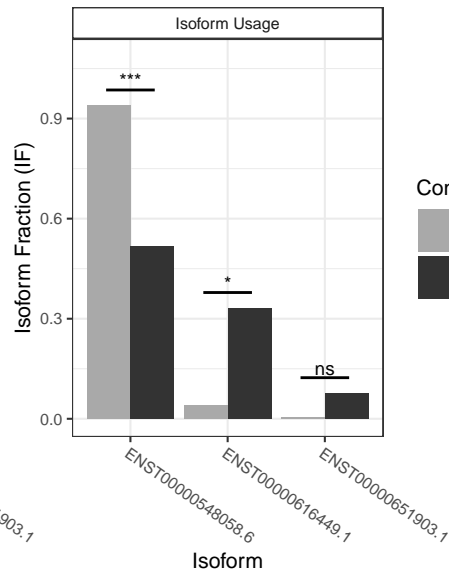
Domain



Condition



Isoform



Isoform

Condition



Domain



ACAS_N

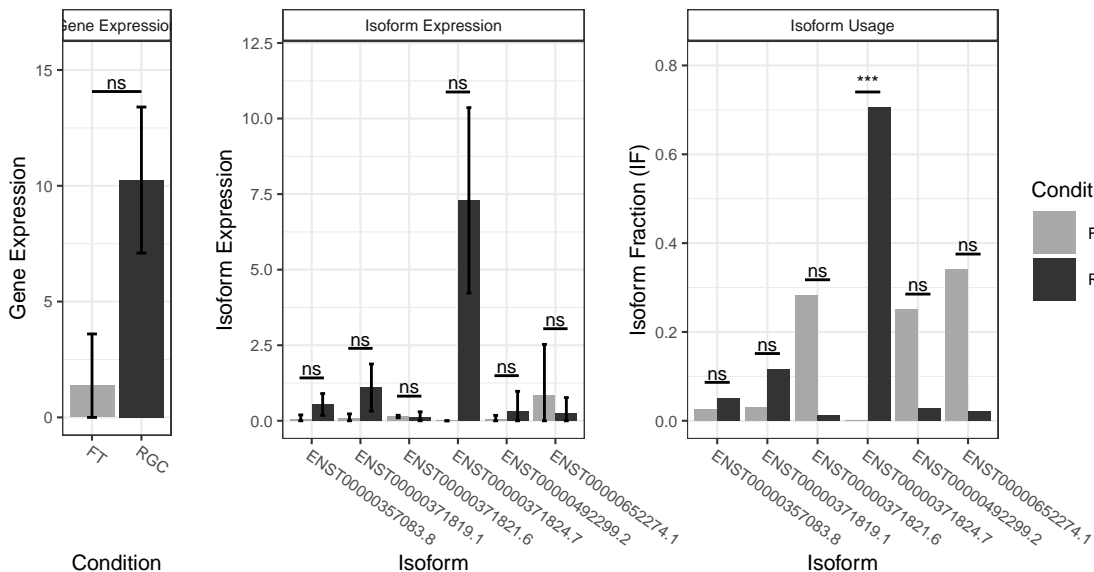
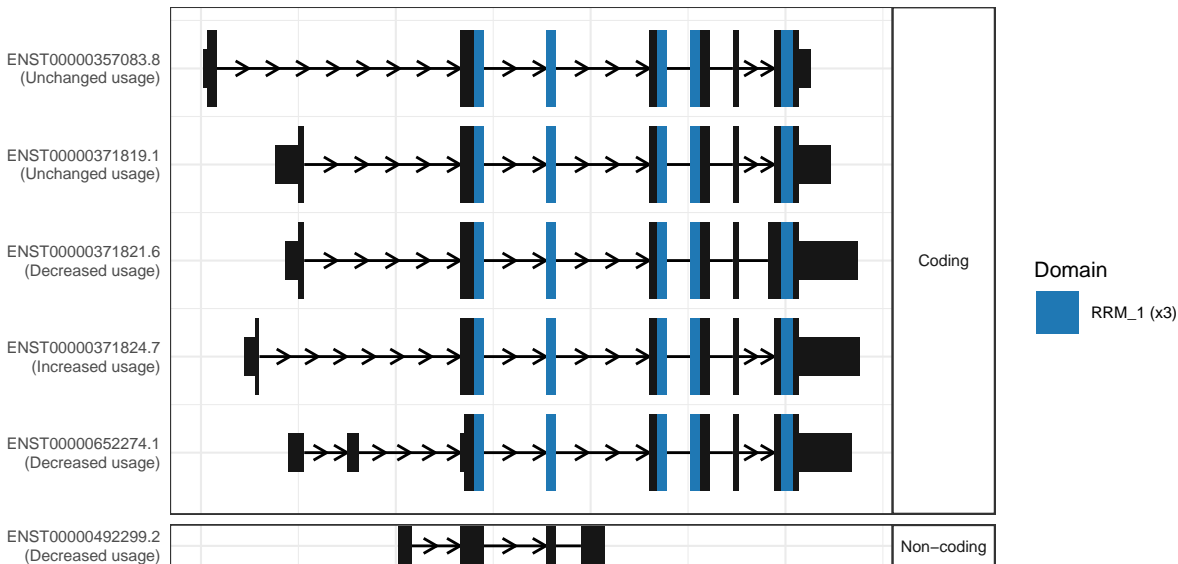


AMP-binding




AMP-binding_C

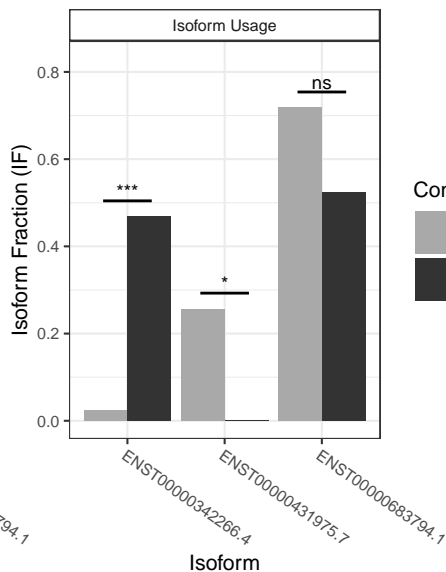
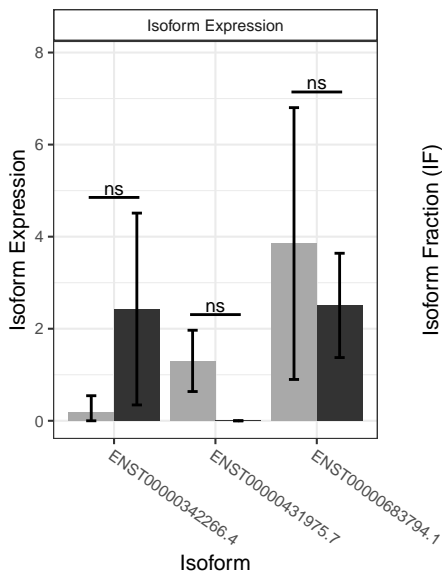
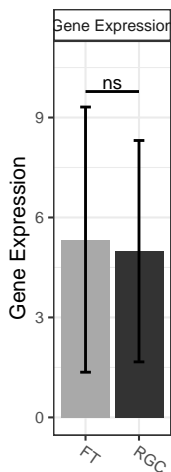
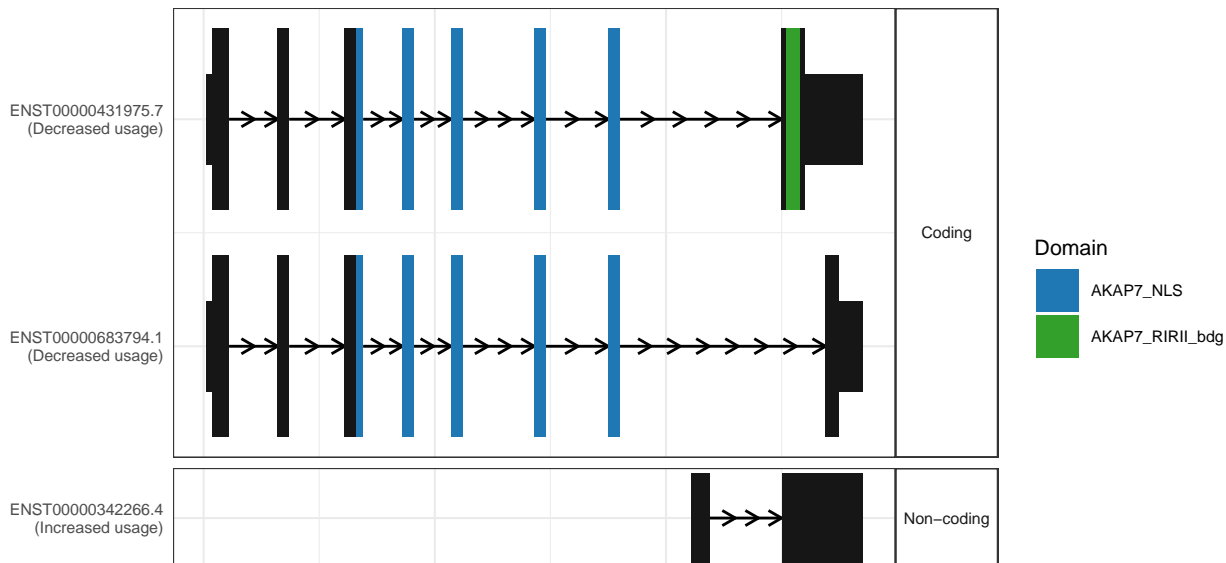
The isoform switch in ELAVL4 (FT vs RGC)



Domain

 RRM_1 (x3)

The isoform switch in AKAP7 (FT vs RGC)



Condition

Isoform

Isoform

Condition



Domain

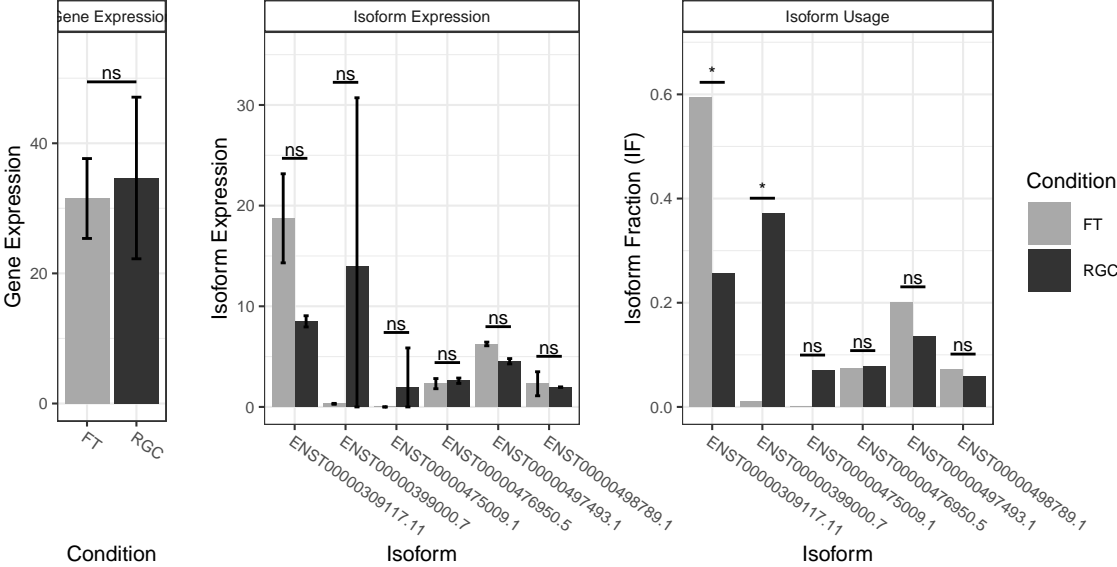
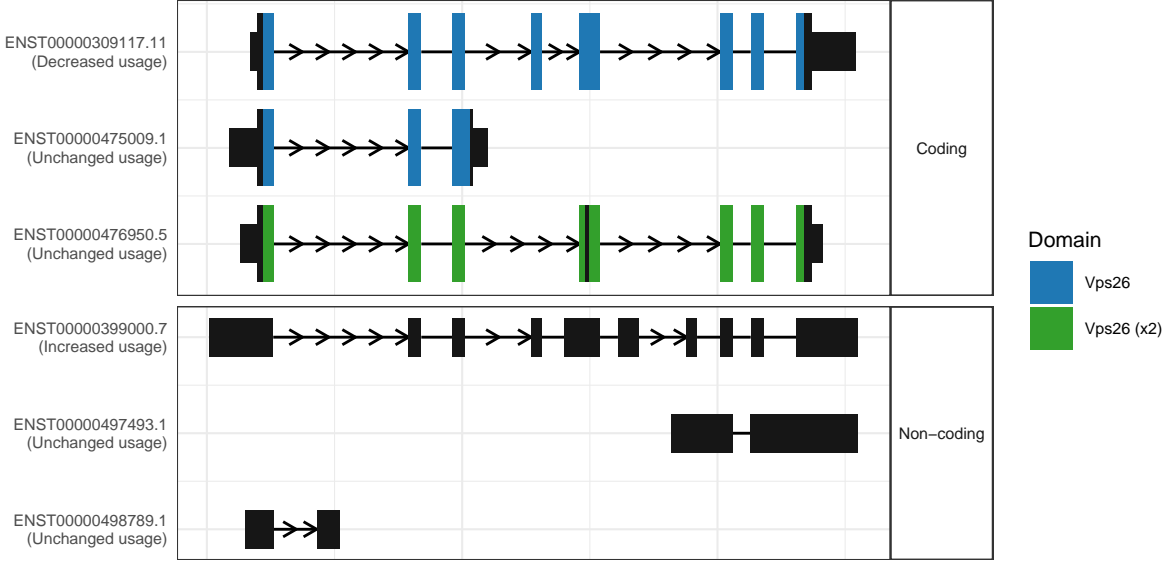


AKAP7_NLS



AKAP7_RIRII_bdg

The isoform switch in VPS26C (FT vs RGC)



Domain

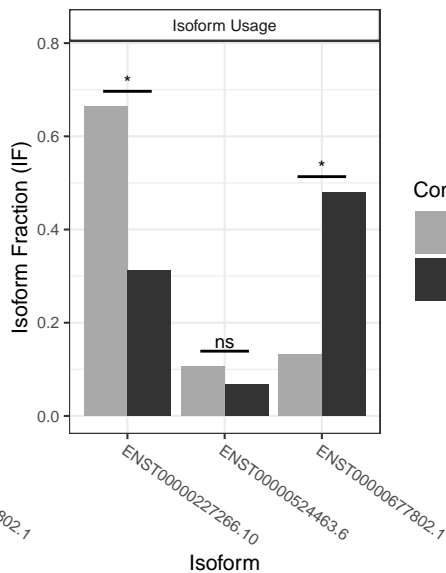
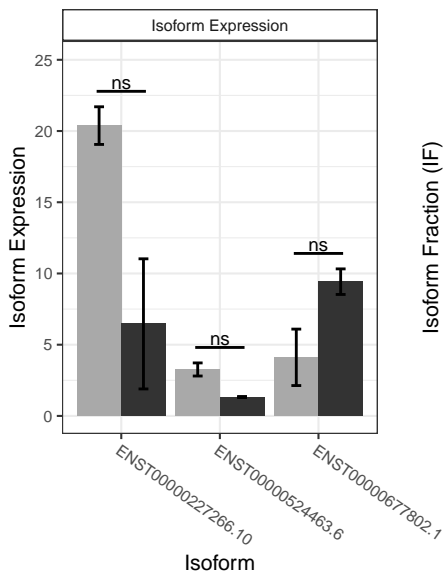
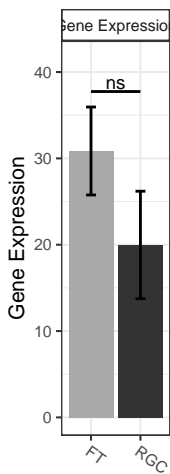
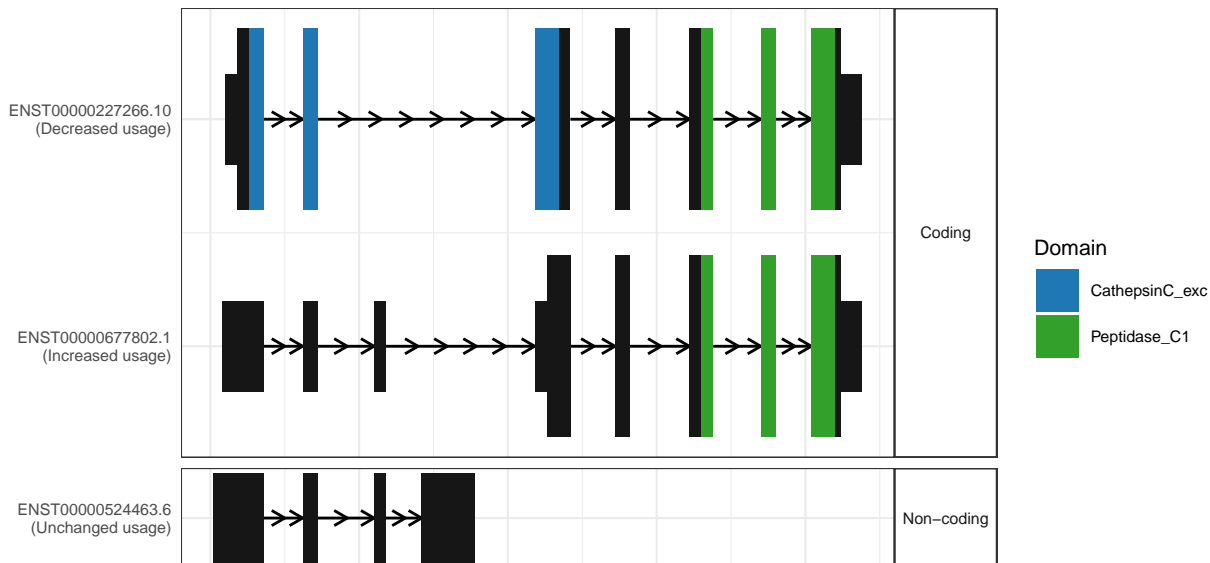


Vps26



Vps26 (x2)

The isoform switch in CTSC (FT vs RGC)



Domain

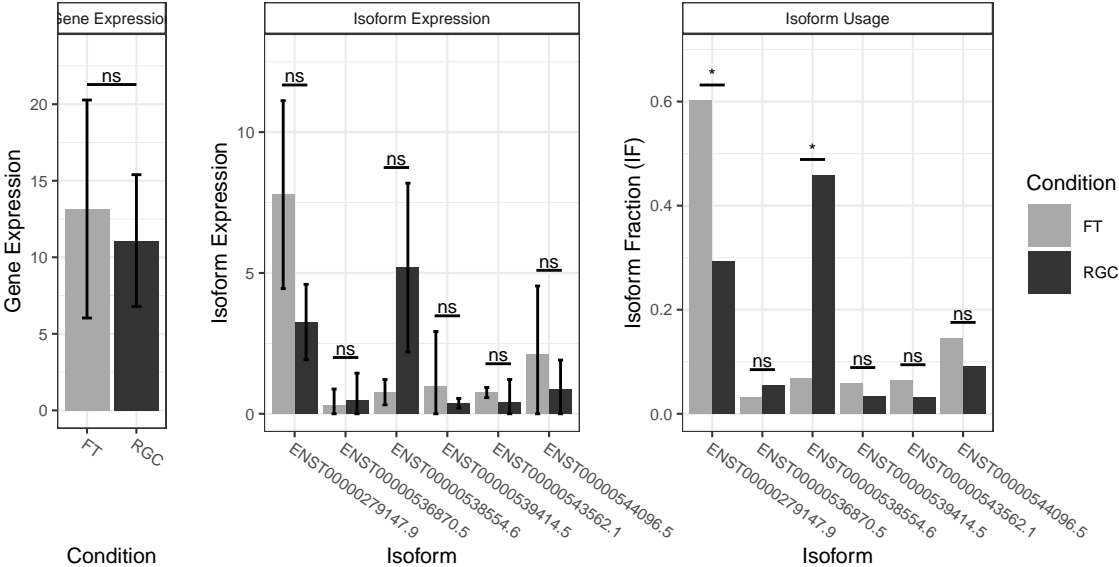
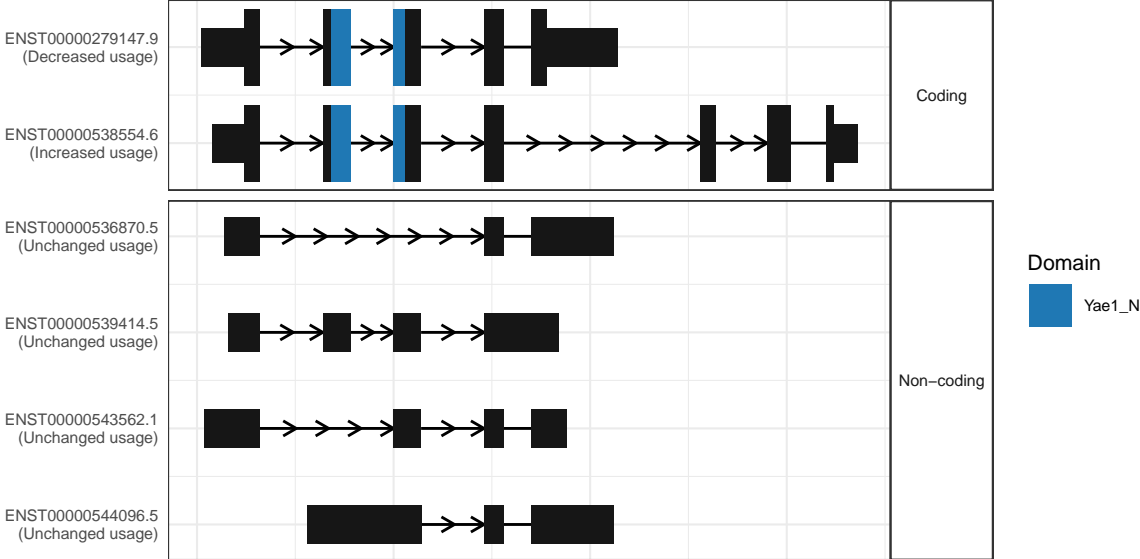


CathepsinC_exc



Peptidase_C1

The isoform switch in LTO1 (FT vs RGC)

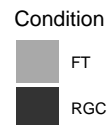
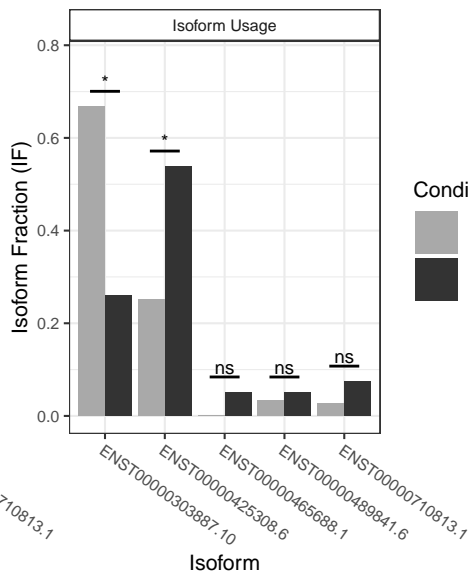
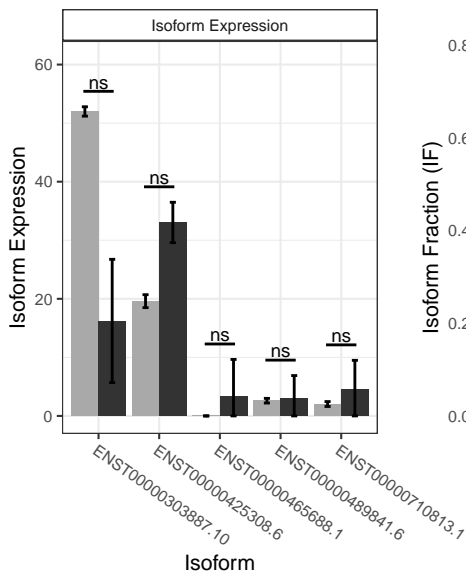
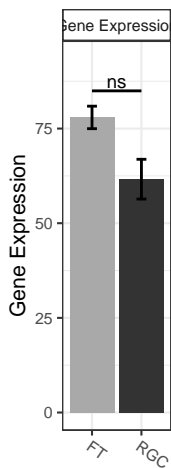
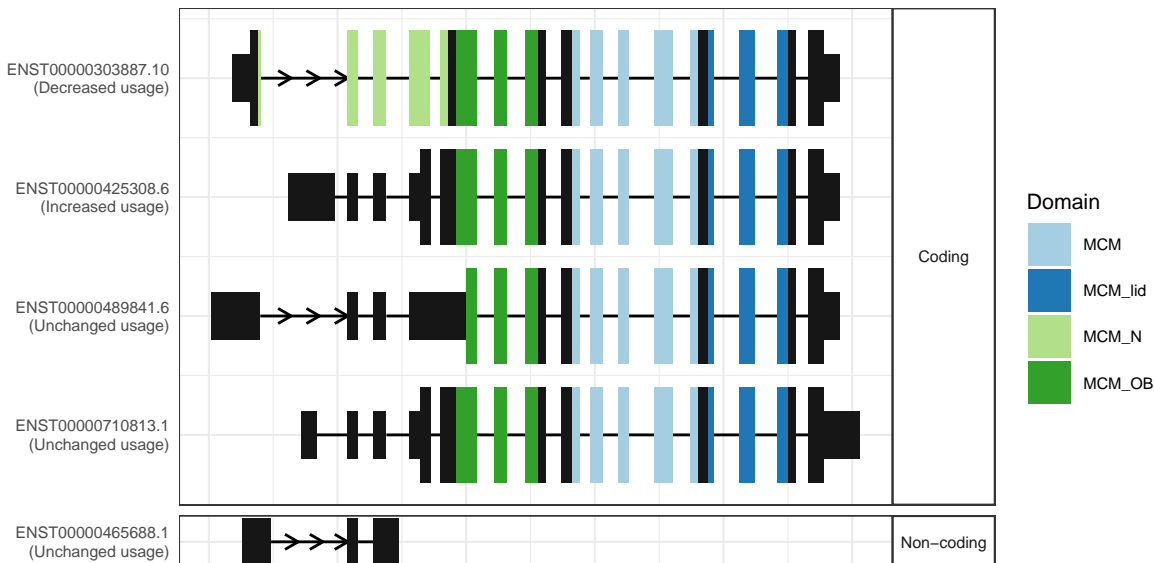


Domain



Yae1_N

The isoform switch in MCM7 (FT vs RGC)



Domain



MCM



MCM_lid

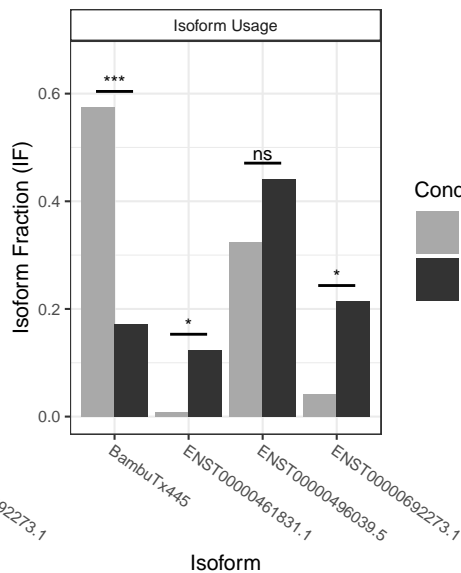
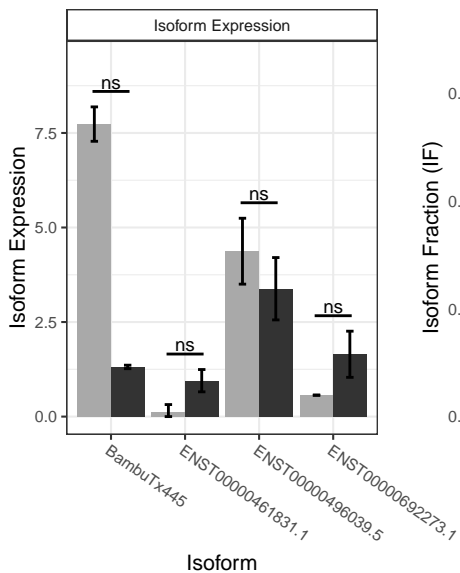
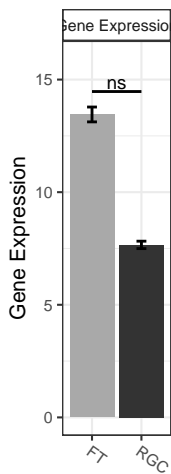
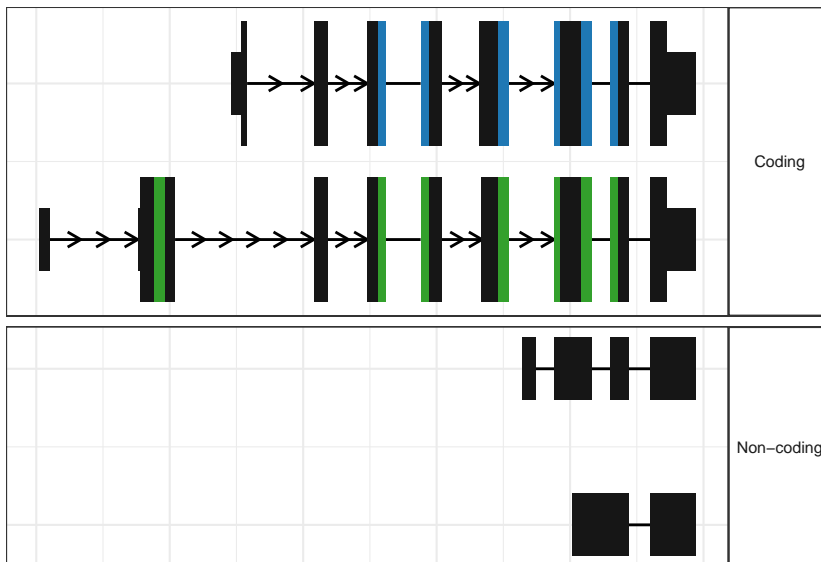


MCM_N



MCM_OB

The isoform switch in ZNF385C (FT vs RGC)



Domain

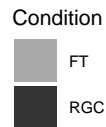
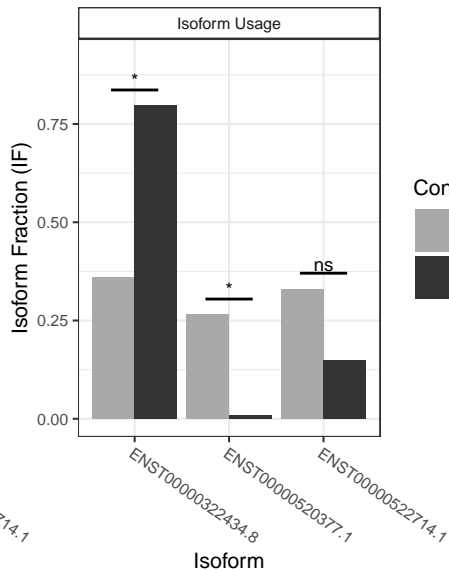
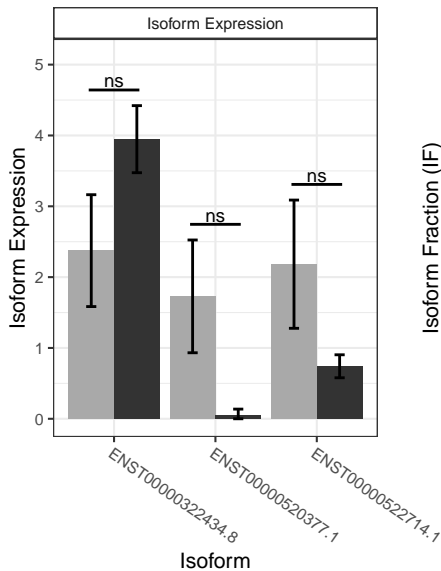
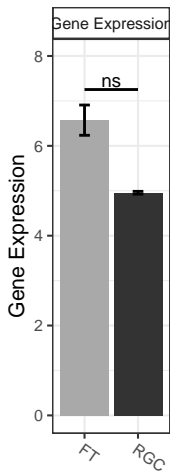
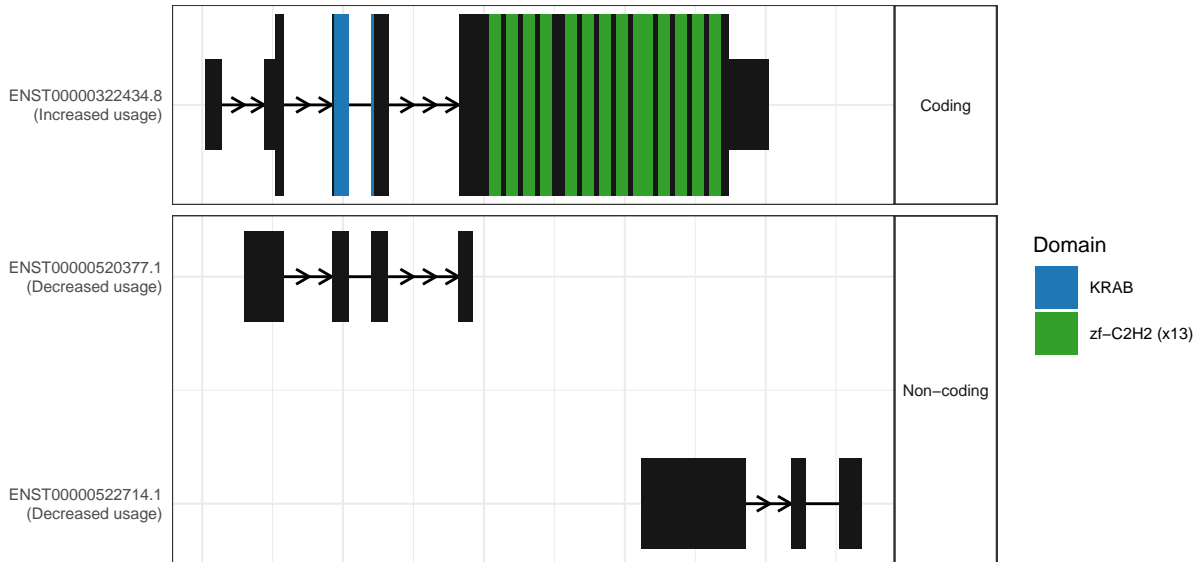


zf_met ($x3$)



zf_met ($x4$)

The isoform switch in ZNF354B (FT vs RGC)



Domain

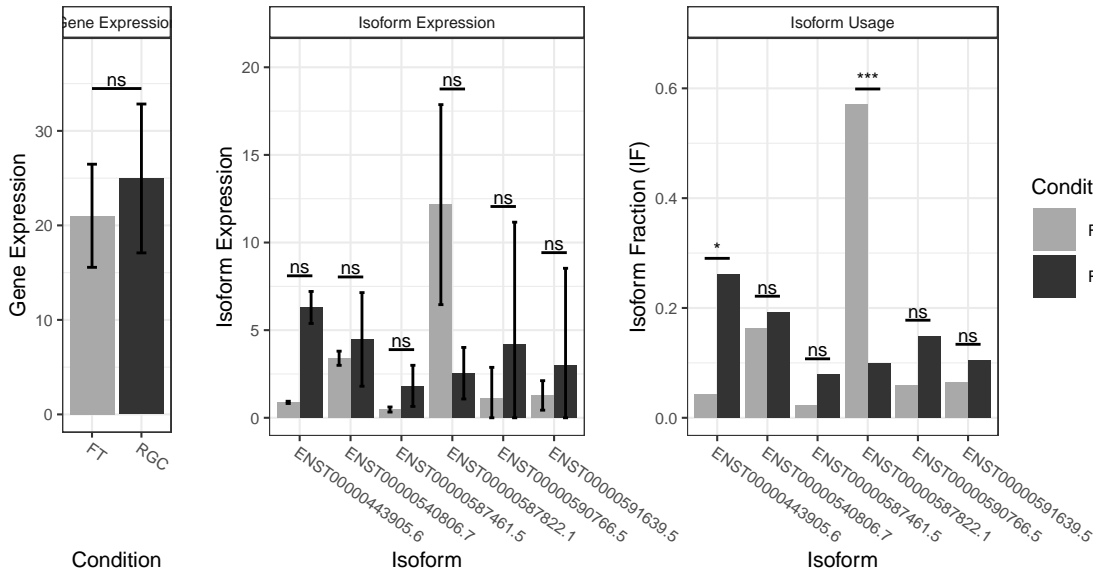
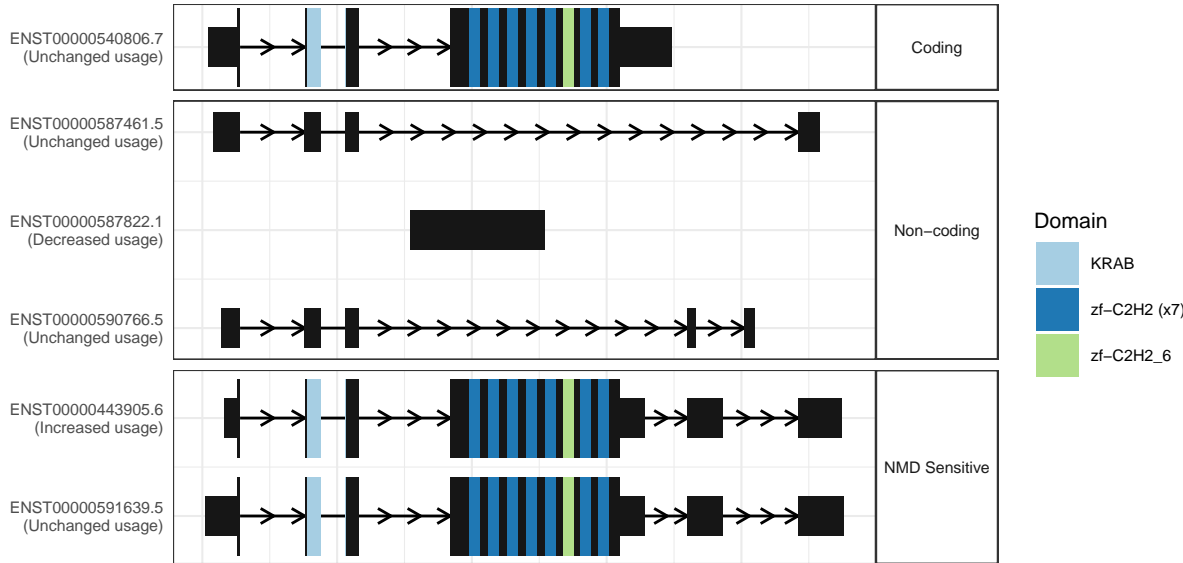


KRAB






zf-C2H2 (x13)

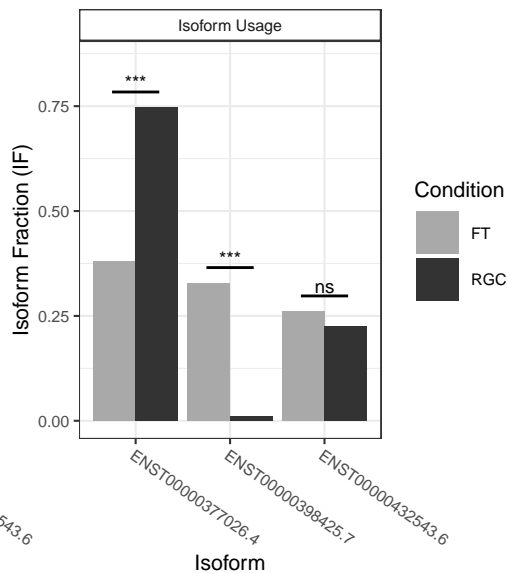
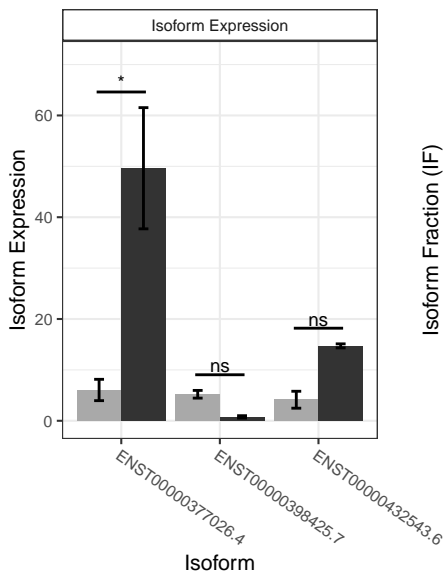
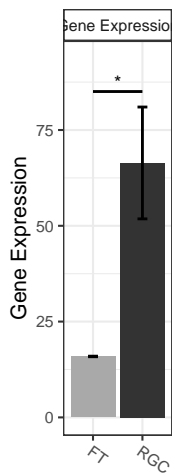
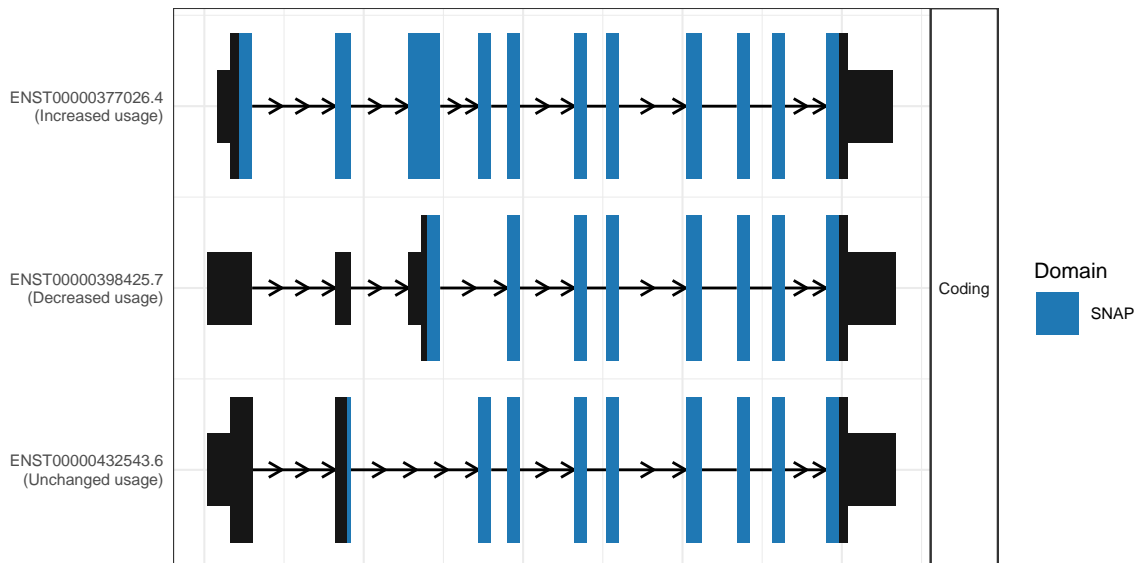
The isoform switch in ZNF506 (FT vs RGC)



Domain

	KRAB
	zf-C2H2 (x7)
	zf-C2H2_6

The isoform switch in NAPB (FT vs RGC)

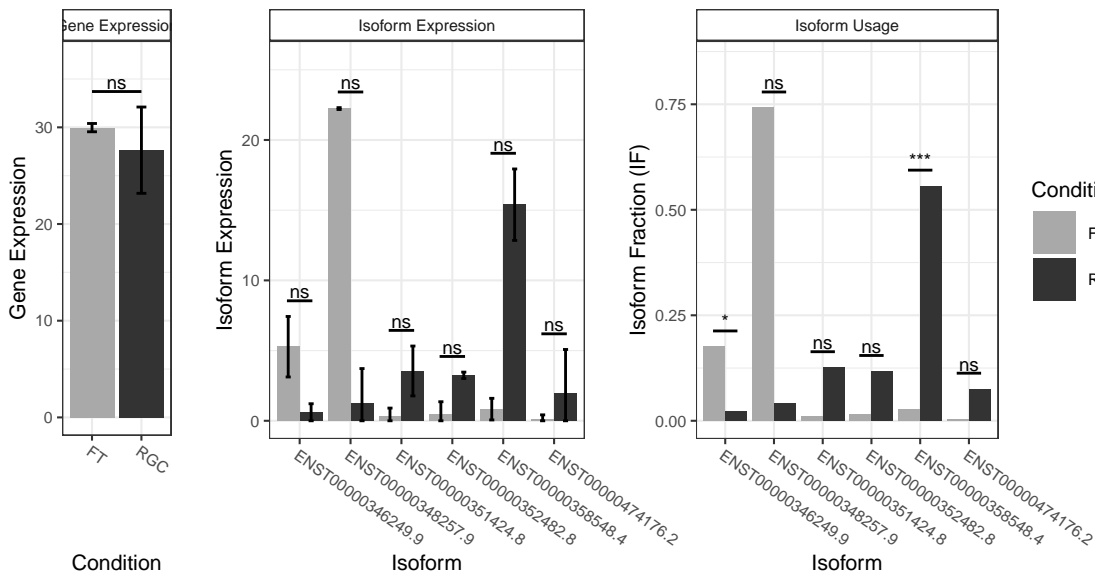
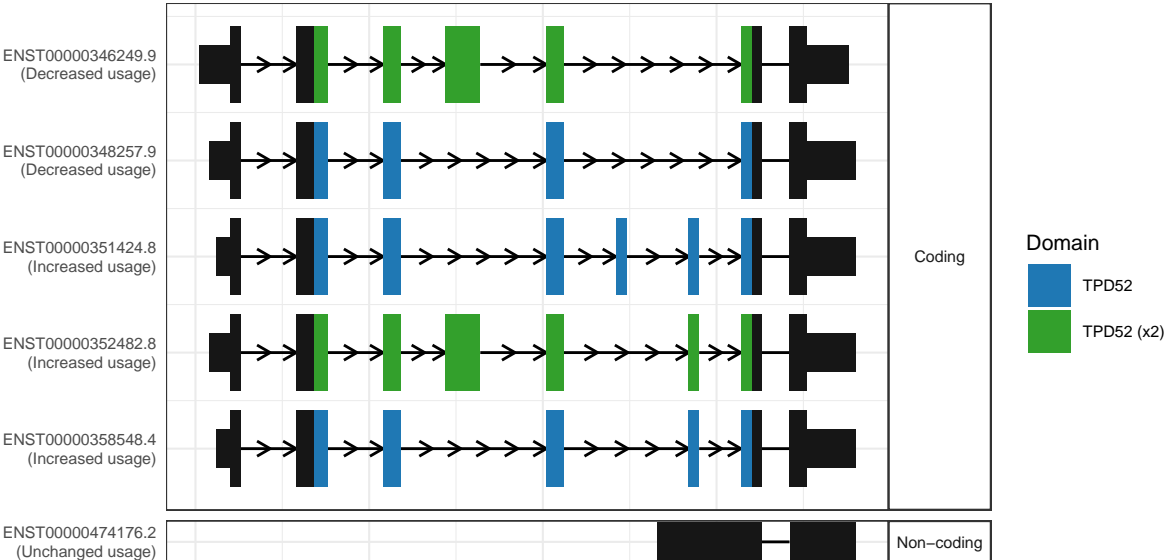


Domain



SNAP

The isoform switch in TPD52L2 (FT vs RGC)



Domain

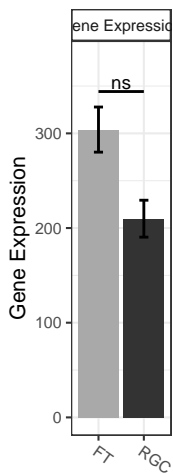
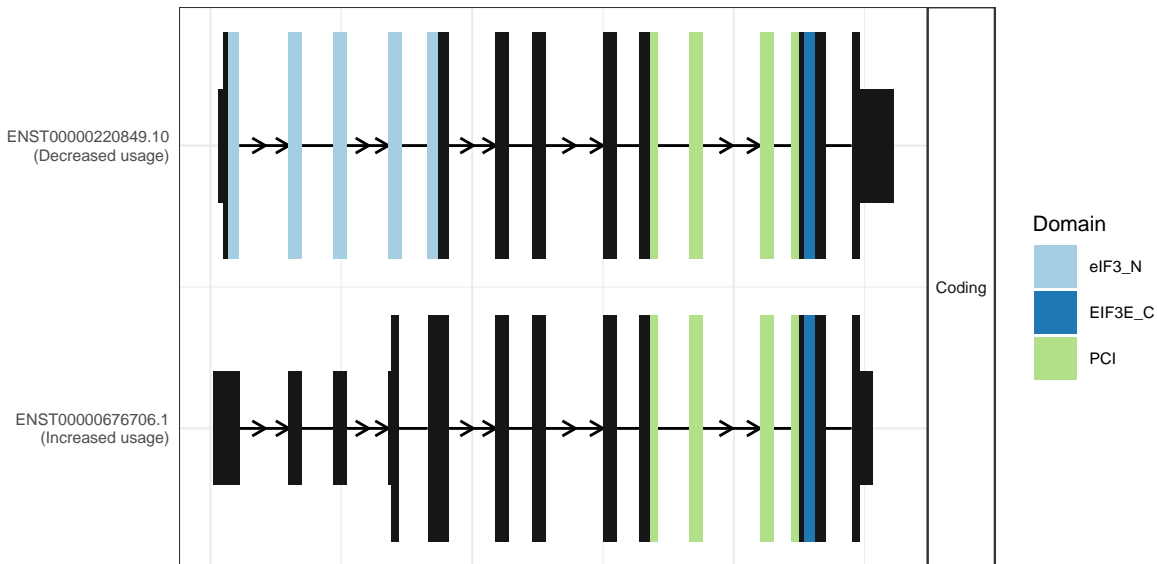


TPD52

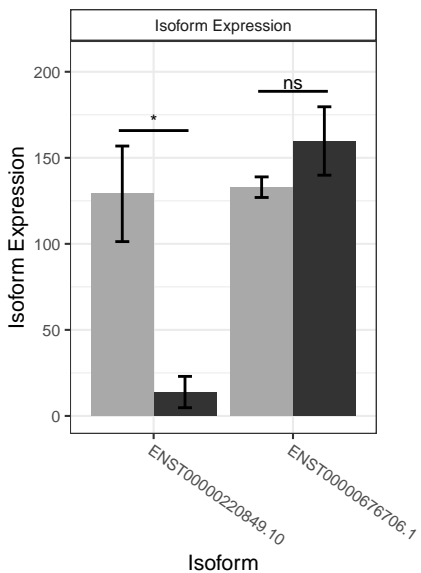


TPD52 (x2)

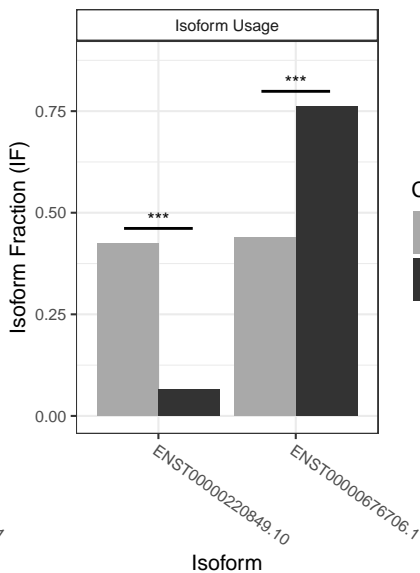
The isoform switch in EIF3E (FT vs RGC)



Condition



Isoform






Isoform

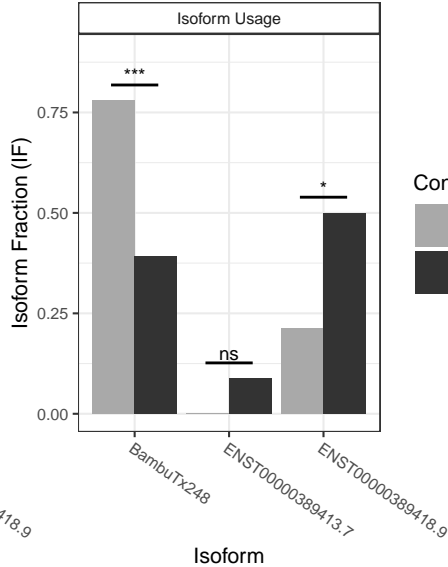
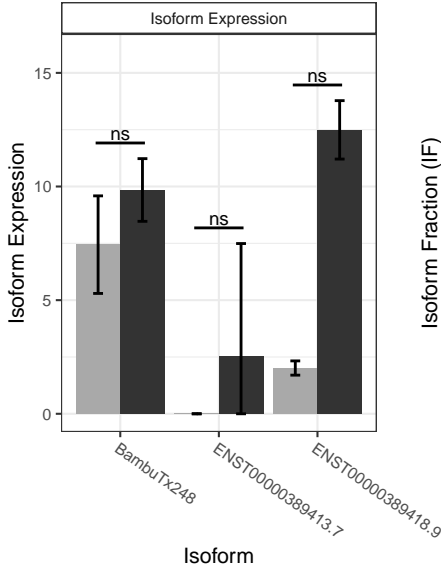
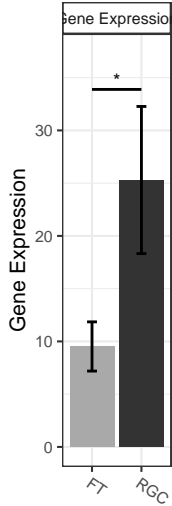
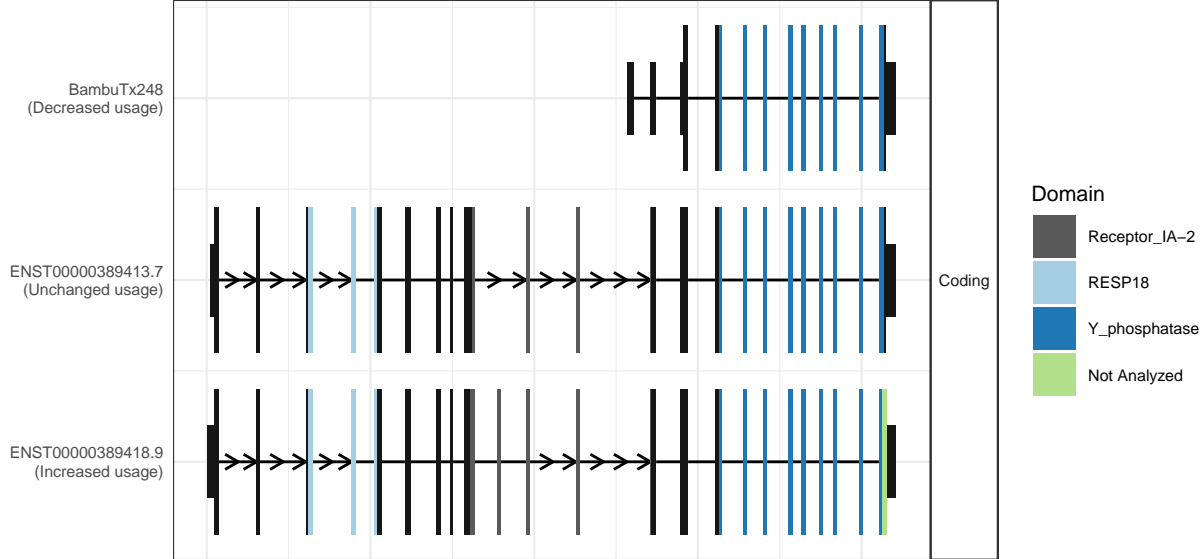
Condition



Domain

	eIF3_N
	EIF3E_C
	PCI

The isoform switch in PTPRN2 (FT vs RGC)



Domain



Receptor_IA-2



RESP18

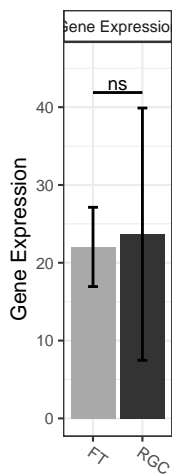
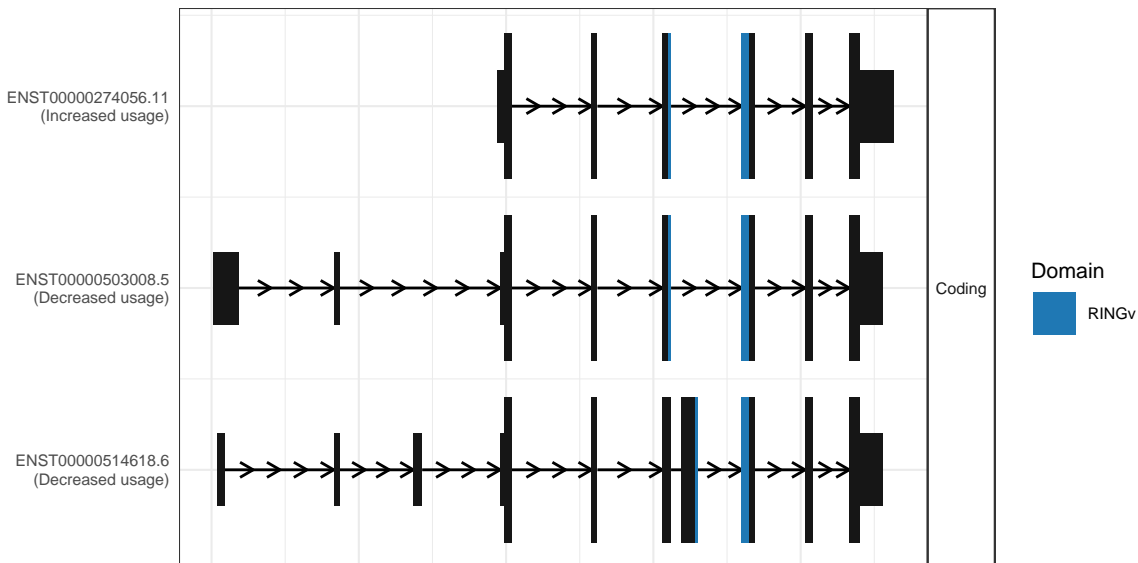


Y_phosphatase

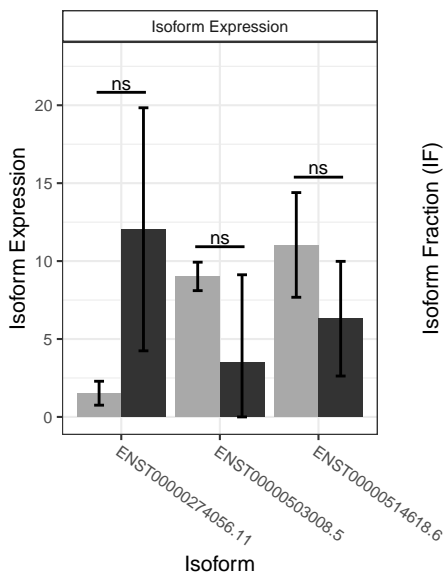


Not Analyzed

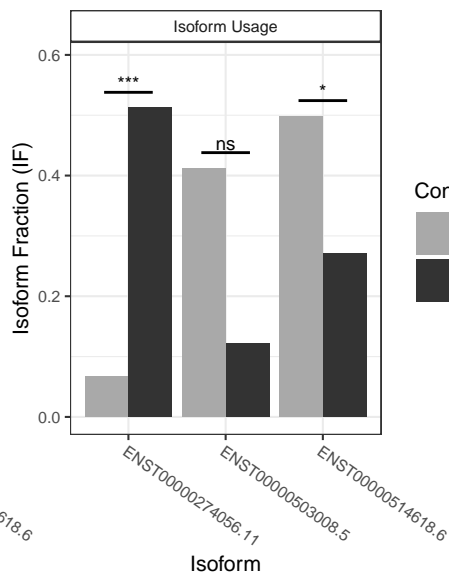
The isoform switch in MARCHF1 (FT vs RGC)



Condition



Isoform



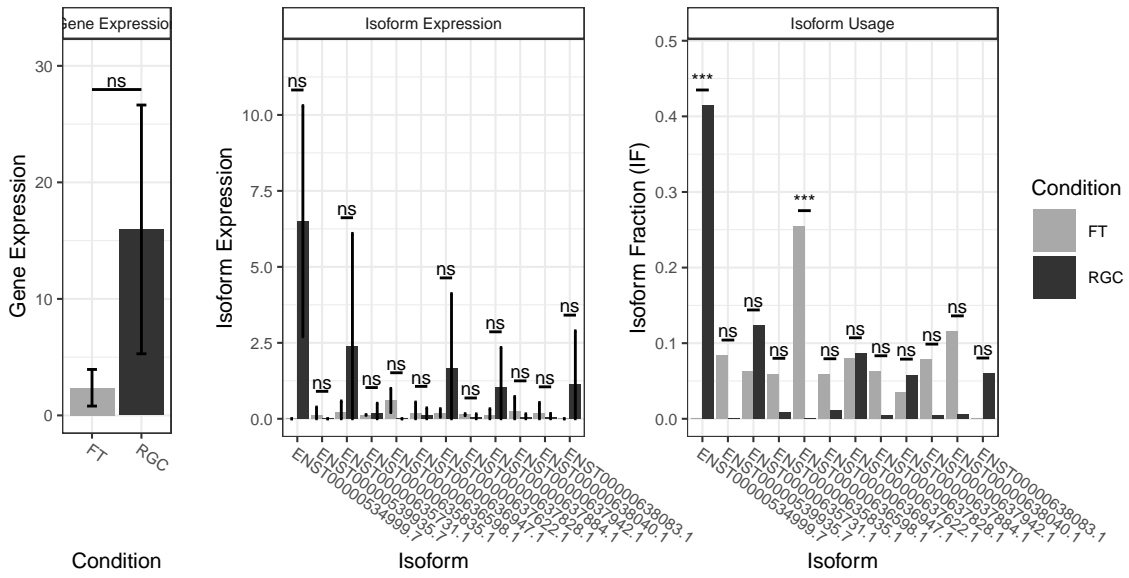
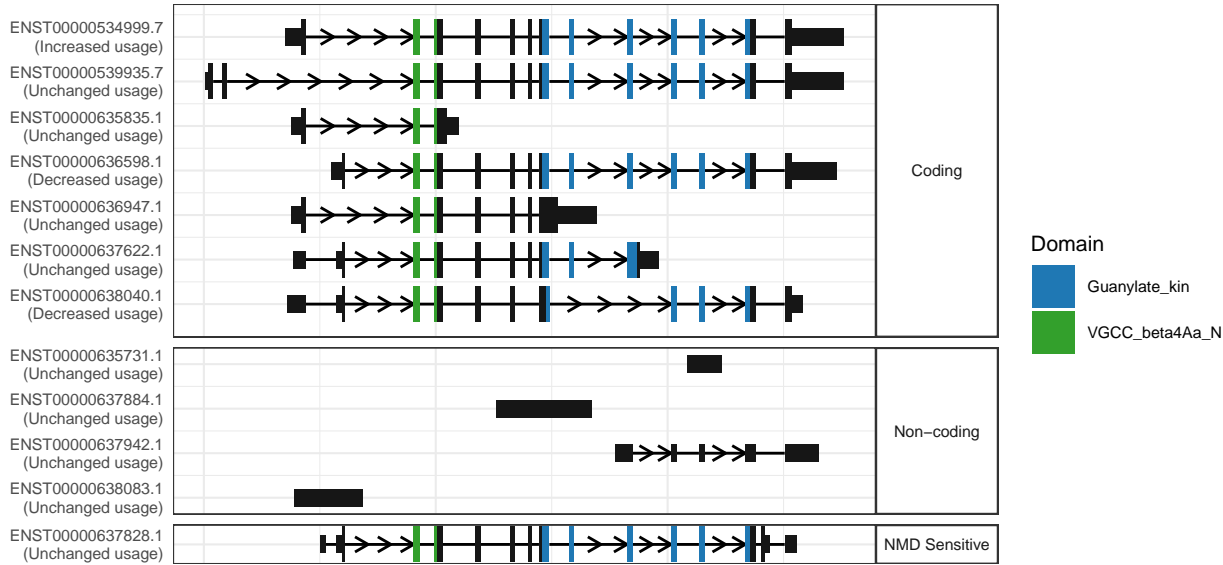
Isoform

Domain



RINGv

The isoform switch in CACNB4 (FT vs RGC)



Domain

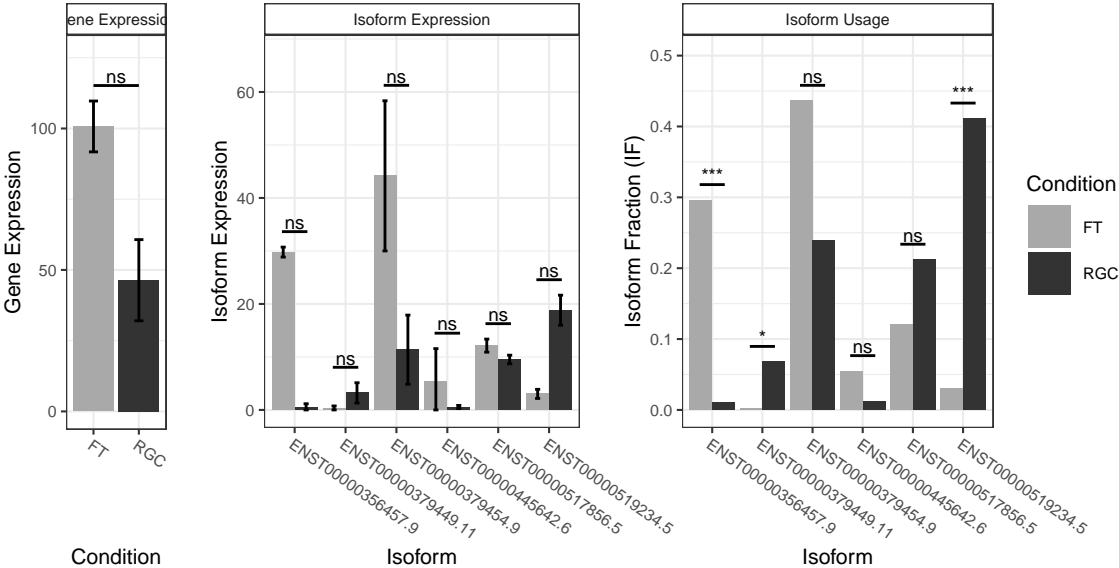
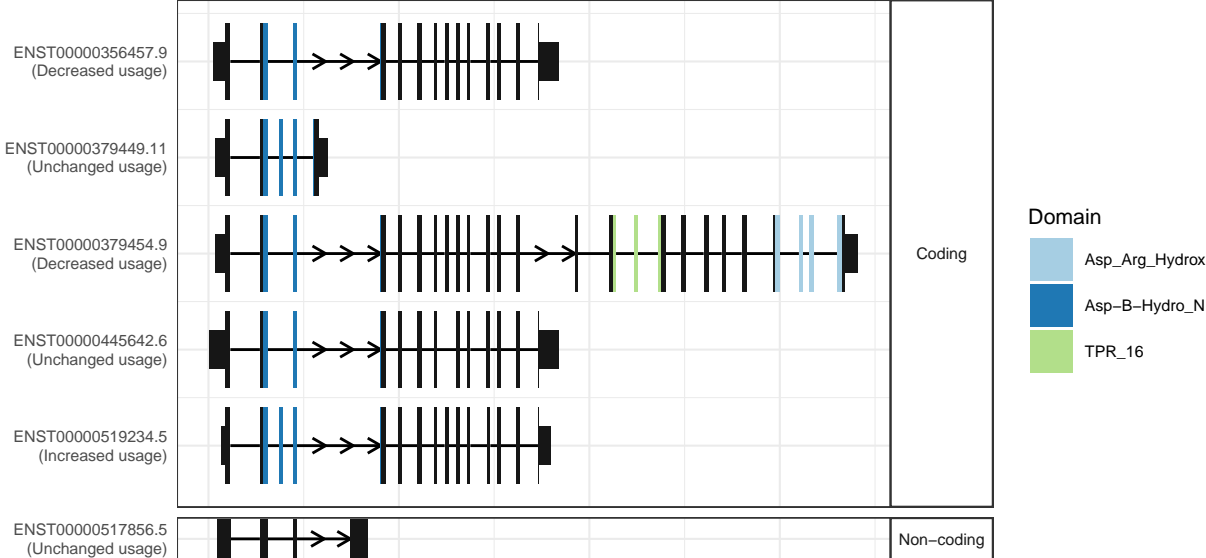


Guanylate_kin






VGCC_beta4Aa_N

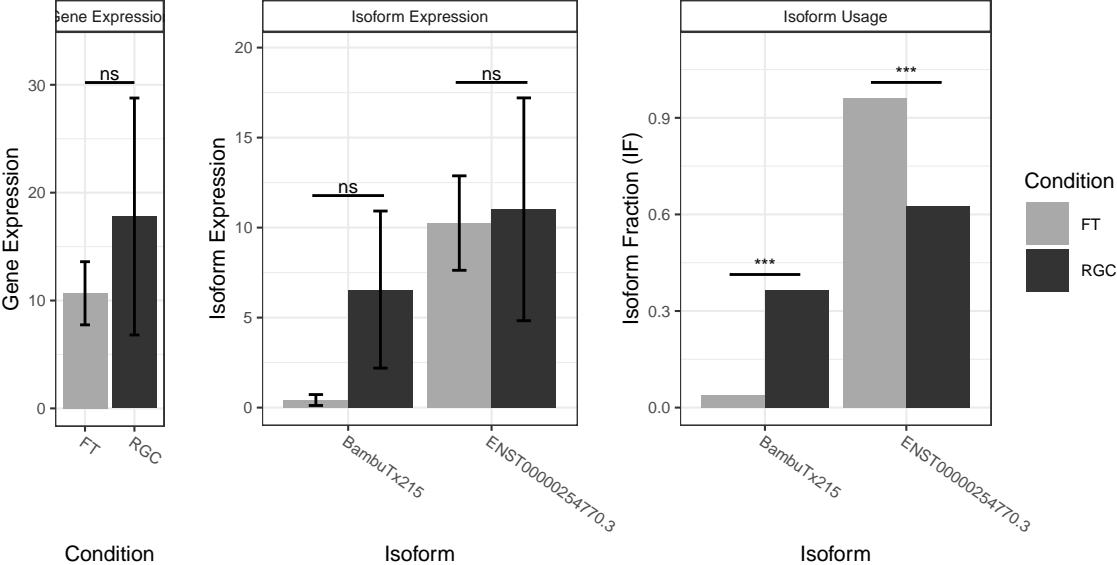
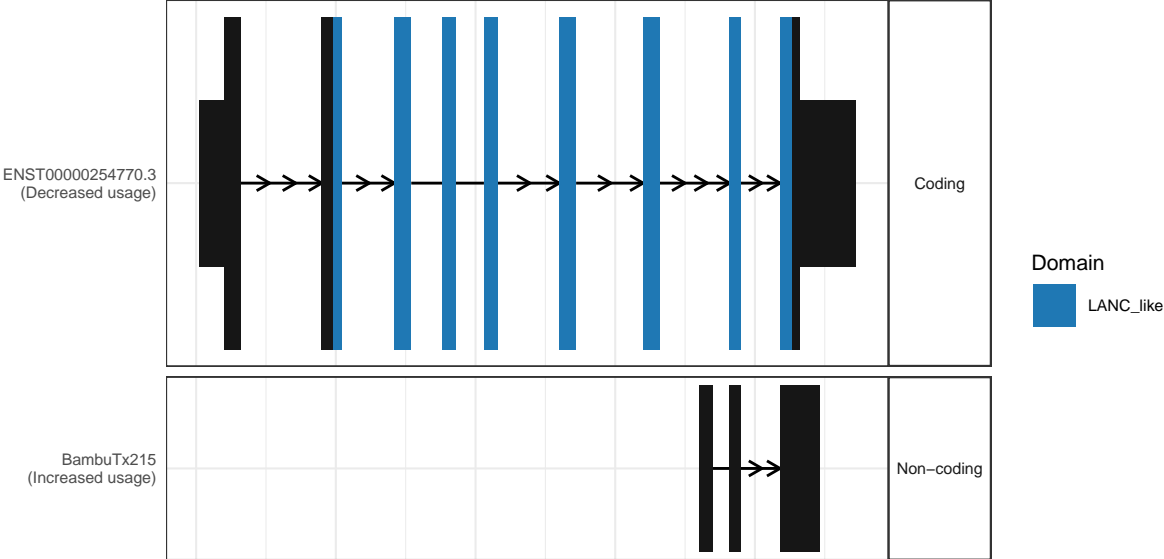
The isoform switch in ASPH (FT vs RGC)



Domain

	Asp_Arg_Hydrox
	Asp-B-Hydro_N
	TPR_16

The isoform switch in LANCL2 (FT vs RGC)

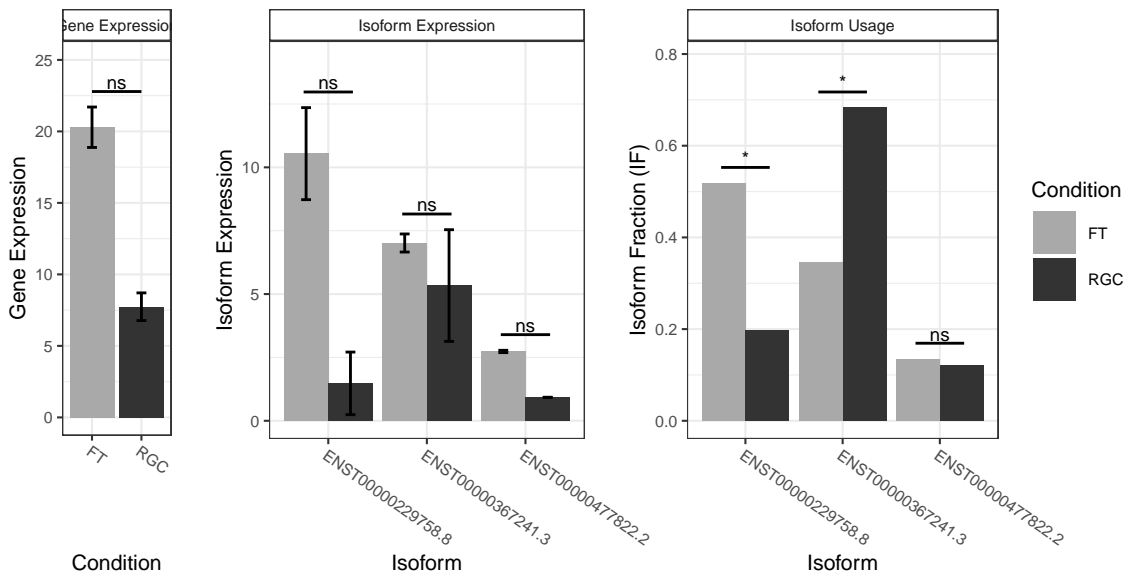
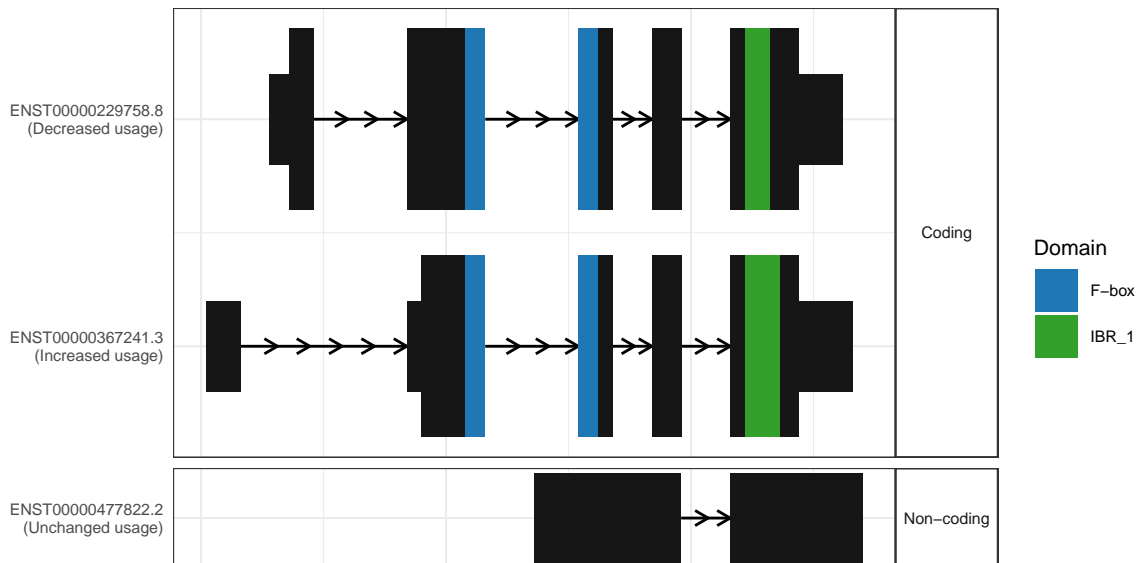


Domain



LANC_like

The isoform switch in FBXO5 (FT vs RGC)



Domain

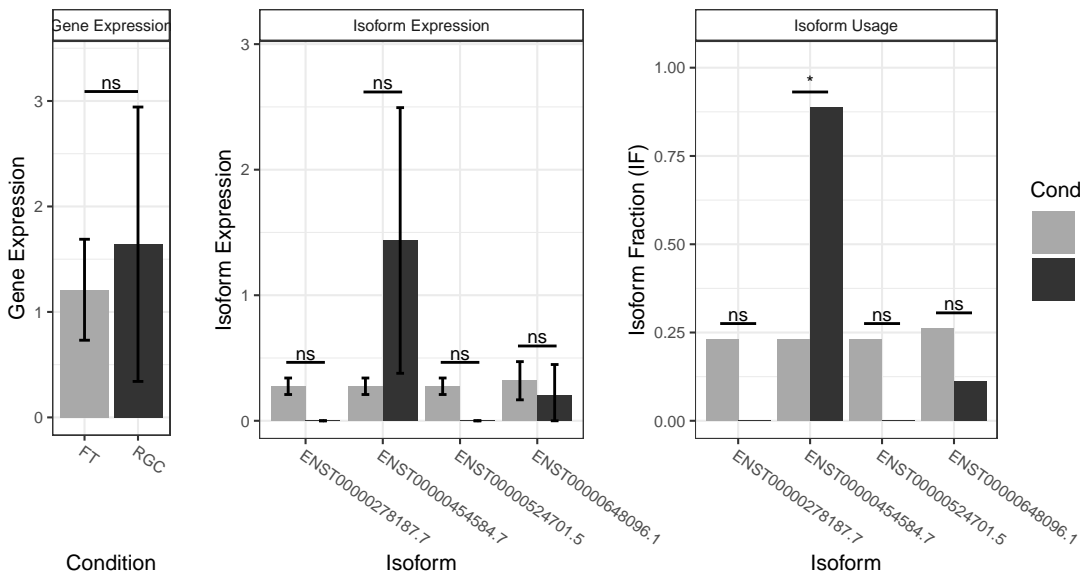
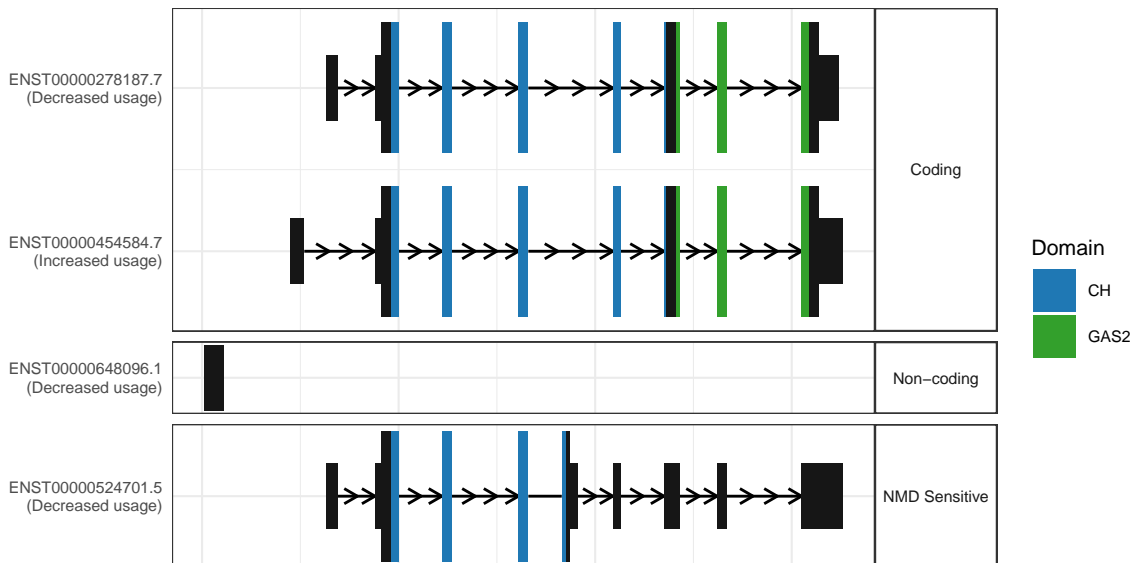


F-box



IBR_1

The isoform switch in GAS2 (FT vs RGC)



Domain

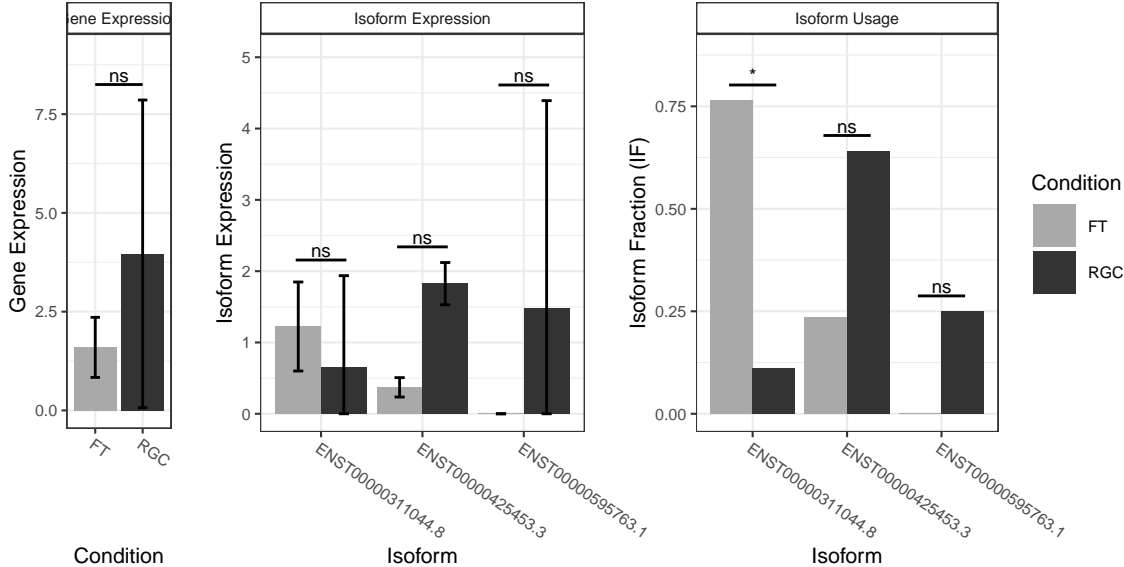
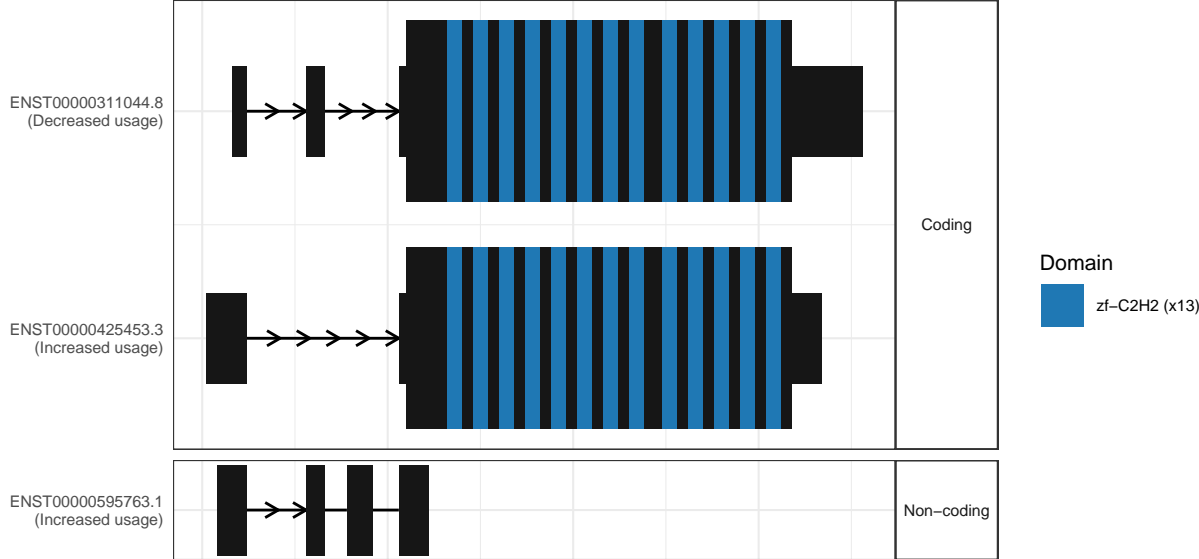


CH



GAS2

The isoform switch in ZNF497 (FT vs RGC)

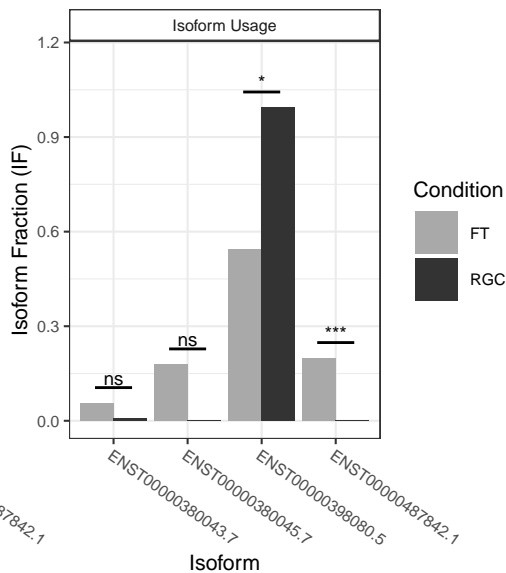
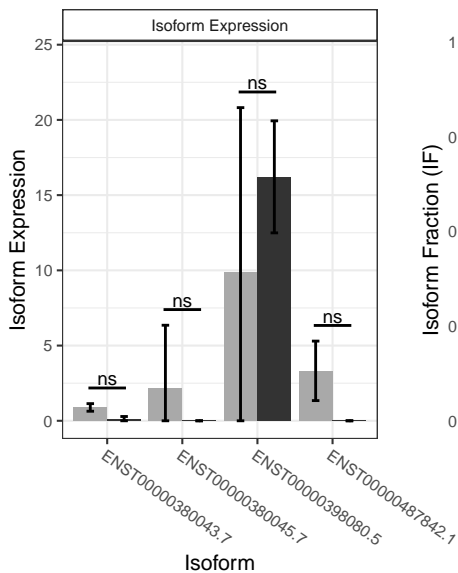
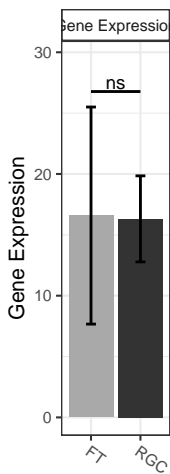
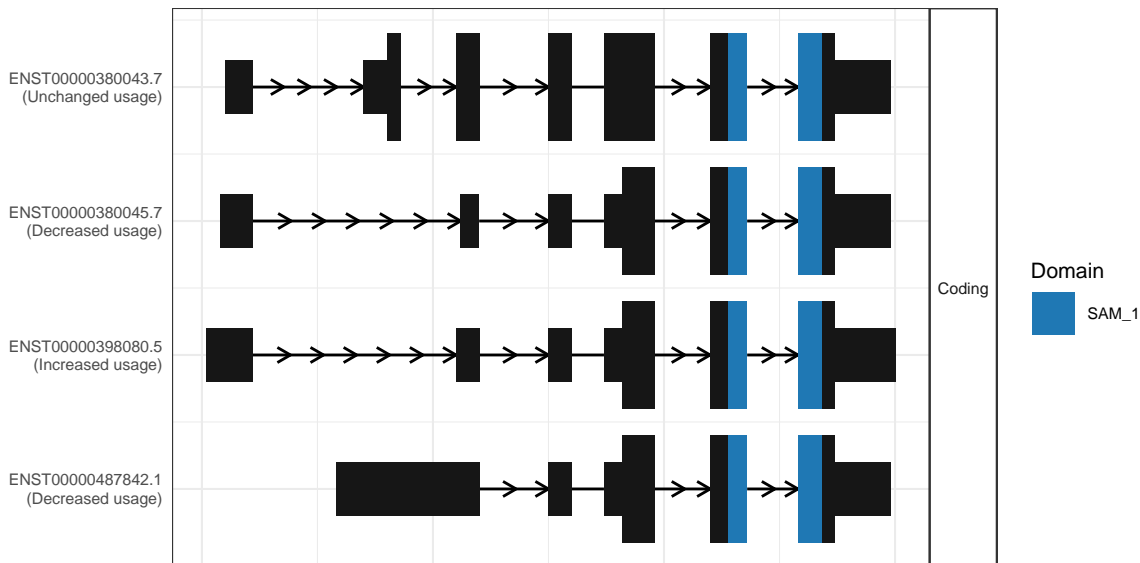


Domain



zf-C2H2 (x13)

The isoform switch in SCML1 (FT vs RGC)

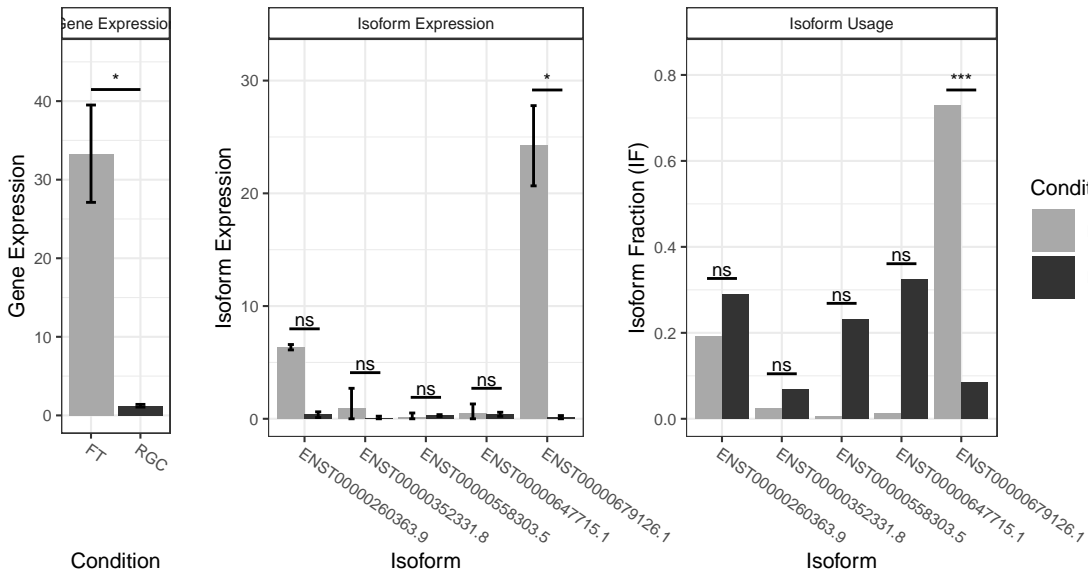
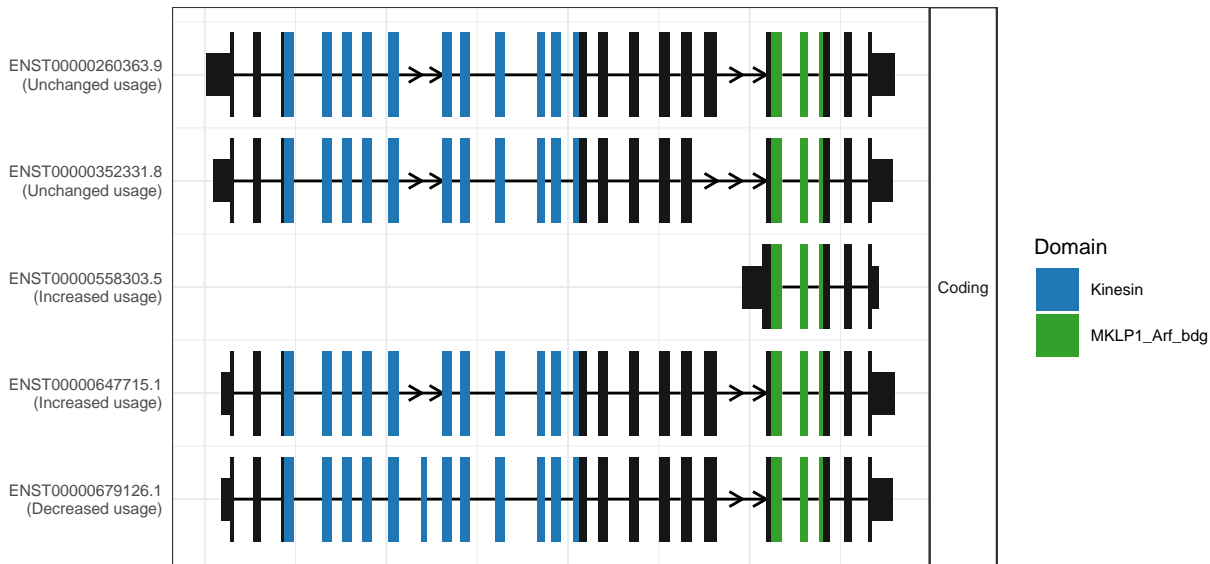


Domain



SAM_1

The isoform switch in KIF23 (FT vs RGC)



Domain

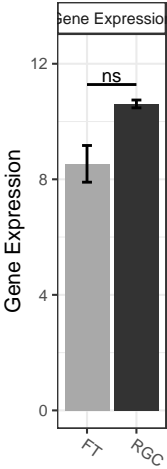
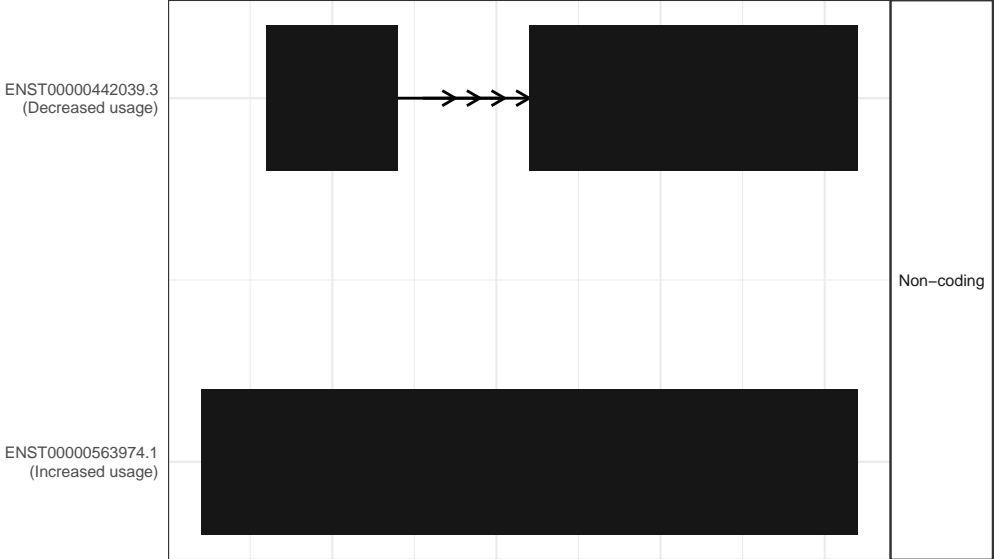


Kinesin

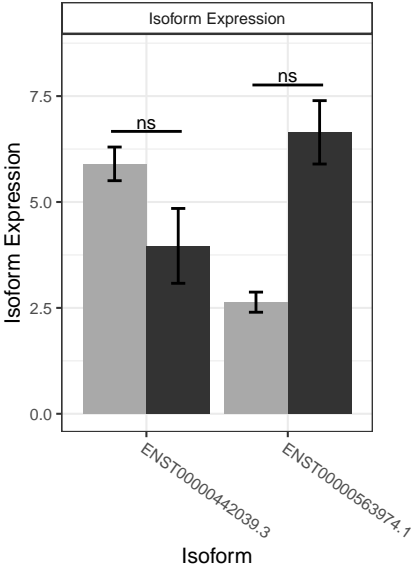


MKLP1_Arf_bdg

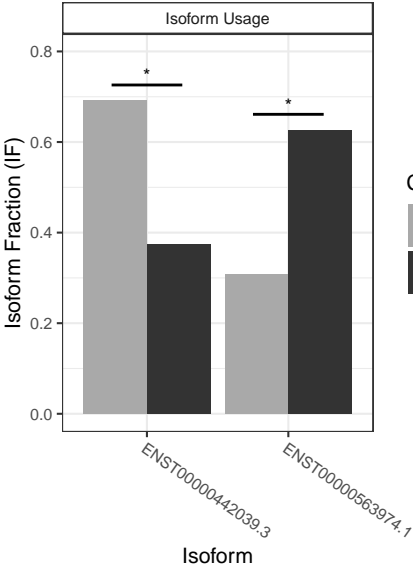
The isoform switch in UQCC4 (FT vs RGC)



Condition



Isoform

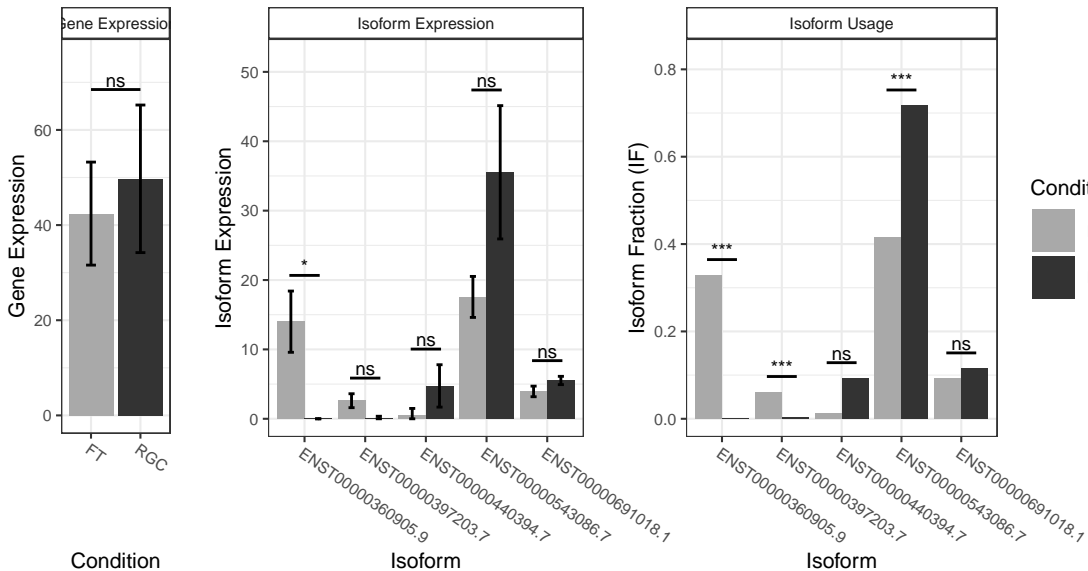
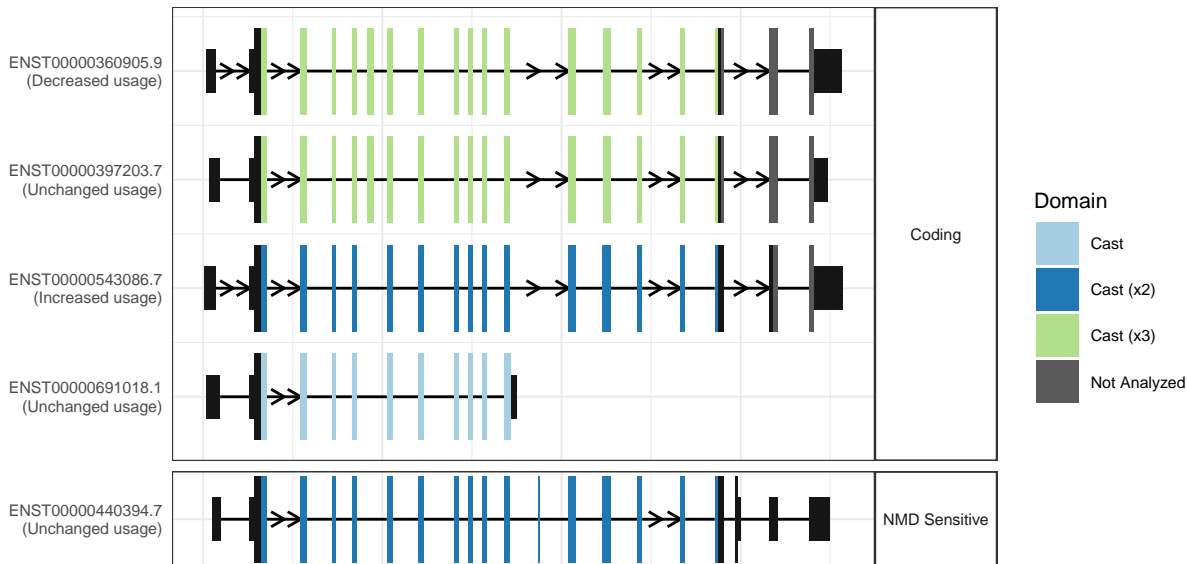


Isoform

Condition



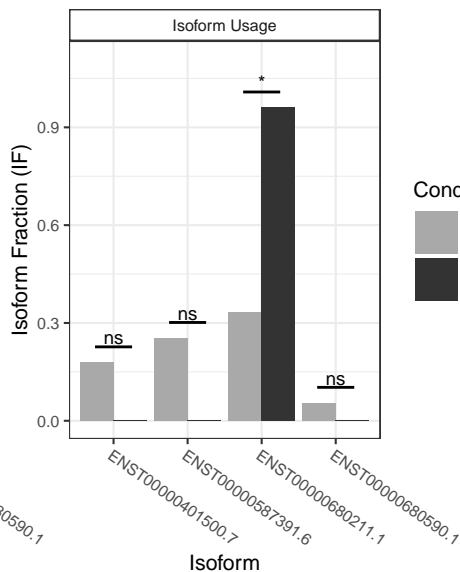
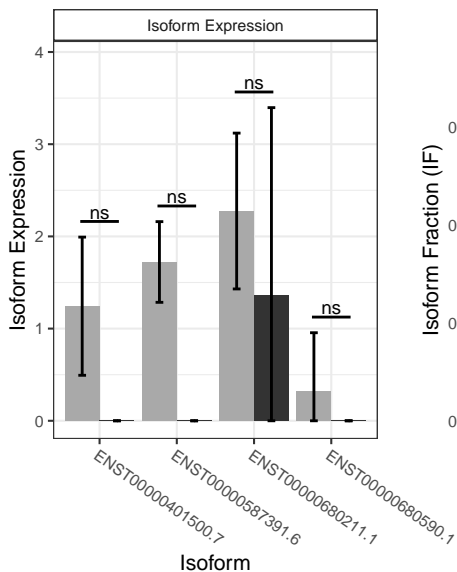
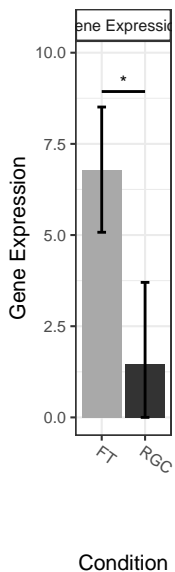
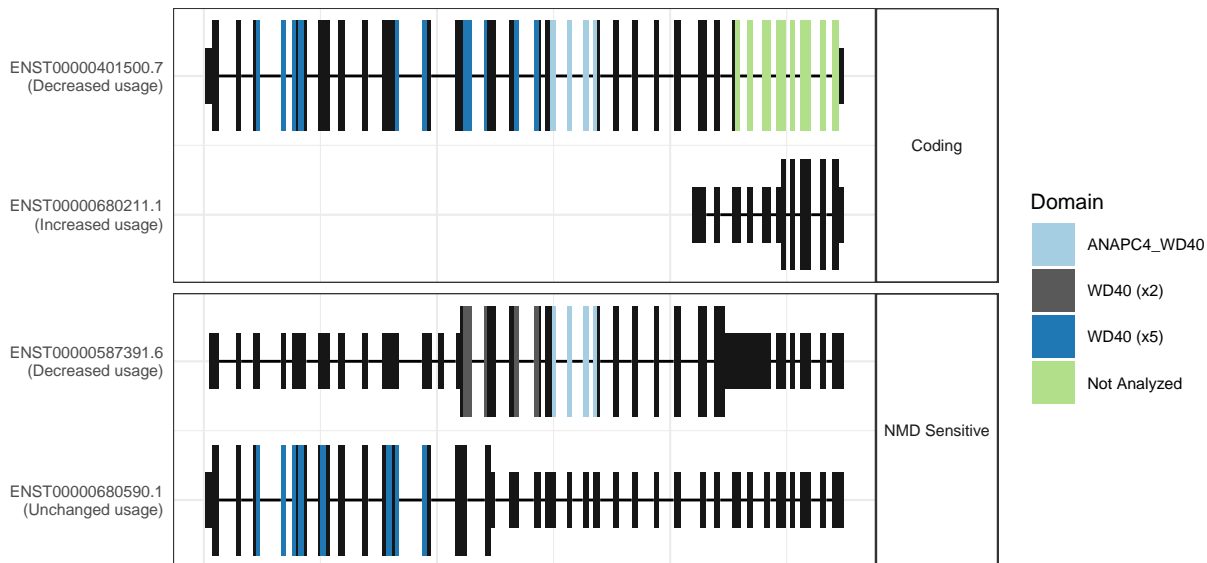
The isoform switch in ERC1 (FT vs RGC)





Domain



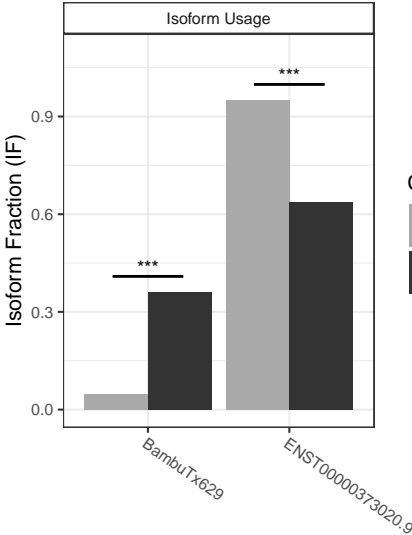
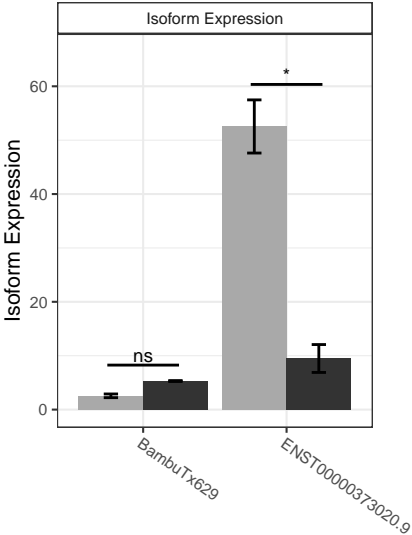
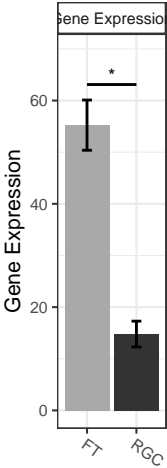
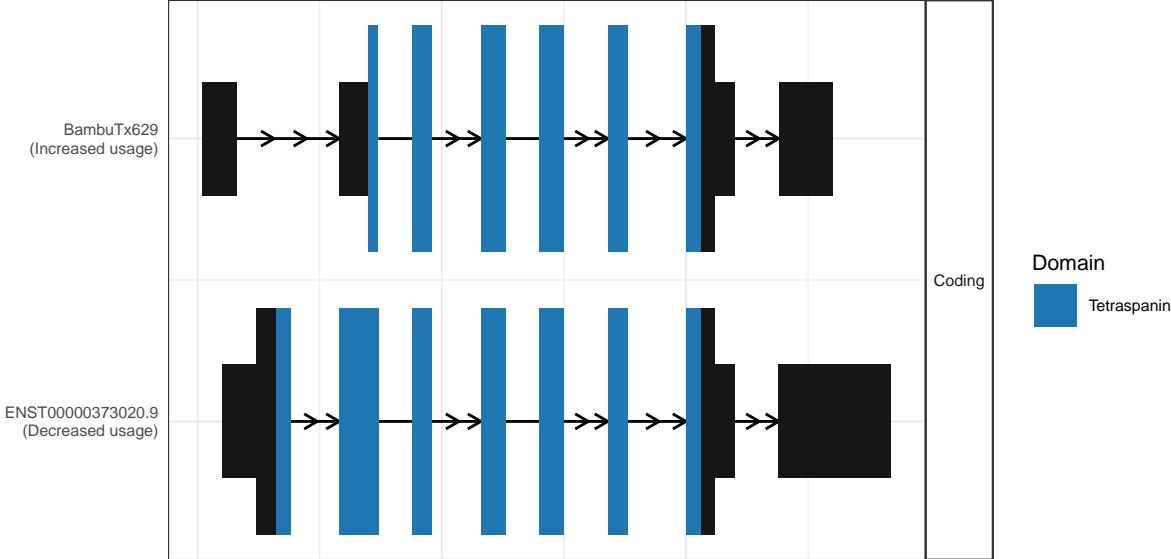
The isoform switch in WDR62 (FT vs RGC)



Domain

	ANAPC4_WD40
	WD40 (x2)
	WD40 (x5)
	Not Analyzed

The isoform switch in TSPAN6 (FT vs RGC)



Condition

Isoform

Isoform

Condition

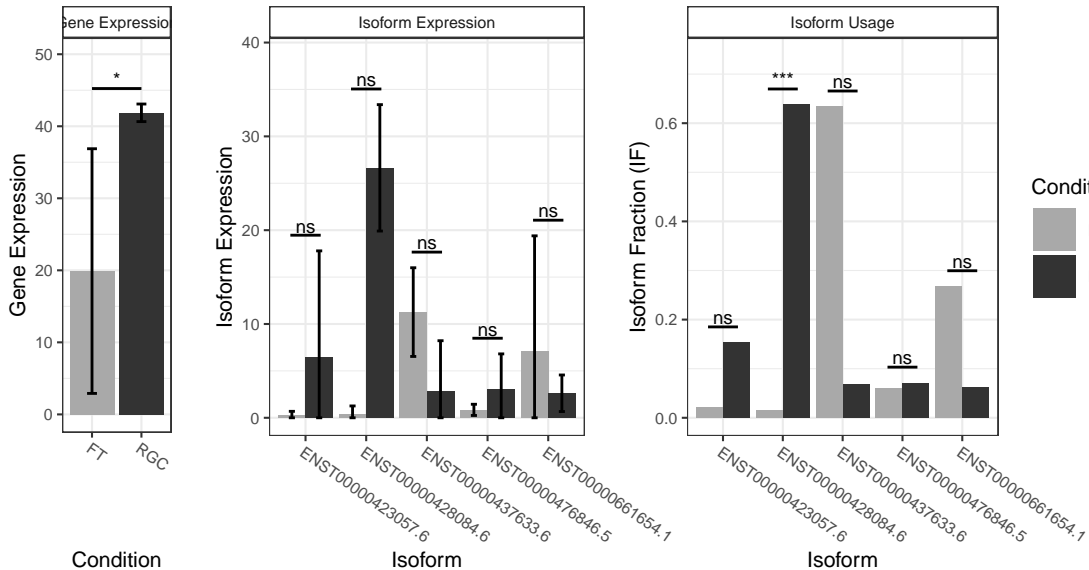
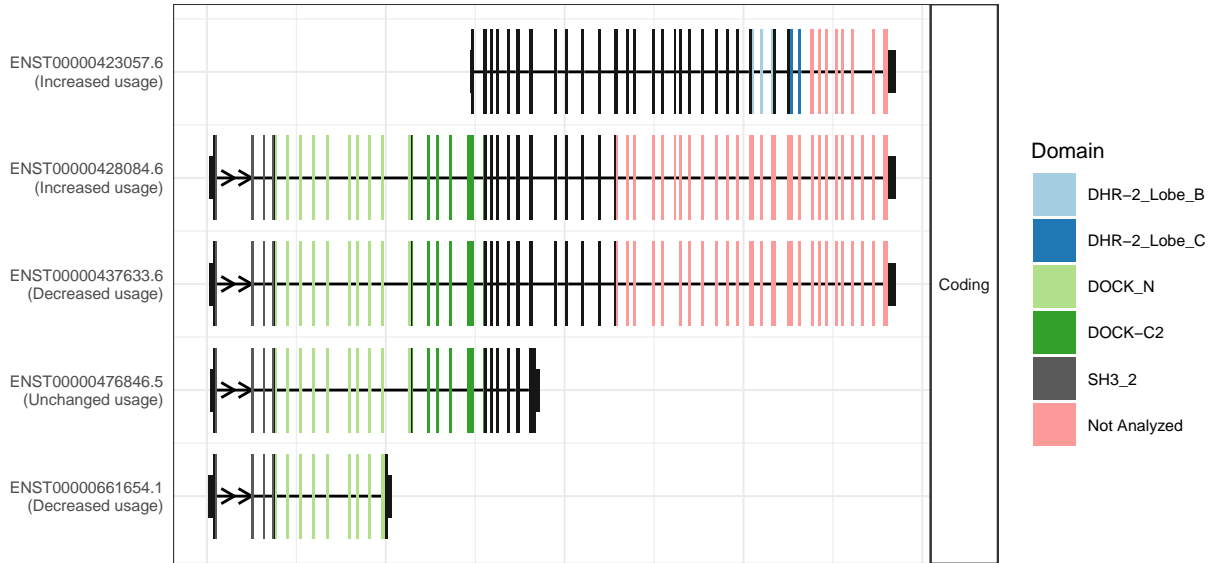


Domain









Tetraspanin

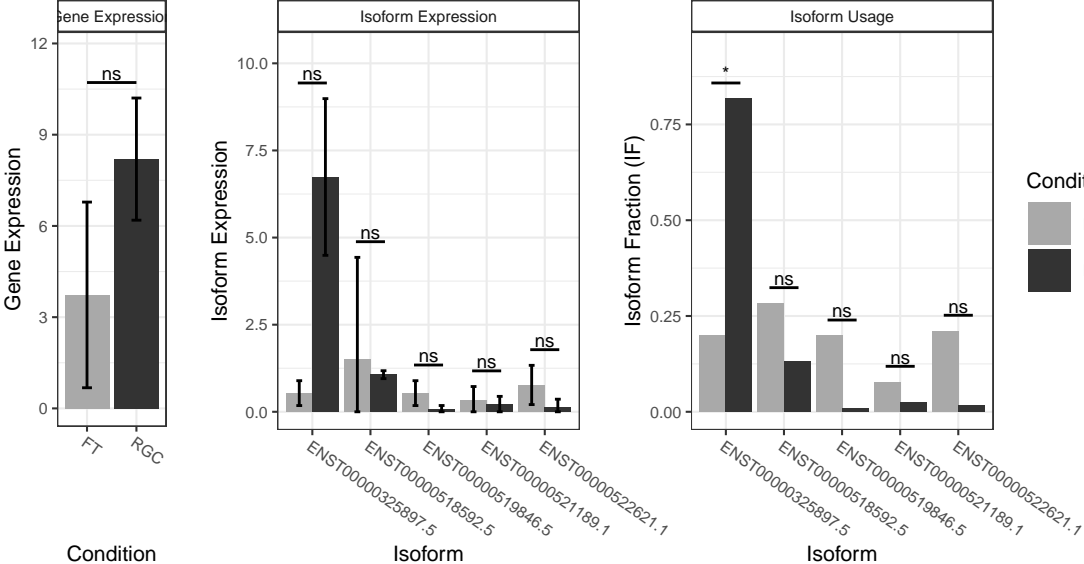
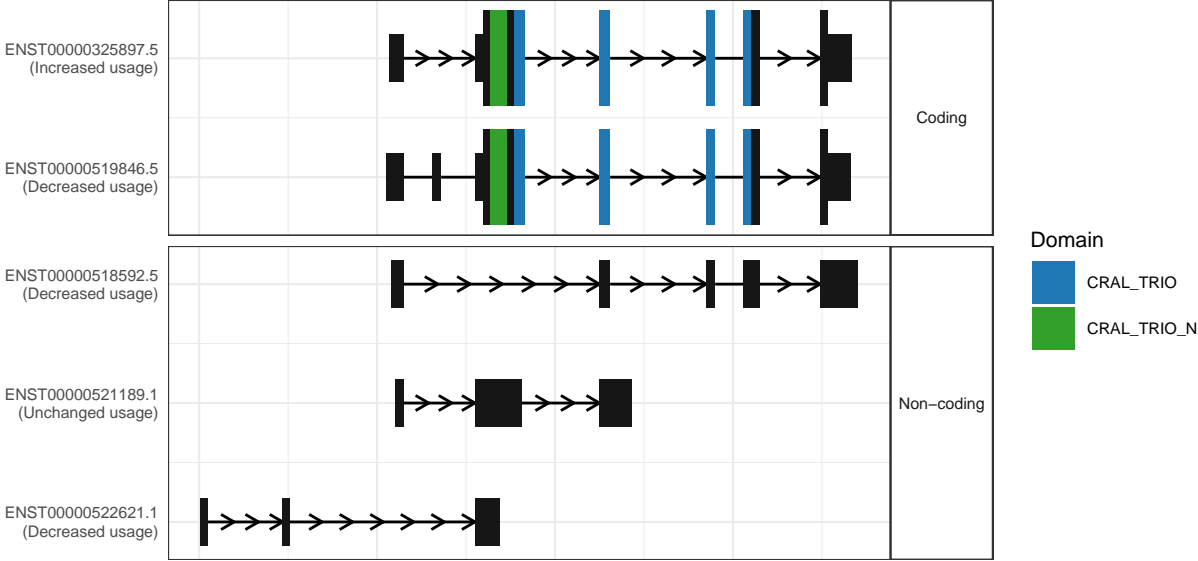
The isoform switch in DOCK4 (FT vs RGC)



Domain

	DHR-2_Lobe_B
	DHR-2_Lobe_C
	DOCK_N
	DOCK-C2
	SH3_2
	Not Analyzed

The isoform switch in CLVS1 (FT vs RGC)



Domain

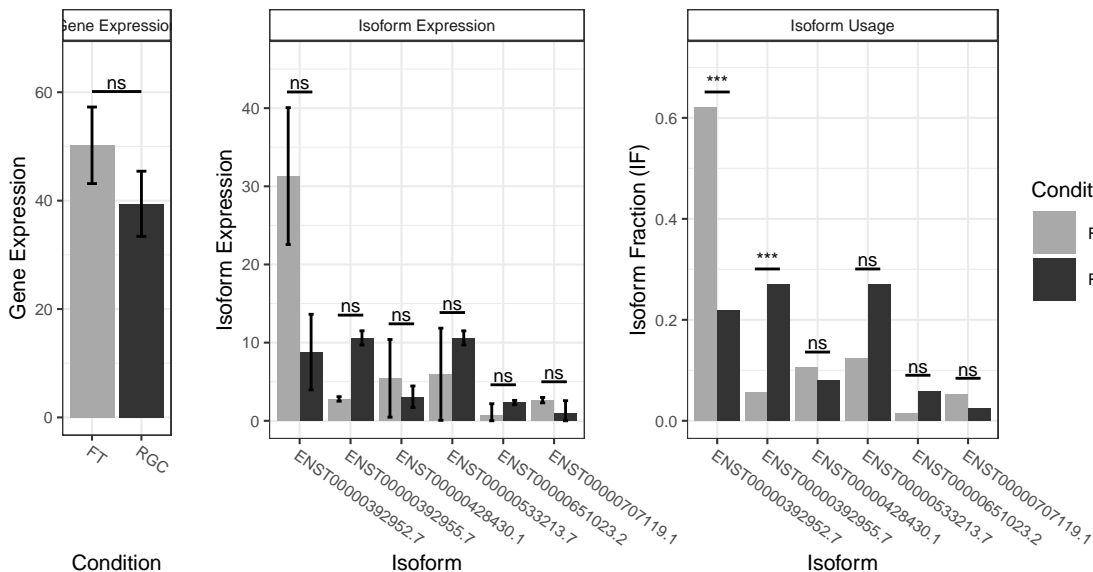
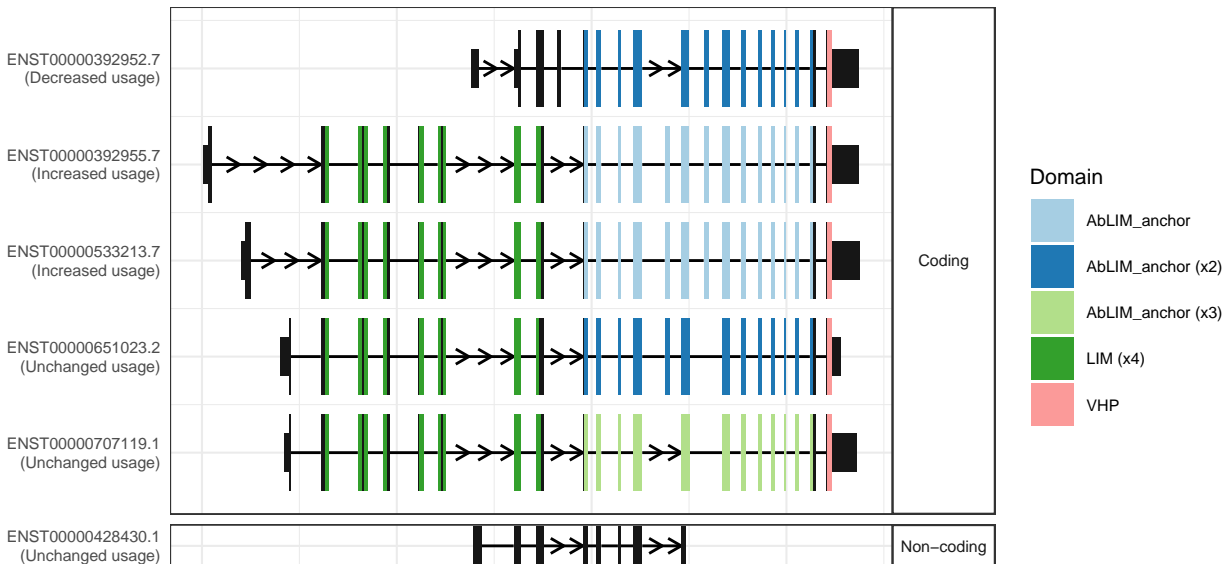


CRAL_TRIO








CRAL_TRIO_N

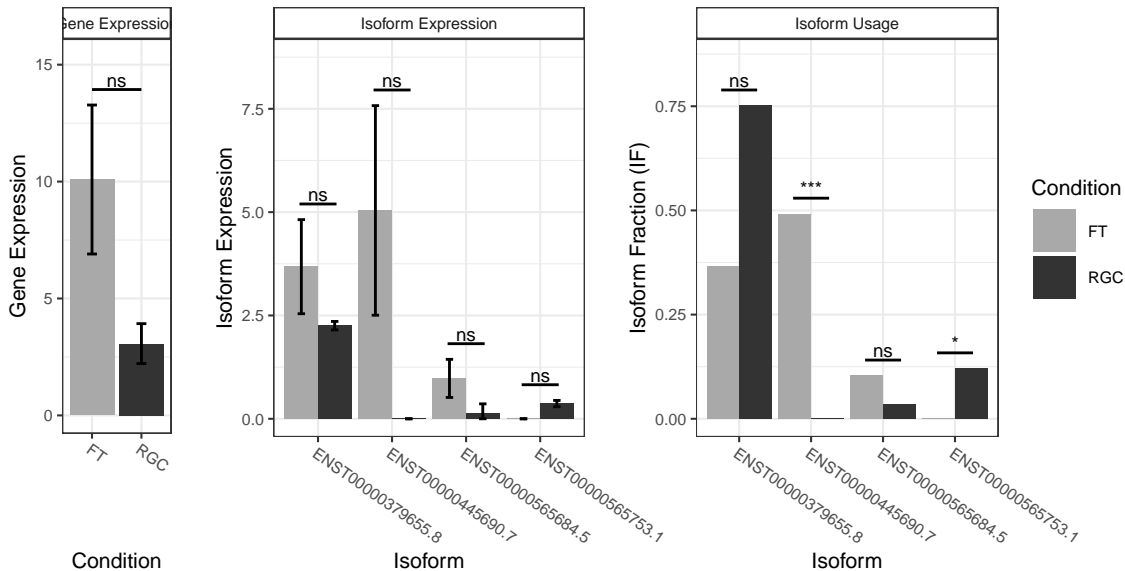
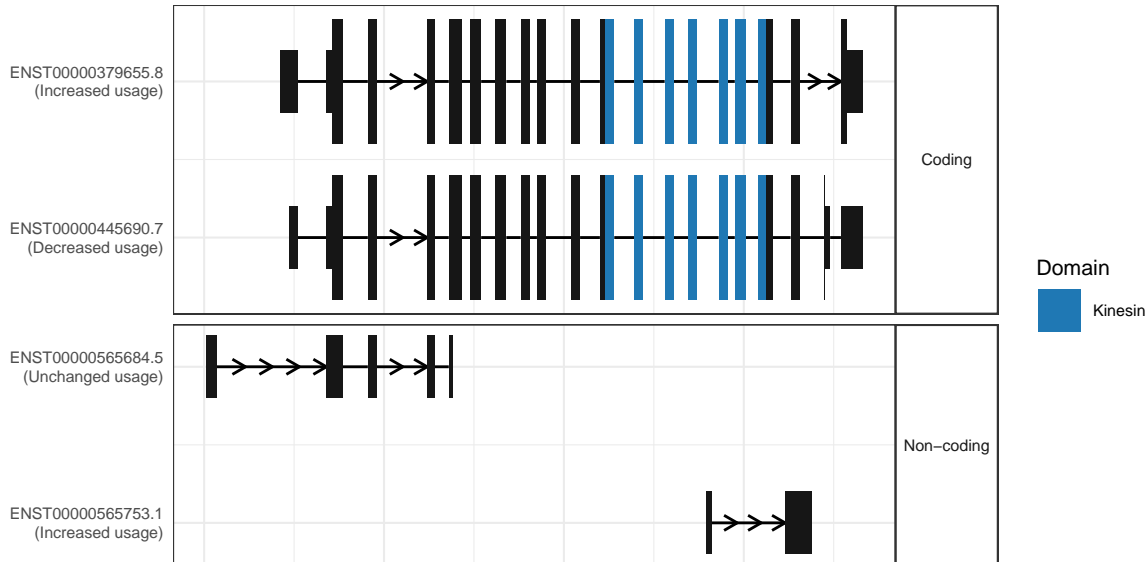
The isoform switch in ABLIM1 (FT vs RGC)



Domain

	AbLIM_anchor
	AbLIM_anchor (x2)
	AbLIM_anchor (x3)
	LIM (x4)
	VHP

The isoform switch in KIFC3 (FT vs RGC)

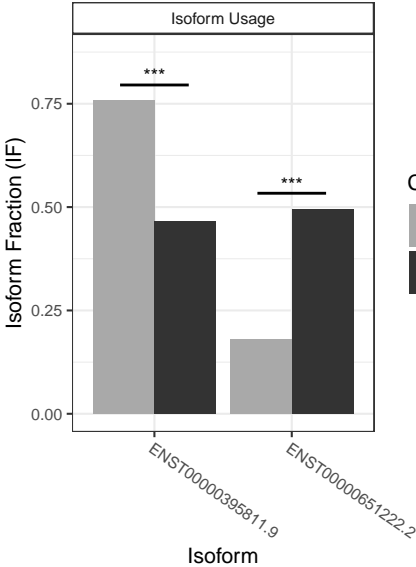
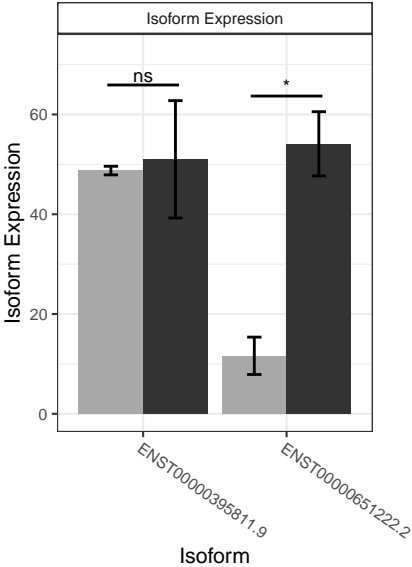
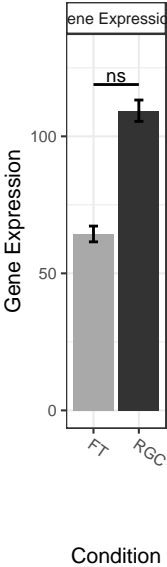
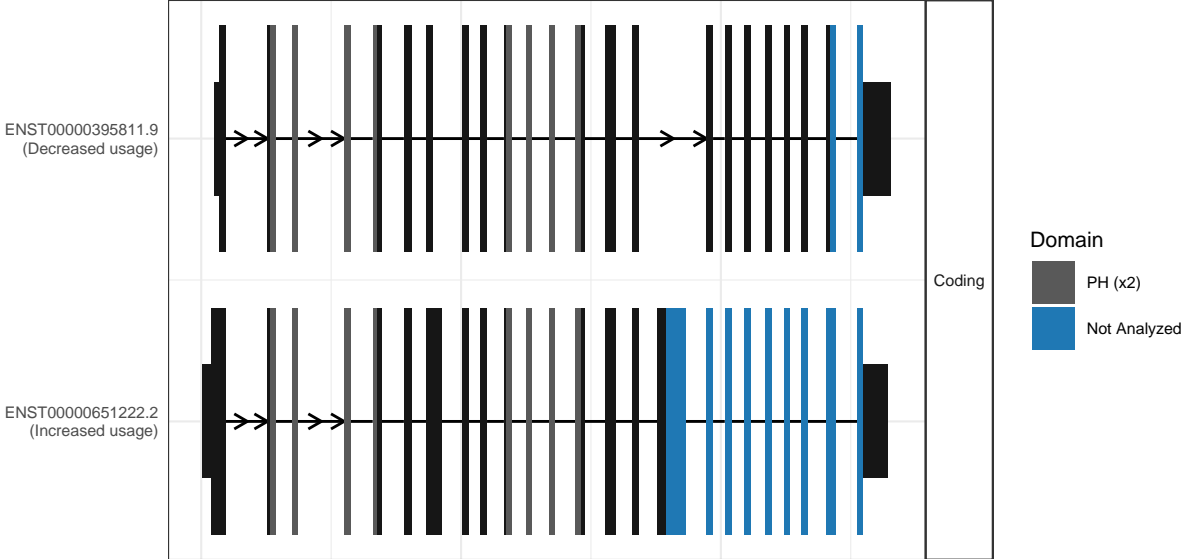


Domain



Kinesin

The isoform switch in MPRIP (FT vs RGC)



Domain

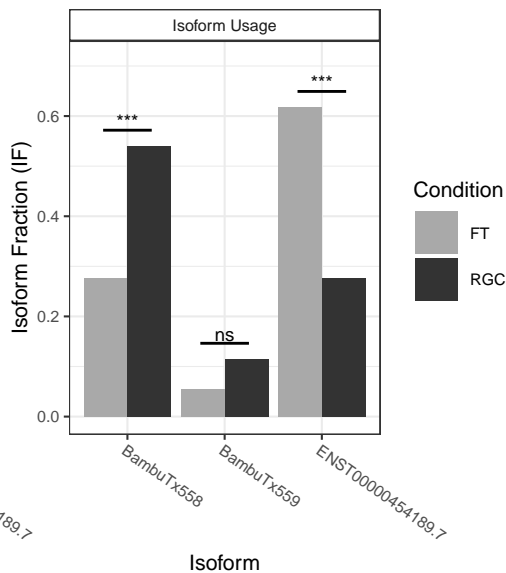
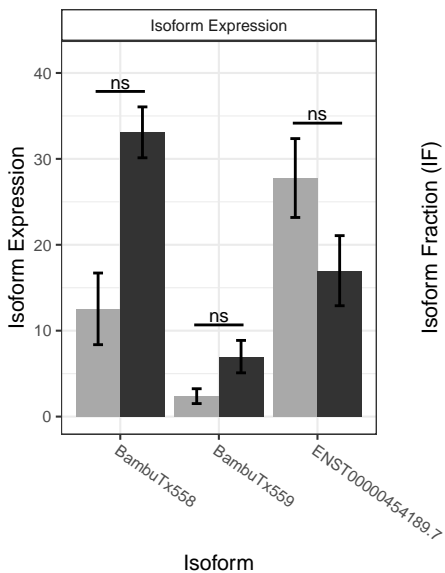
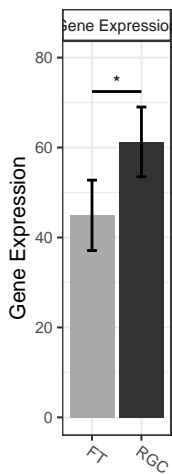
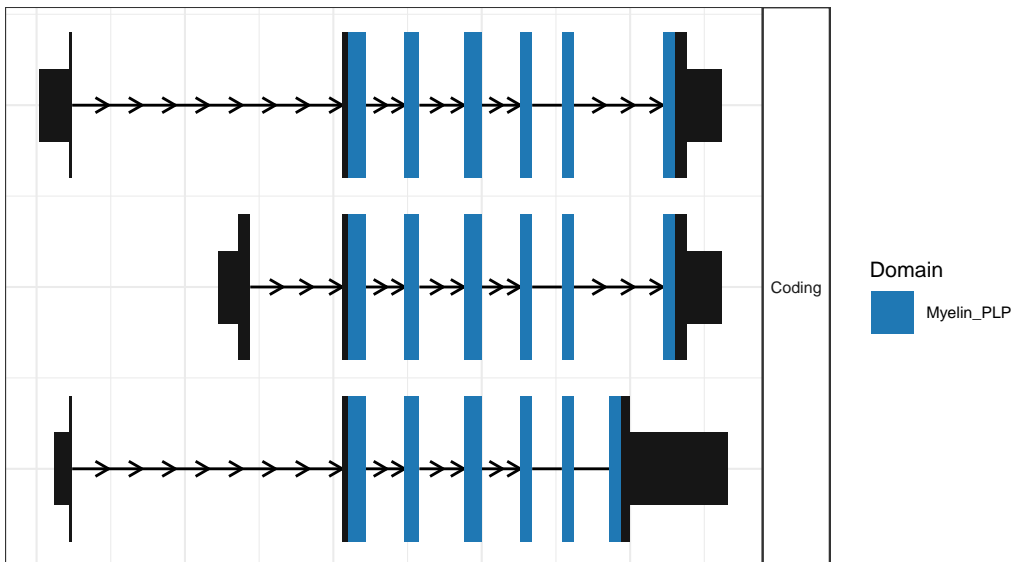


PH (x2)



Not Analyzed

The isoform switch in GPM6B (FT vs RGC)

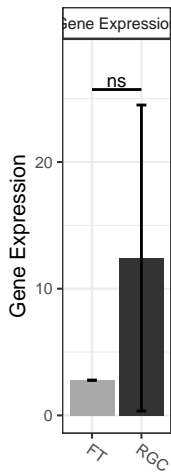
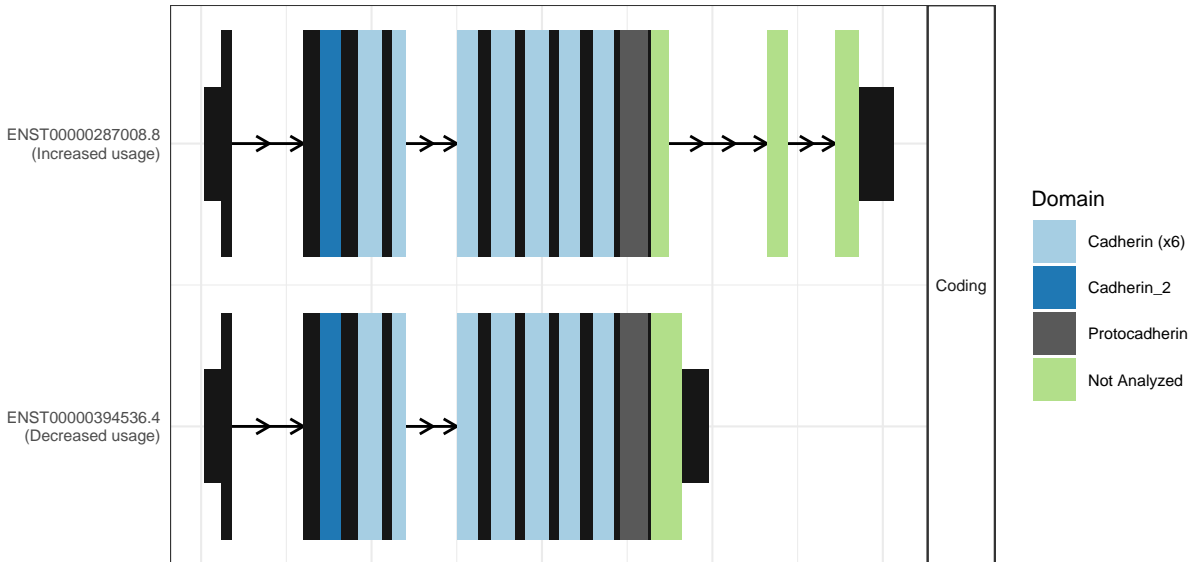


Domain

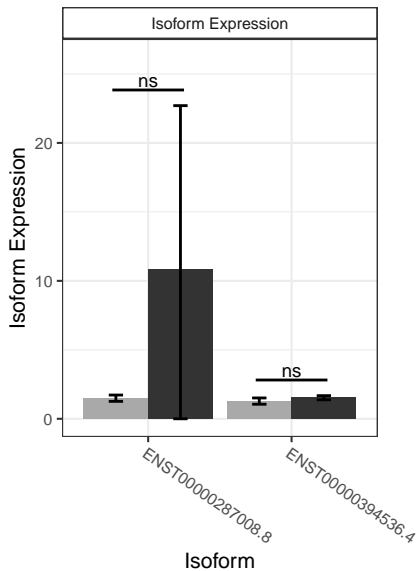


Myelin_PLP

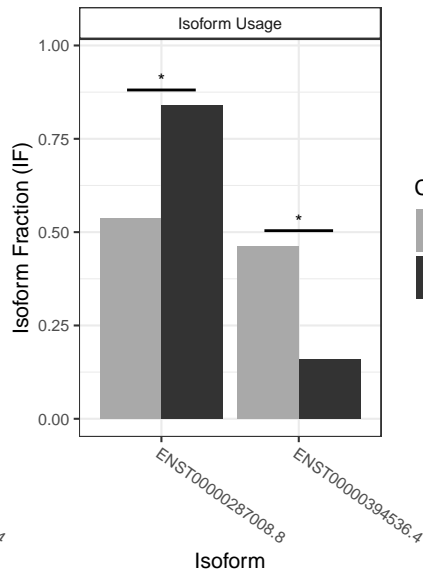
The isoform switch in PCDH1 (FT vs RGC)



Condition



Isoform



Isoform

Condition



Domain



Cadherin (x6)



Cadherin_2

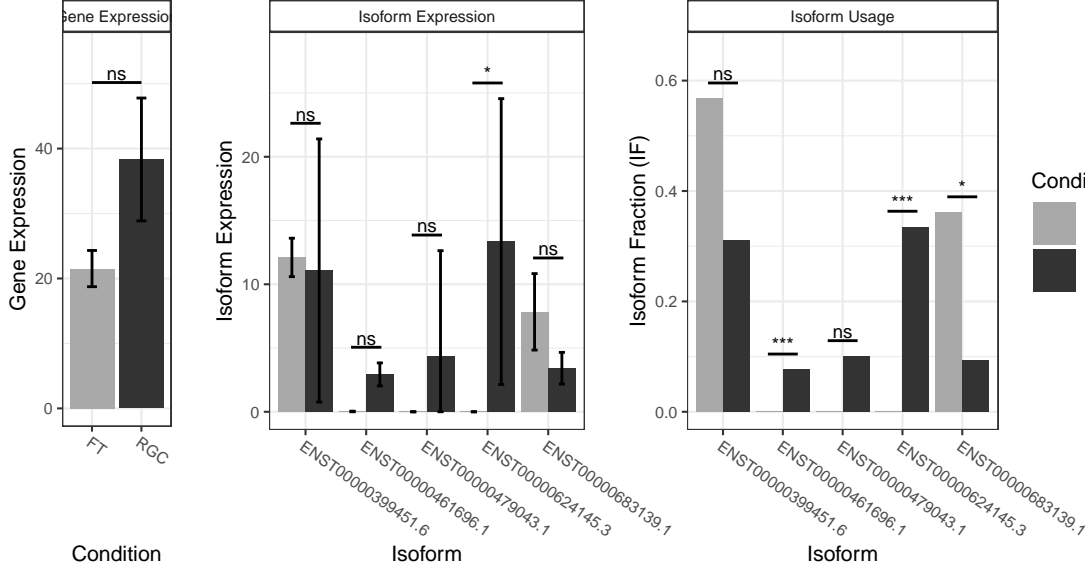
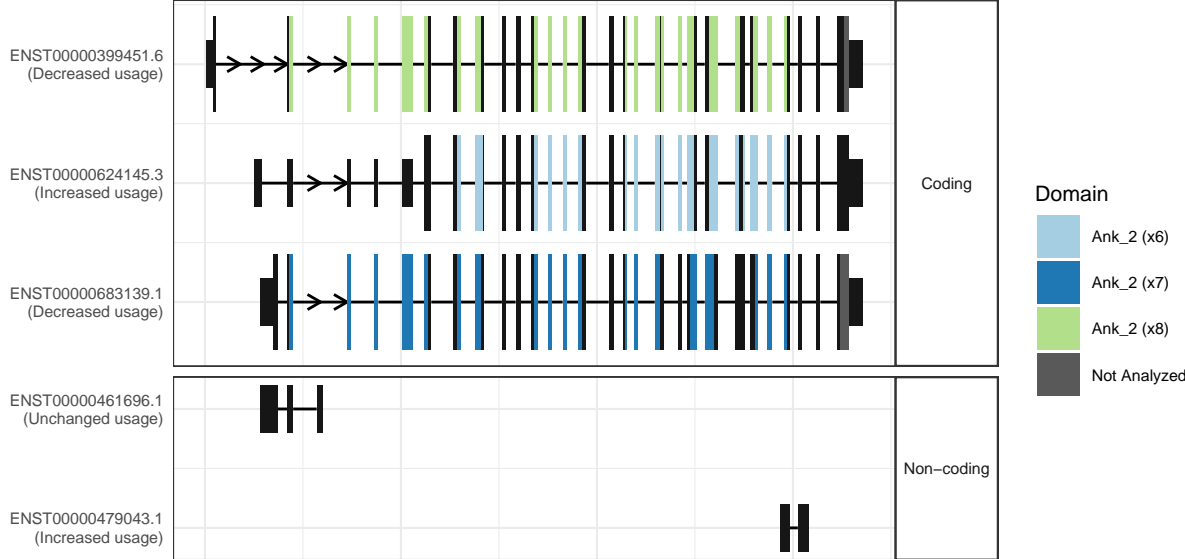


Protocadherin



Not Analyzed

The isoform switch in ANKRD28 (FT vs RGC)



Domain



Ank_2 (x6)



Ank_2 (x7)

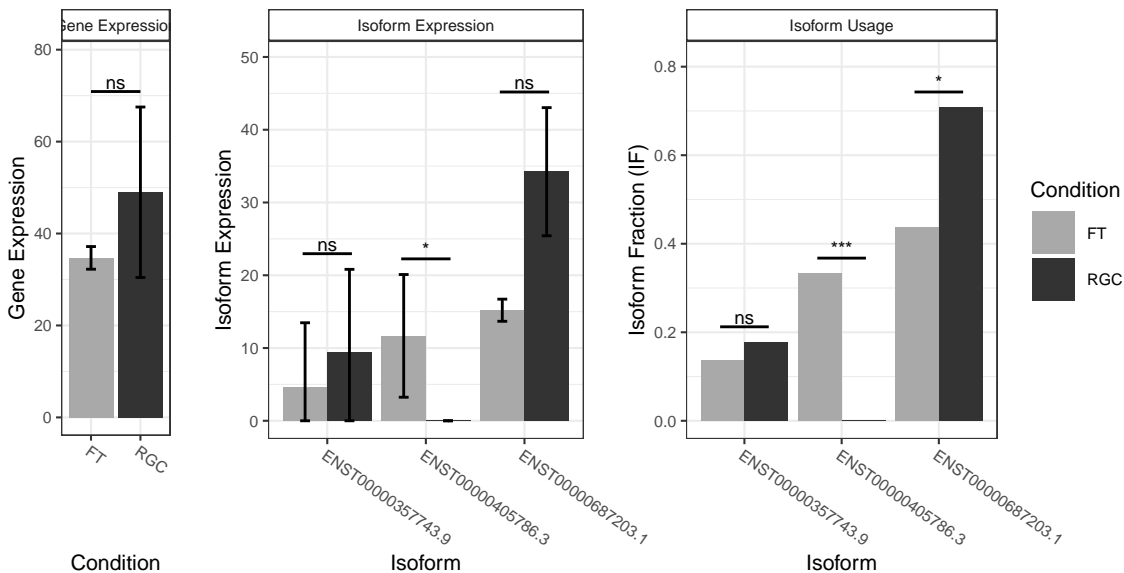
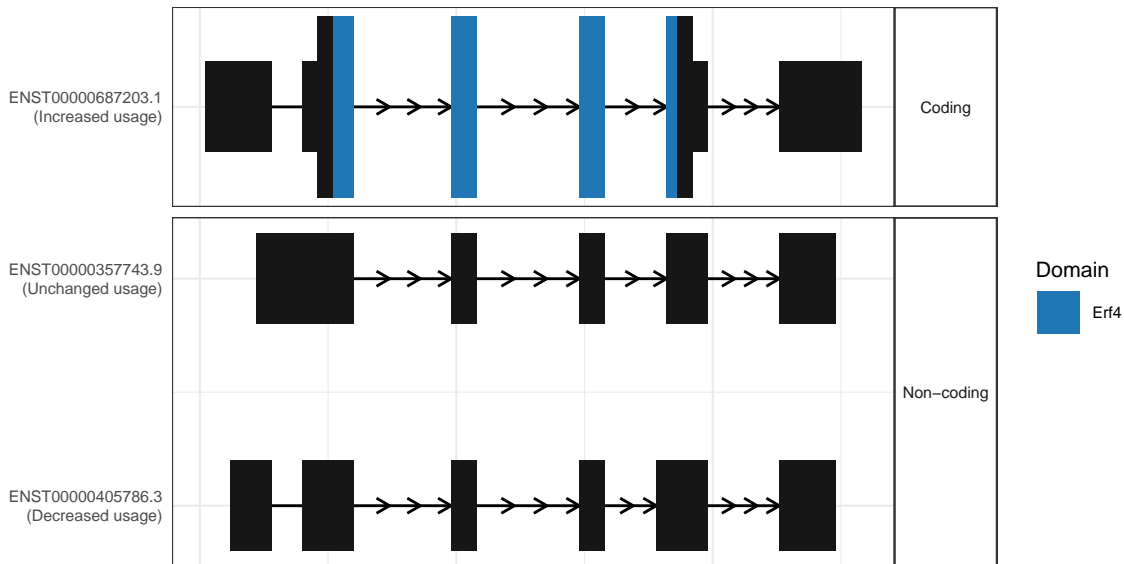


Ank_2 (x8)



Not Analyzed

The isoform switch in GOLGA7 (FT vs RGC)

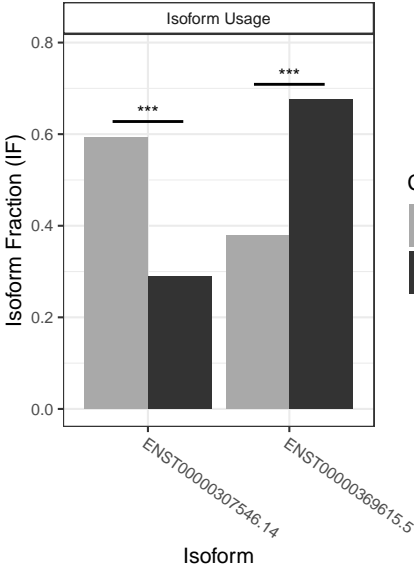
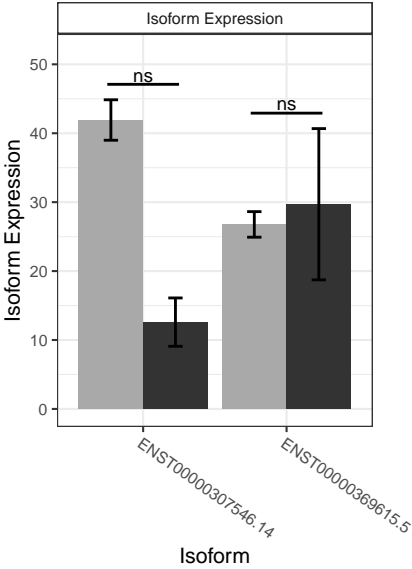
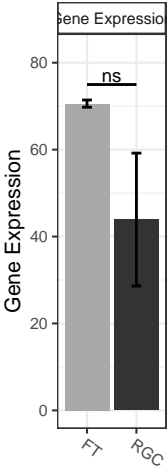
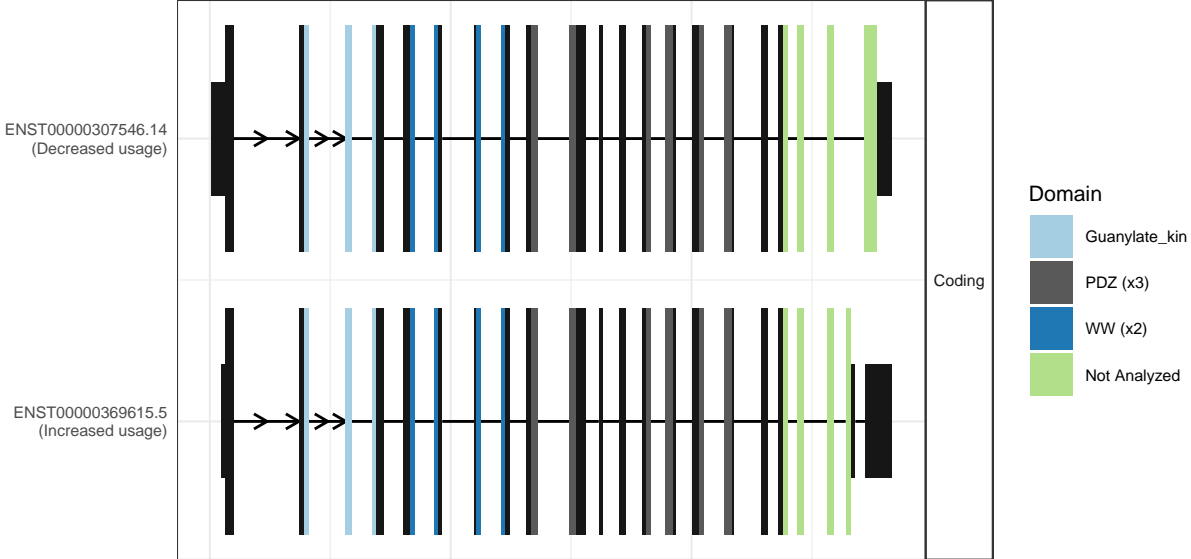


Domain







Erf4

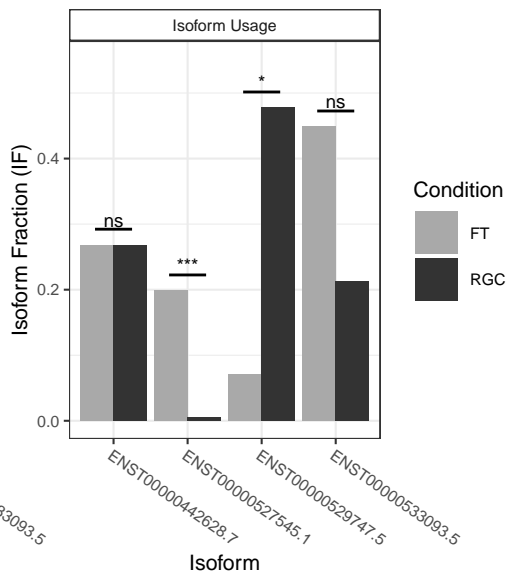
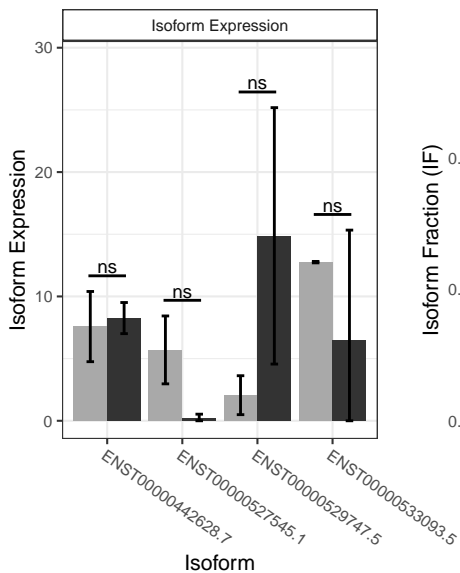
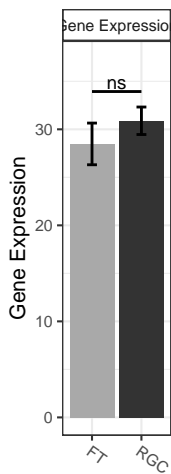
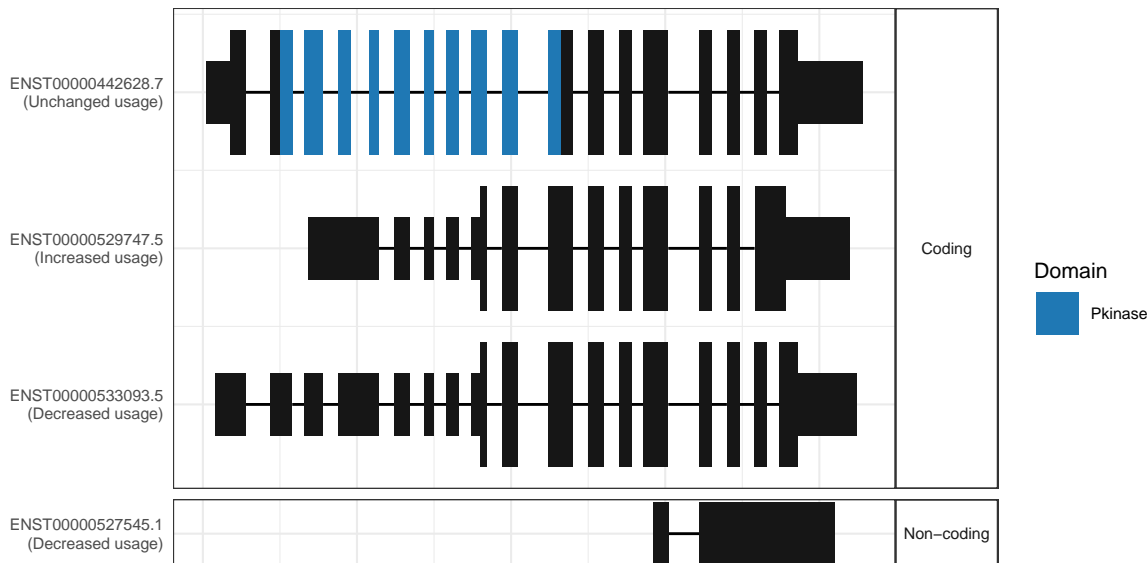
The isoform switch in MAGI3 (FT vs RGC)



Domain

	Guanylate_kin
	PDZ (x3)
	WW (x2)
	Not Analyzed

The isoform switch in NRBP2 (FT vs RGC)

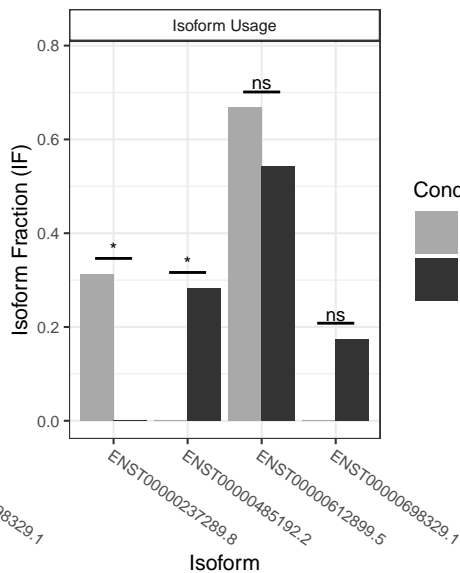
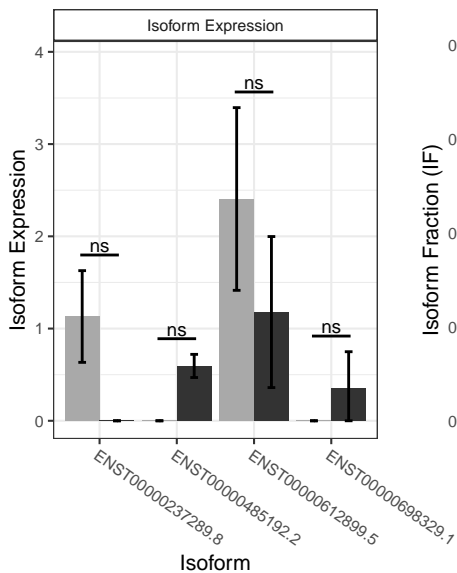
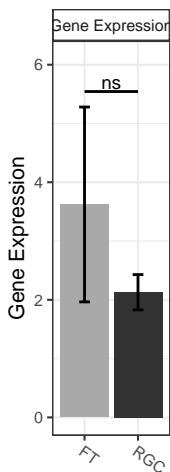
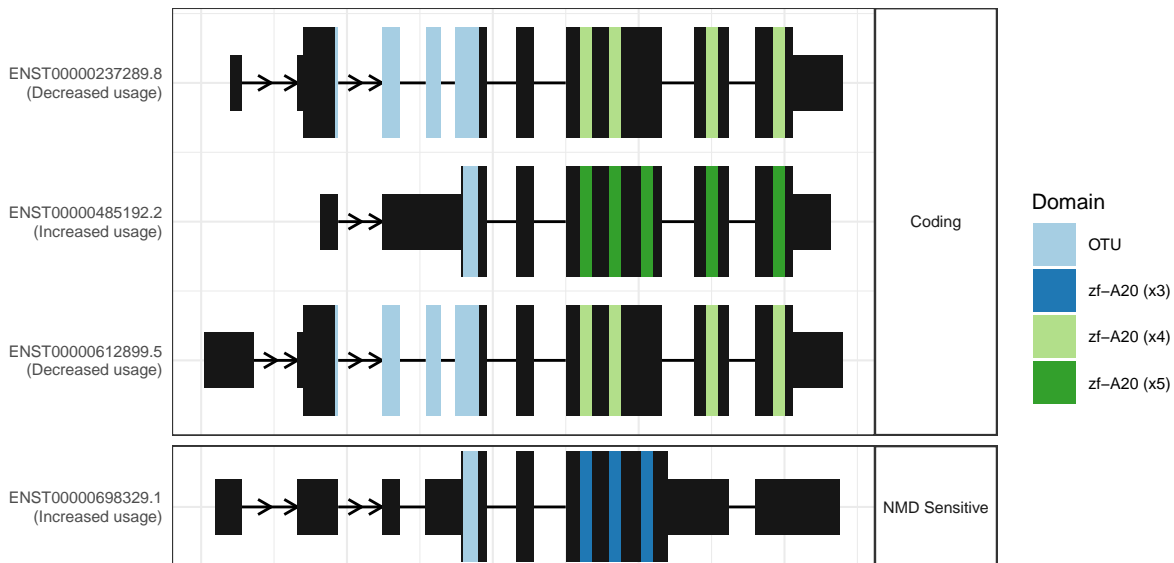


Domain



Pkinase

The isoform switch in TNFAIP3 (FT vs RGC)



Domain



OTU



zf-A20 (x3)

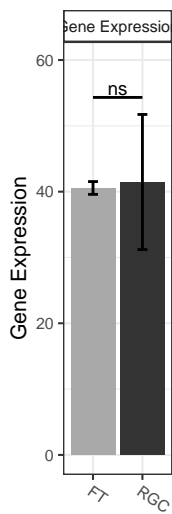
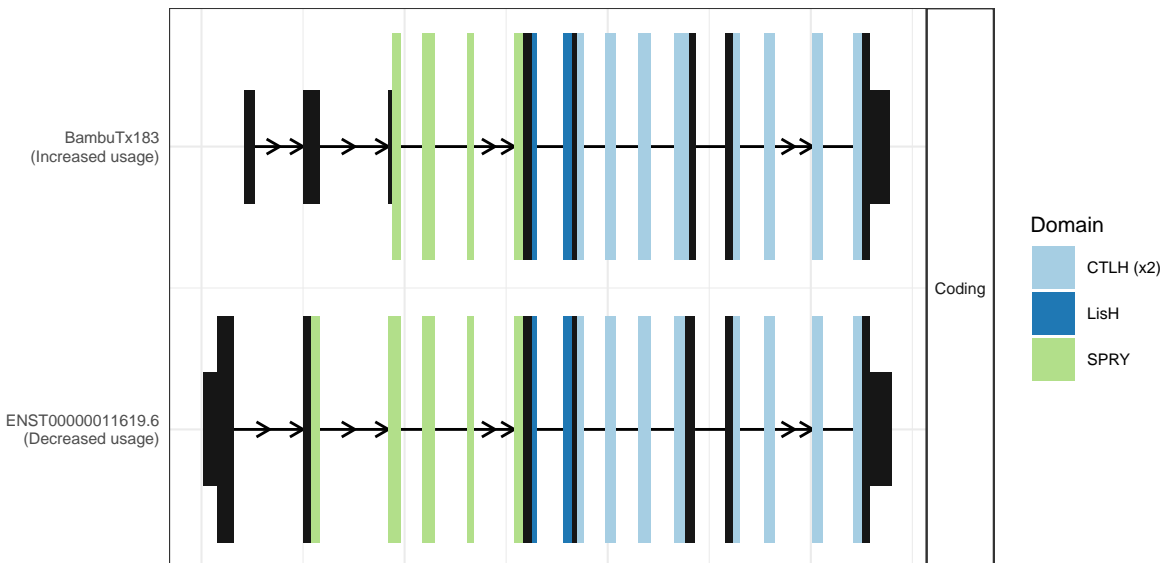


zf-A20 (x4)

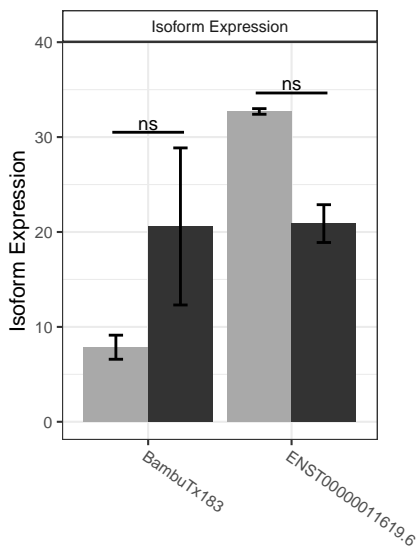


zf-A20 (x5)

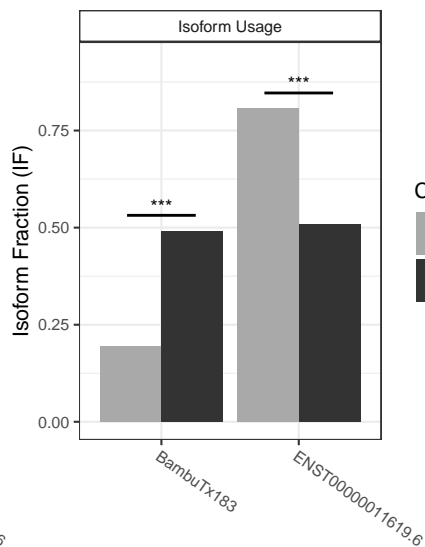
The isoform switch in RANBP9 (FT vs RGC)



Condition



Isoform






Isoform

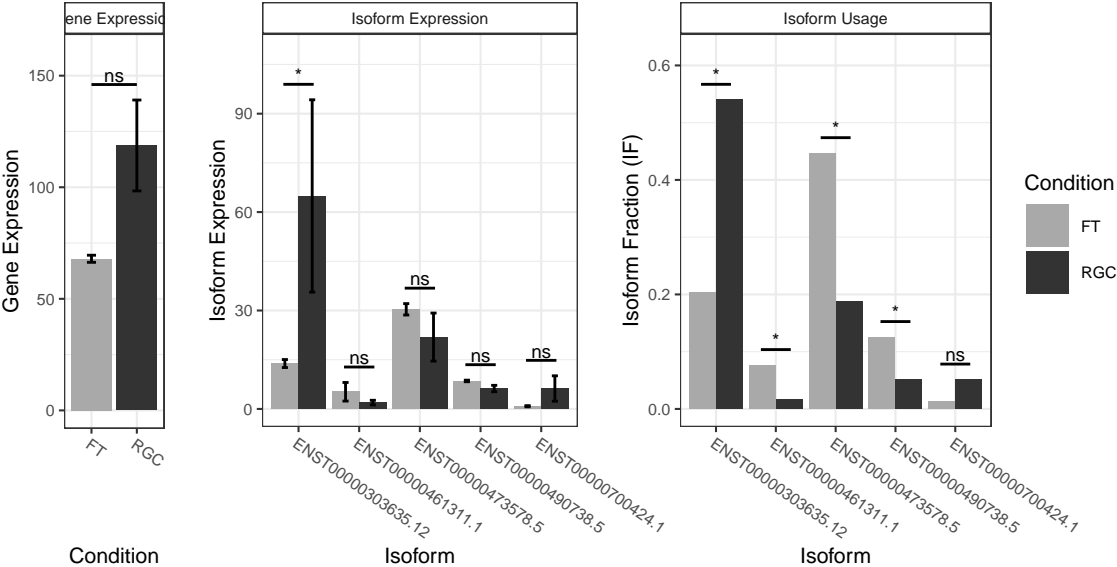
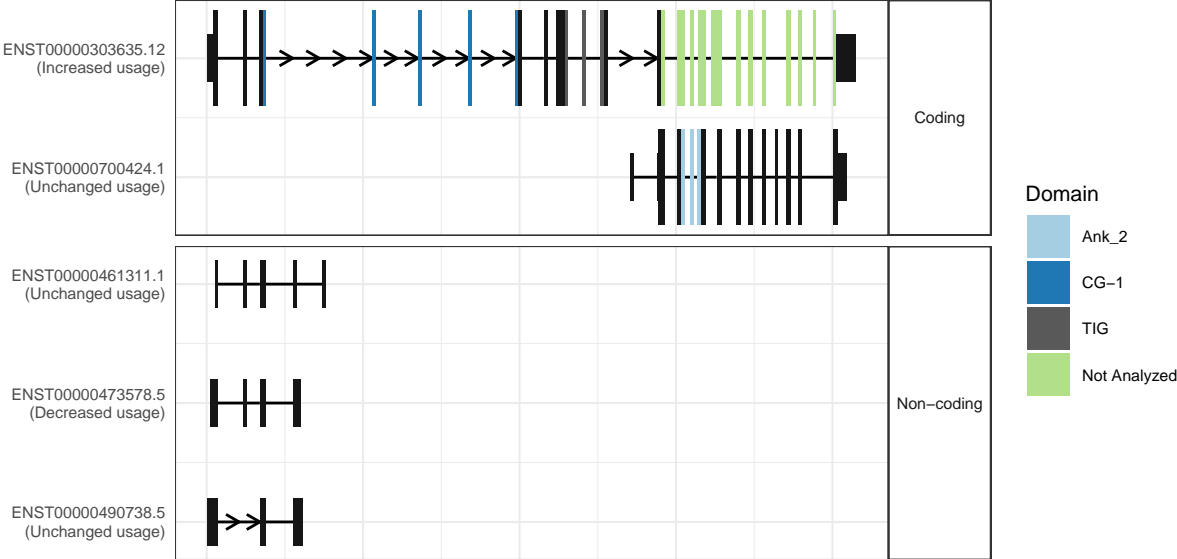
Condition



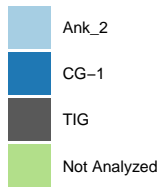
Domain

	CTLH (x2)
	LisH
	SPRY

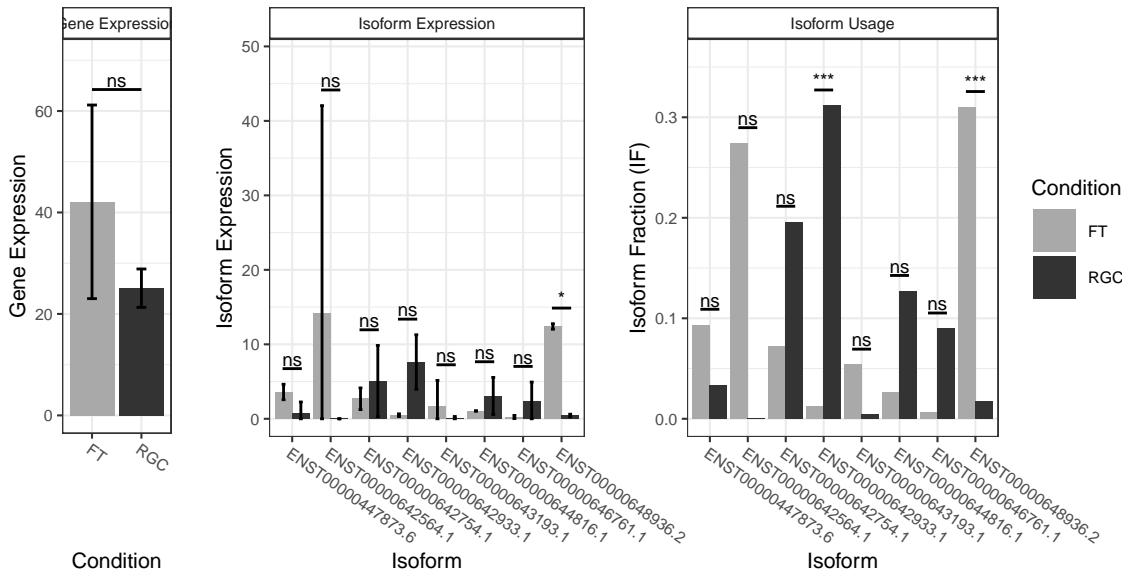
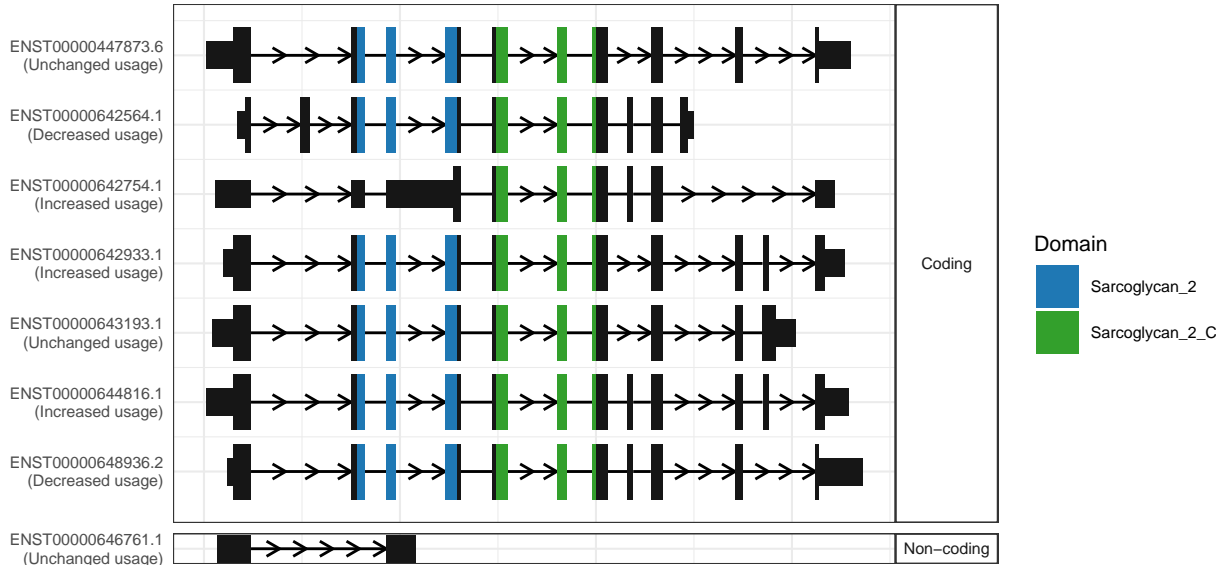
The isoform switch in CAMTA1 (FT vs RGC)



Domain



The isoform switch in SGCE (FT vs RGC)



Domain



Sarcoglycan_2



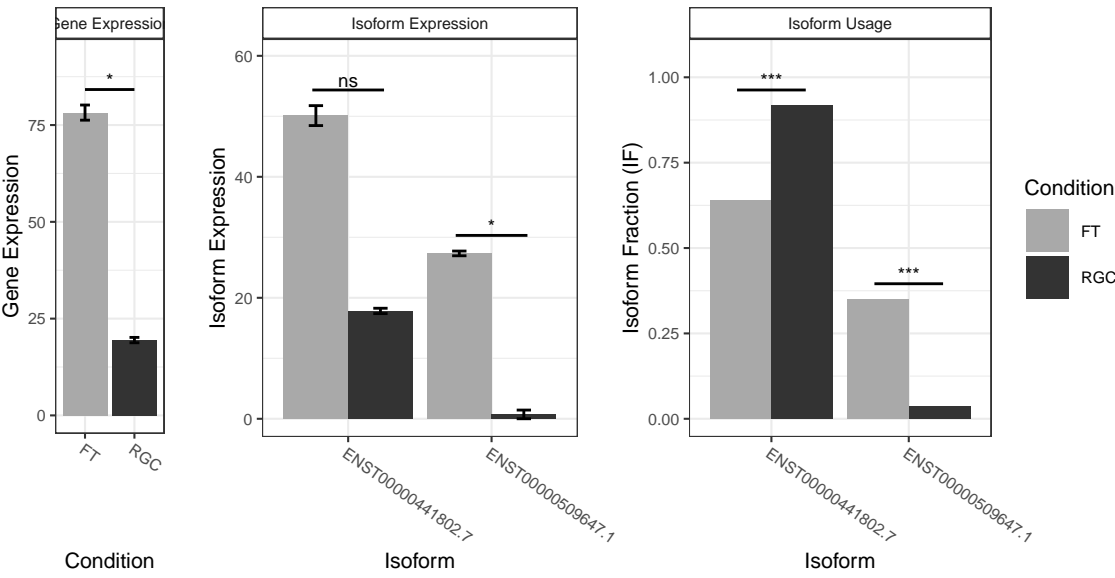
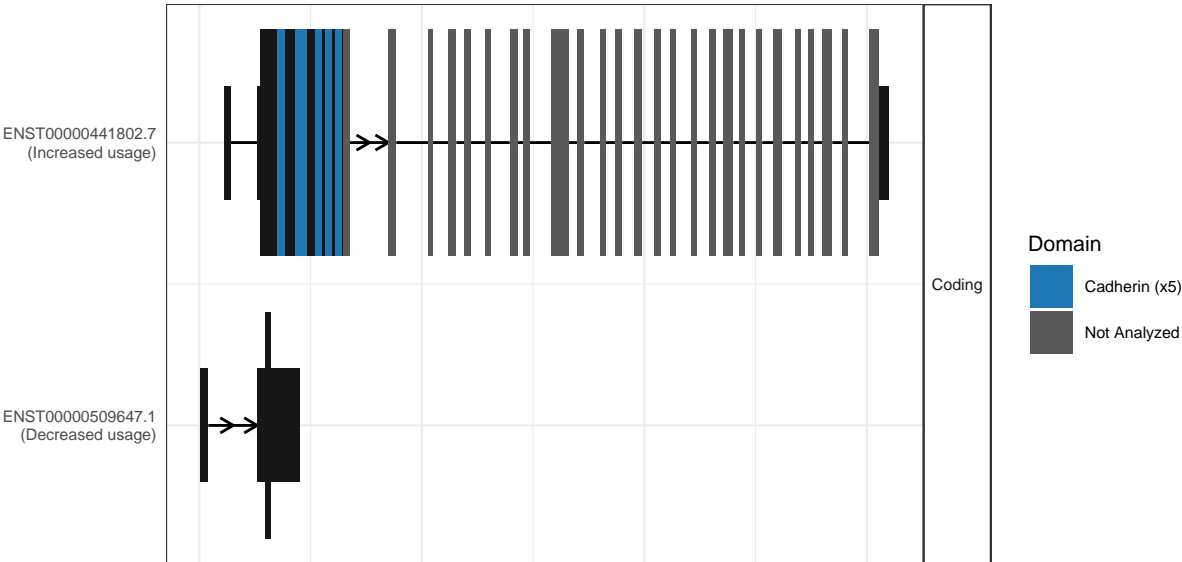
Sarcoglycan_2_C

Domain



ANAPC10

The isoform switch in FAT1 (FT vs RGC)



Domain

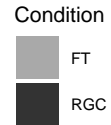
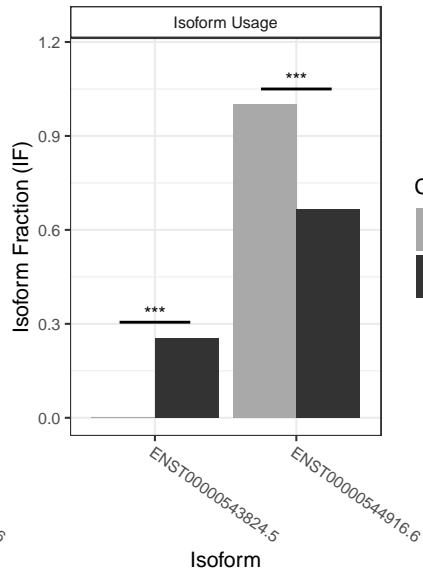
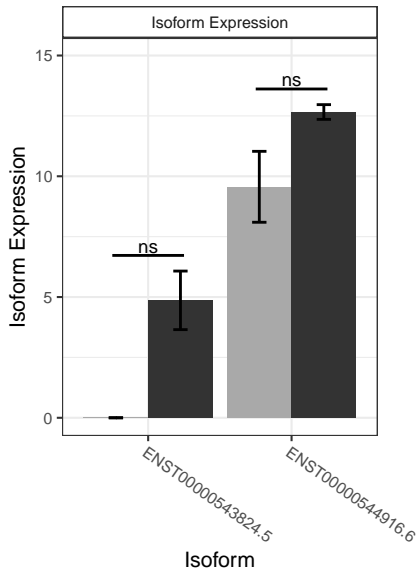
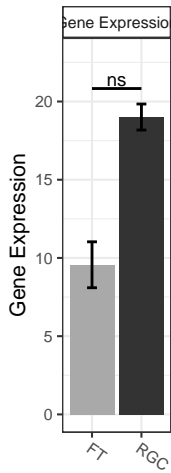
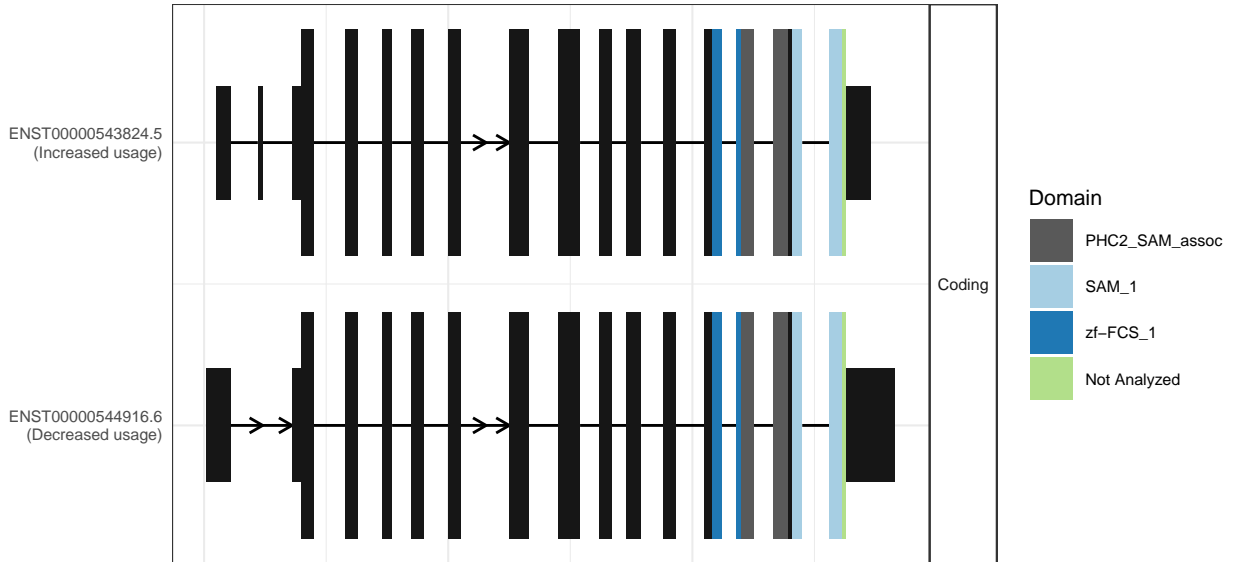


Cadherin (x5)

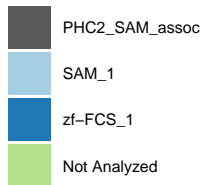


Not Analyzed

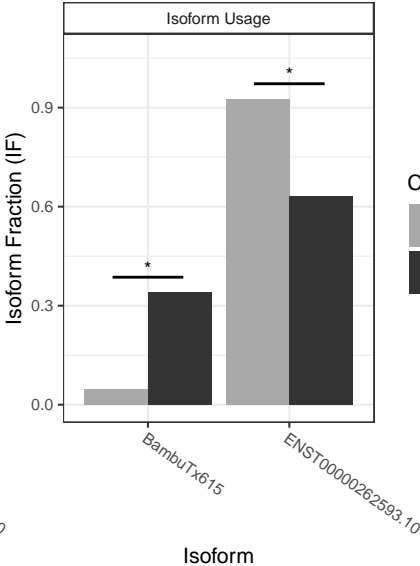
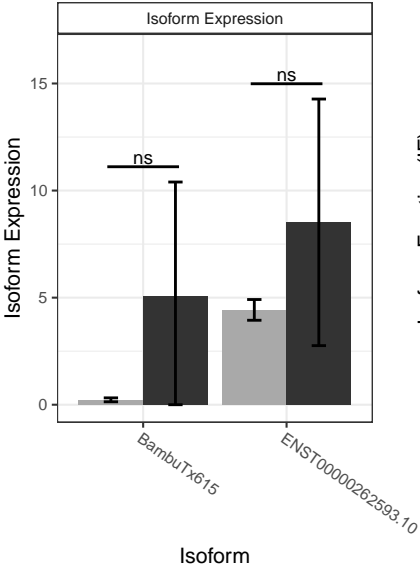
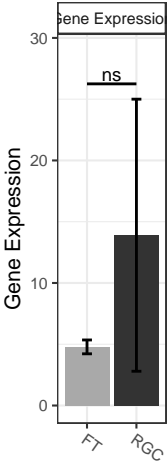
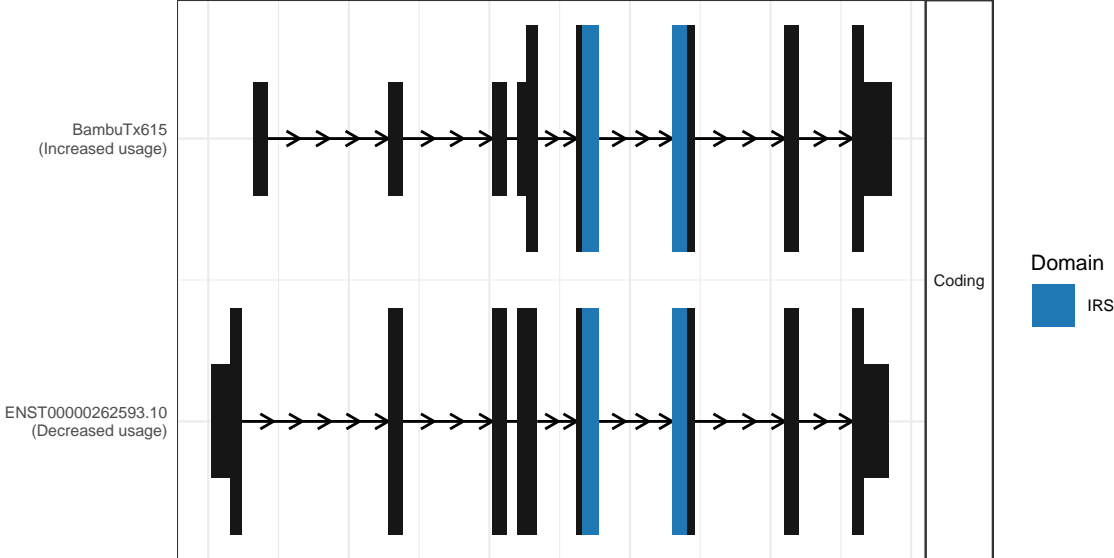
The isoform switch in PHC1 (FT vs RGC)



Domain



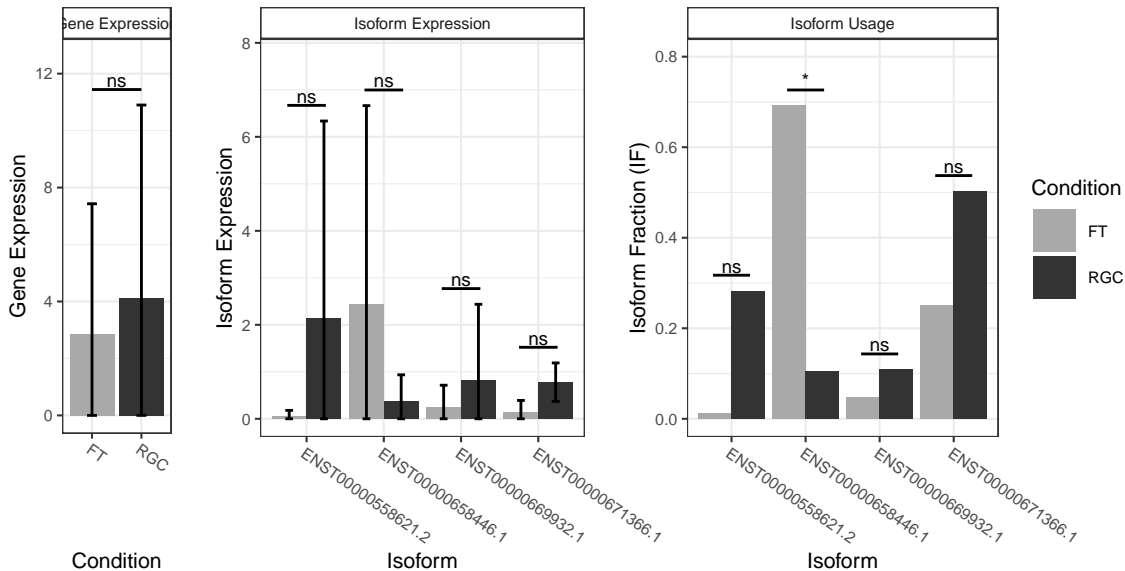
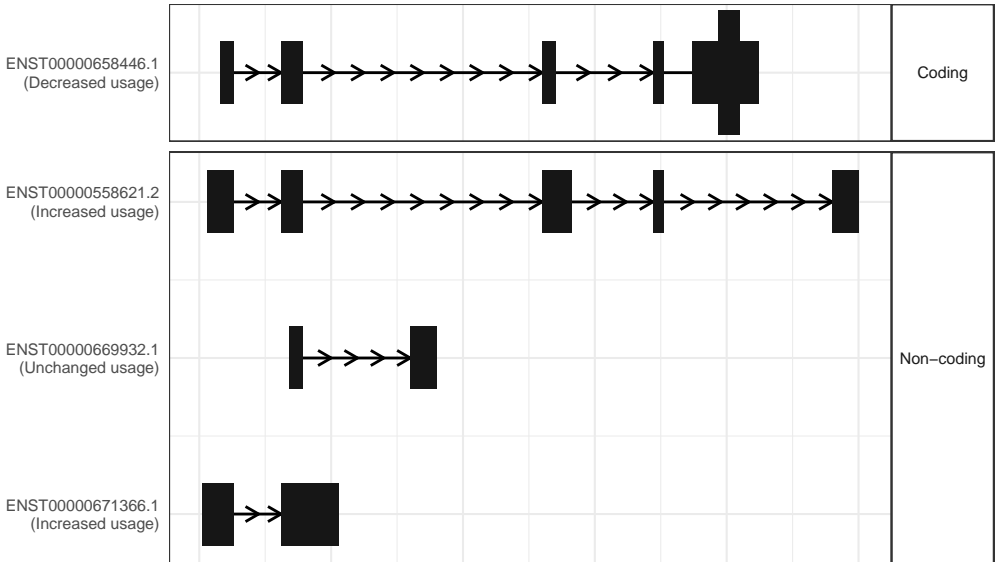
The isoform switch in DOK5 (FT vs RGC)



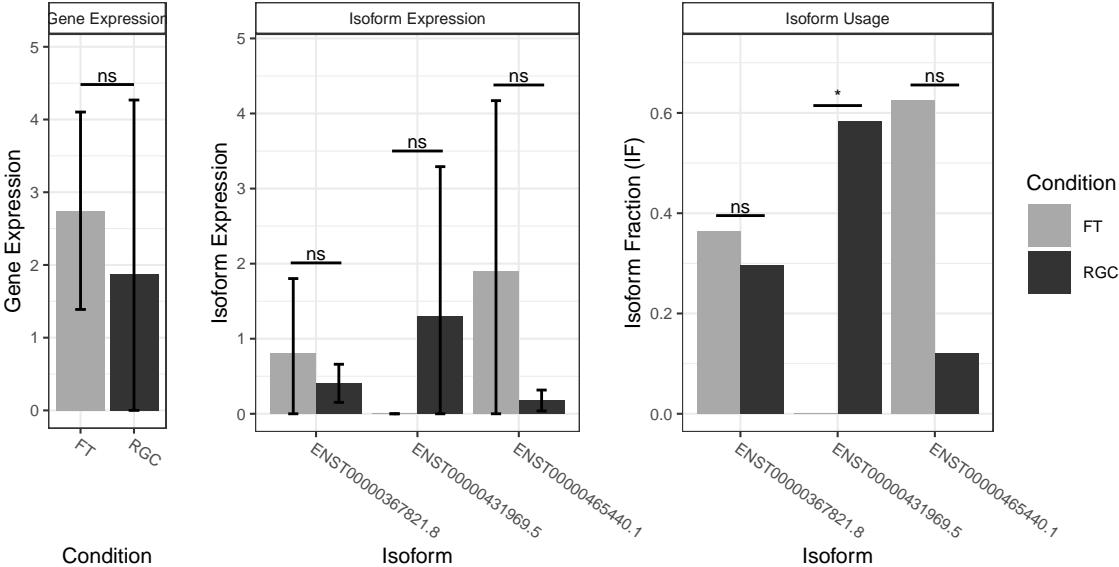
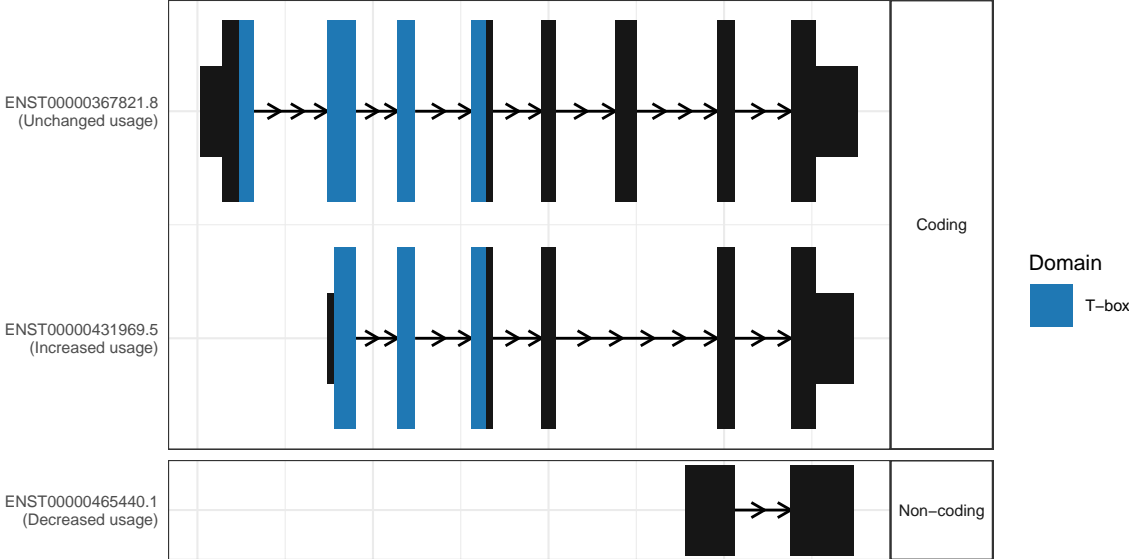
Domain



The isoform switch in unannotated gene (FT vs RGC)



The isoform switch in TBX19 (FT vs RGC)

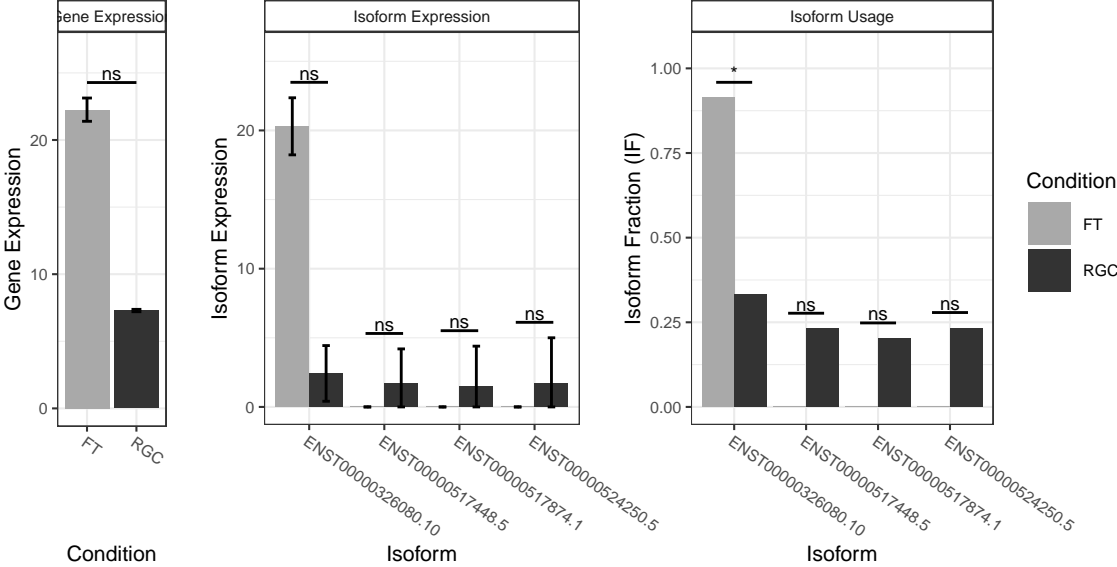
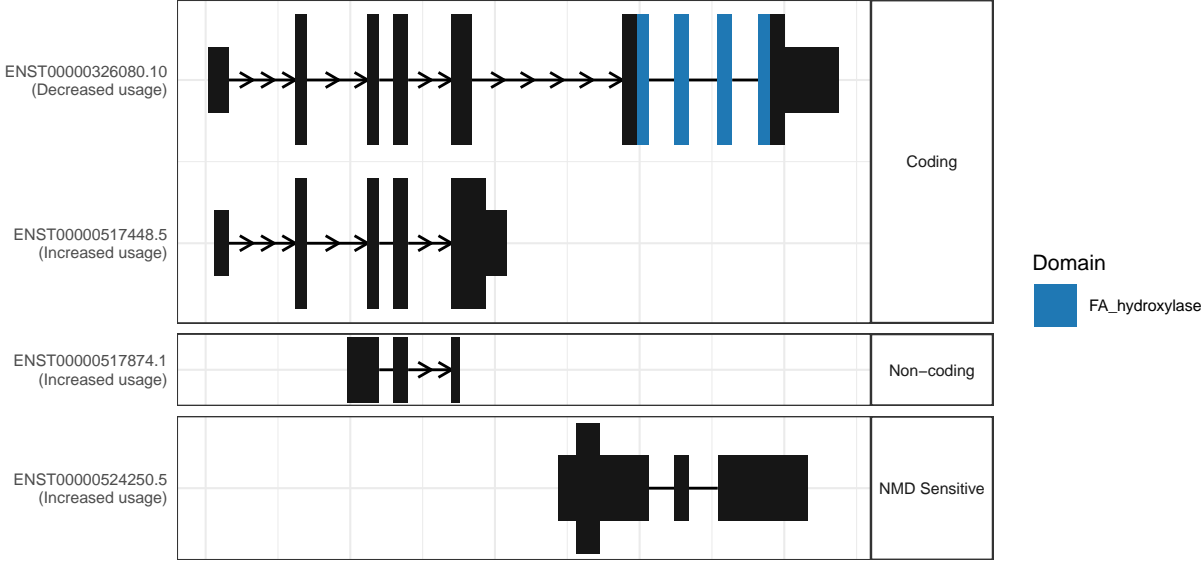


Domain



T-box

The isoform switch in FAXDC2 (FT vs RGC)

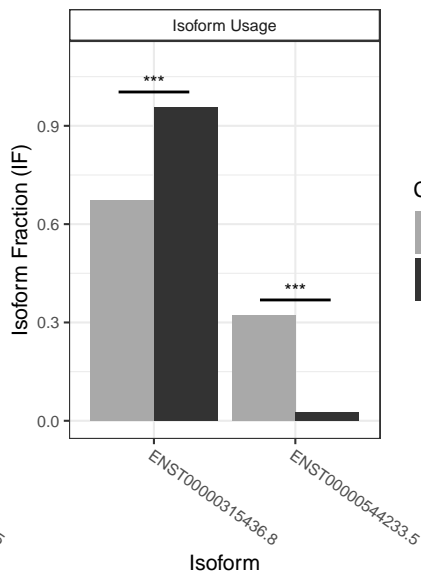
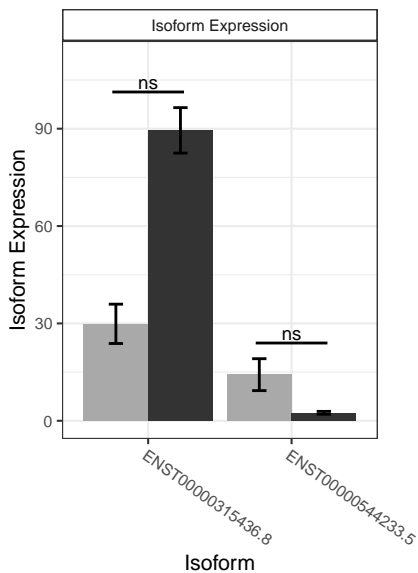
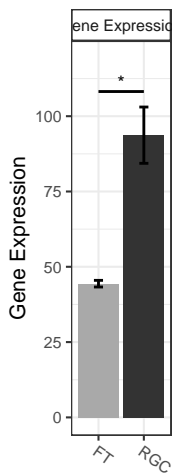
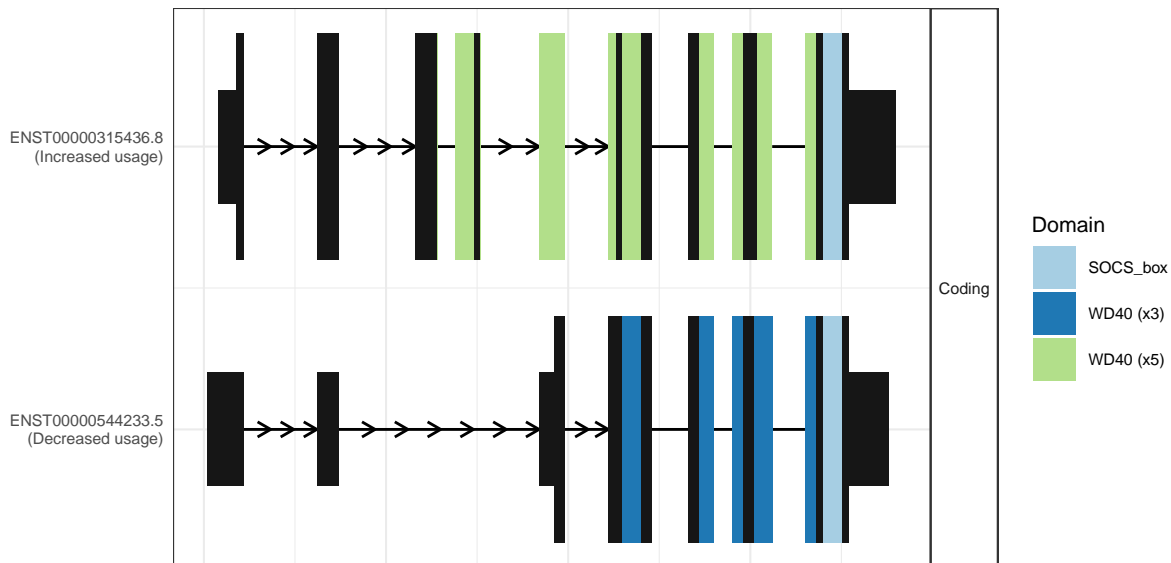


Domain






FA_hydroxylase

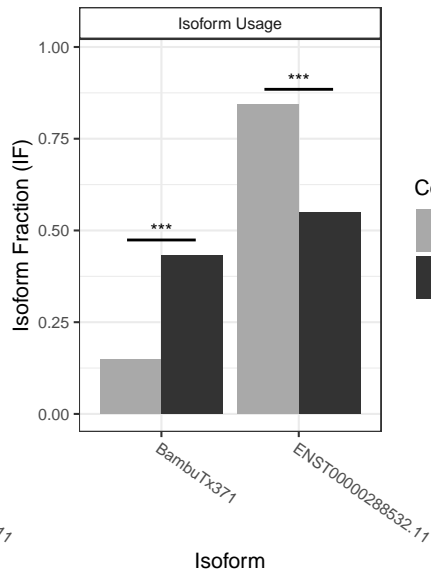
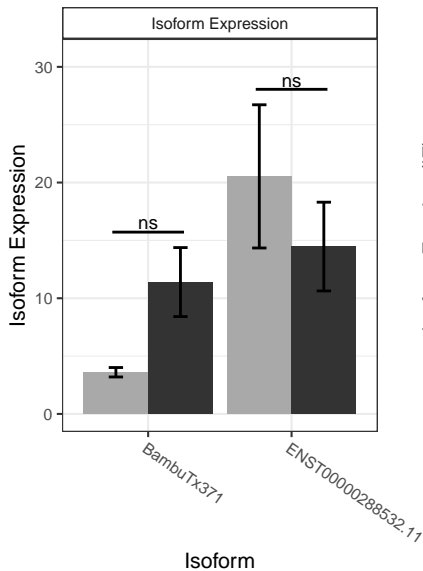
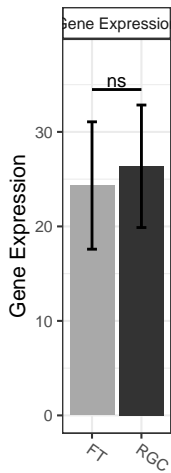
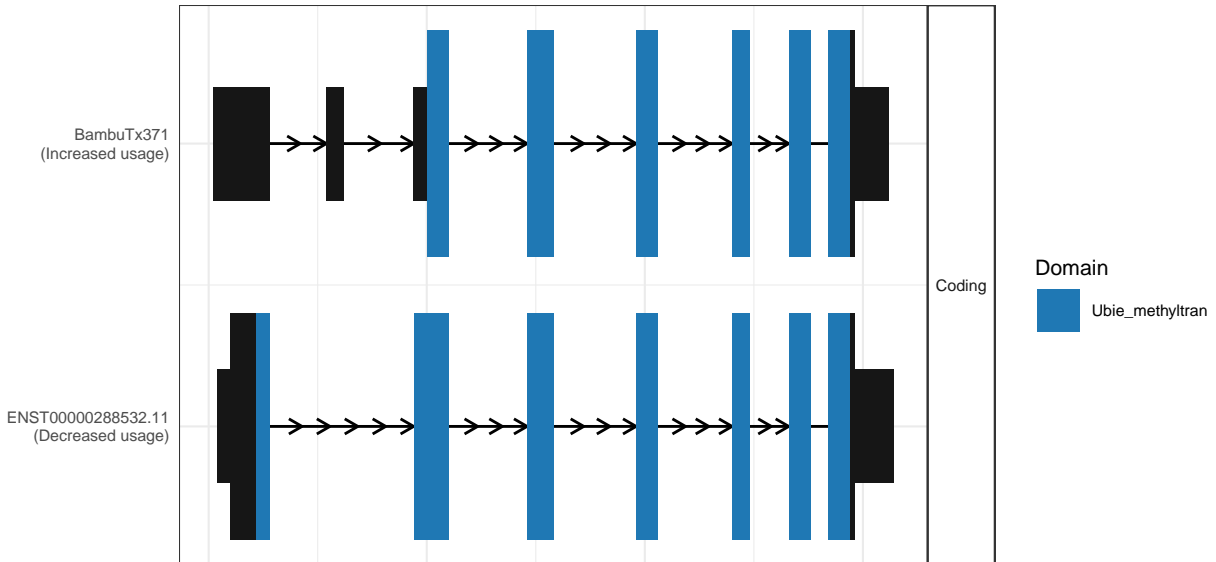
The isoform switch in WSB2 (FT vs RGC)



Domain

	SOCS_box
	WD40 (x3)
	WD40 (x5)

The isoform switch in COQ5 (FT vs RGC)

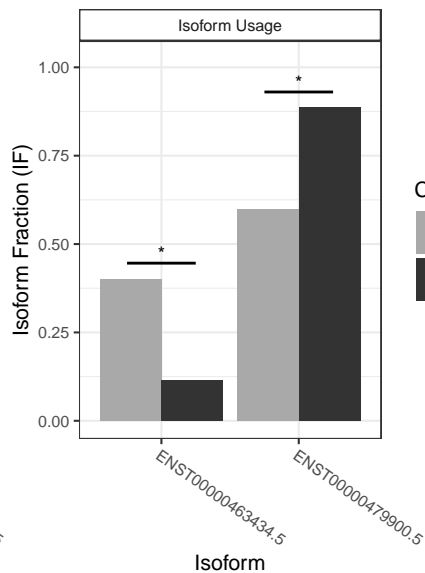
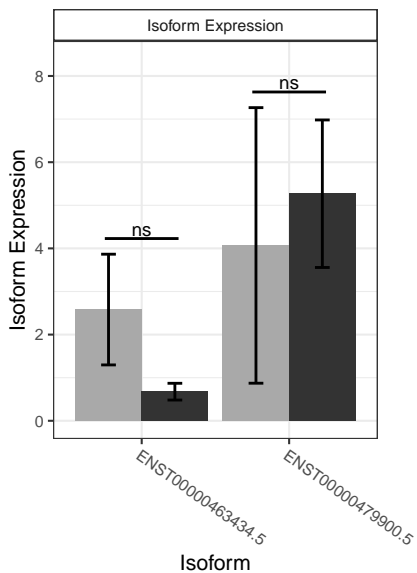
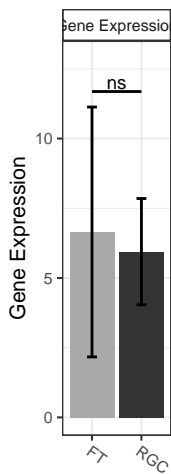
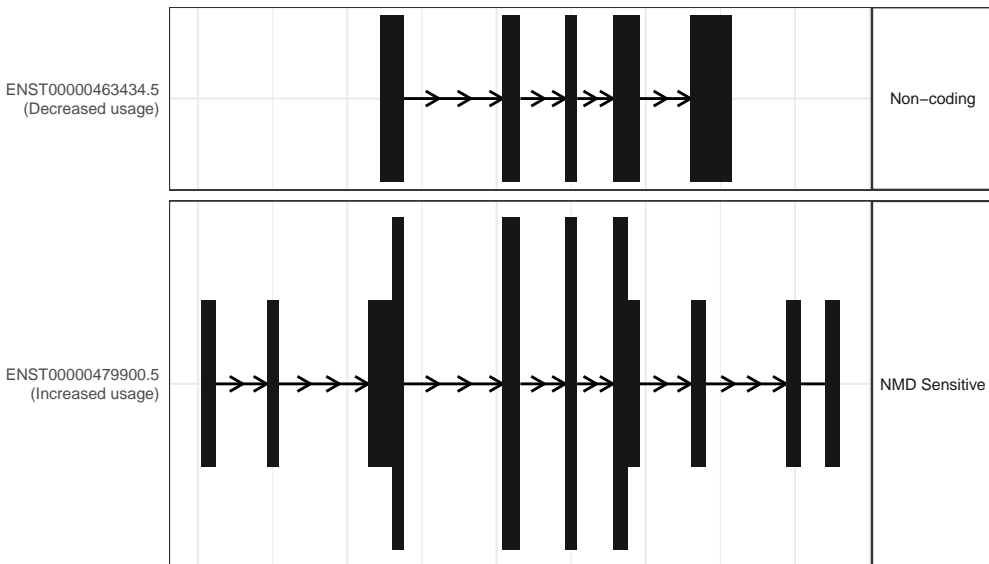


Domain



Ubie_methyltran

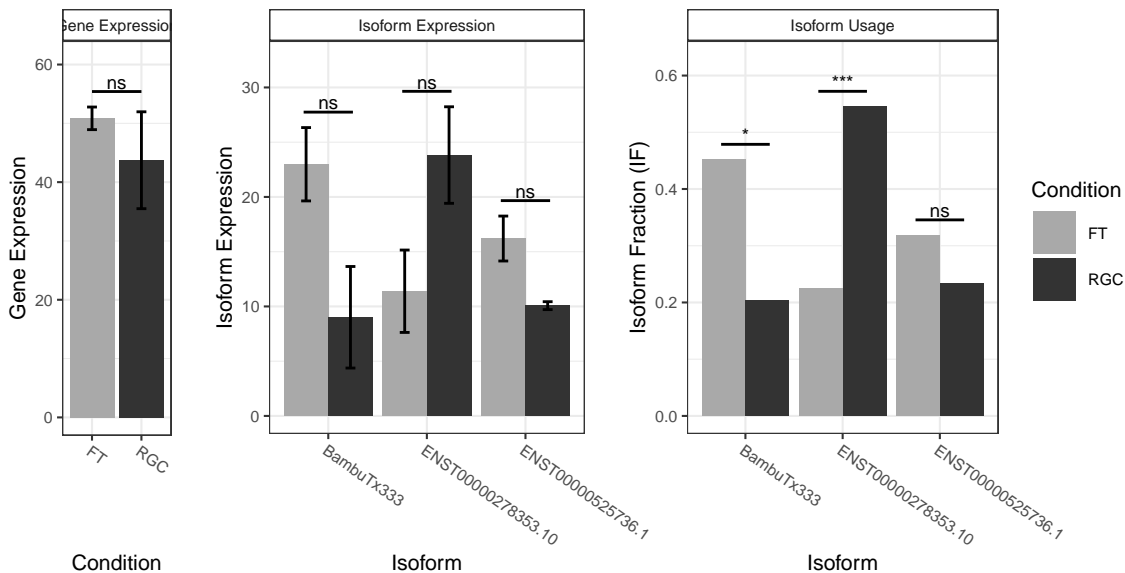
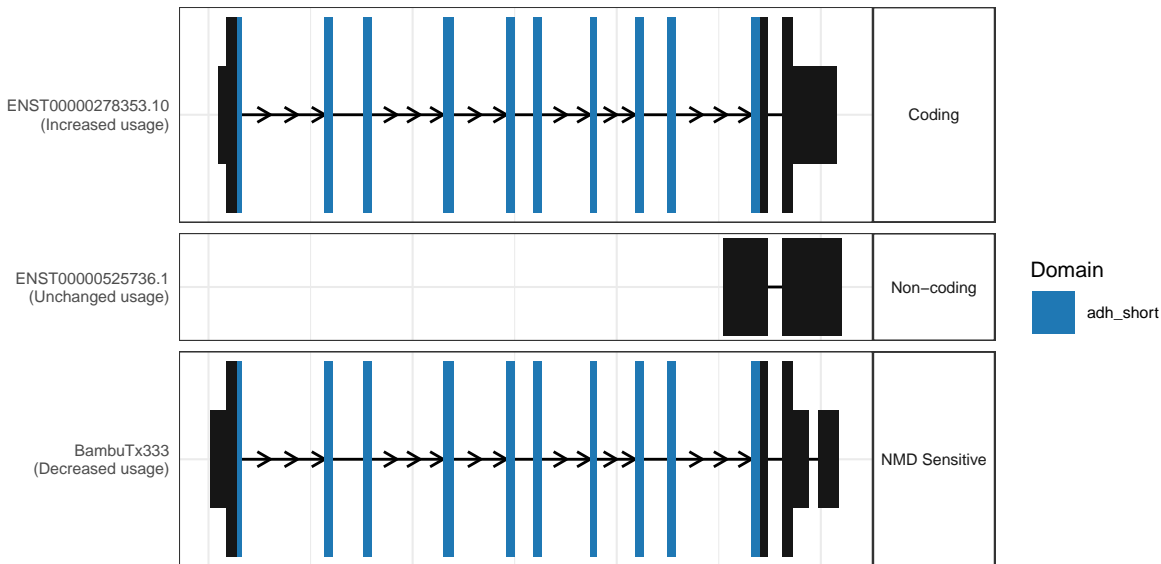
The isoform switch in (FT vs RGC)



Condition



The isoform switch in HSD17B12 (FT vs RGC)

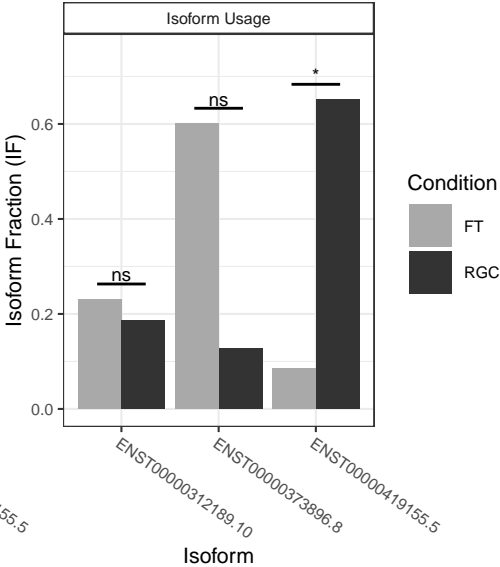
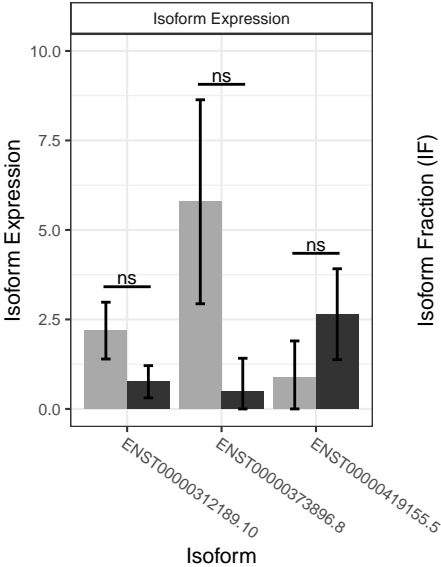
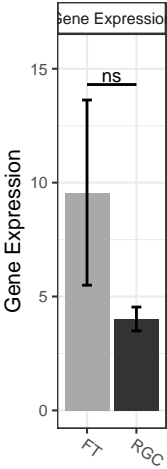
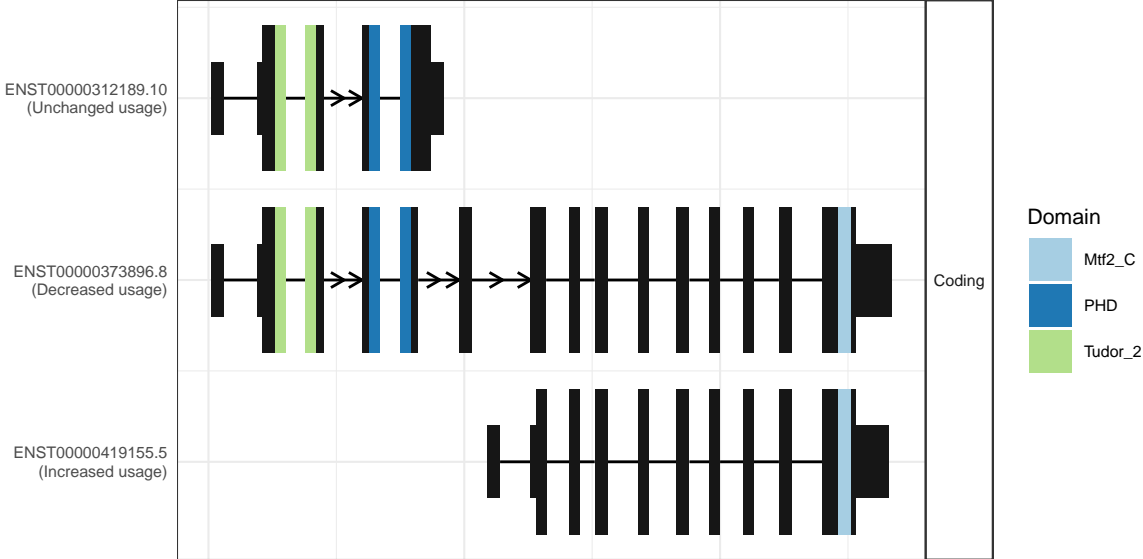


Domain

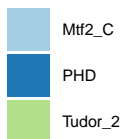


adh_short

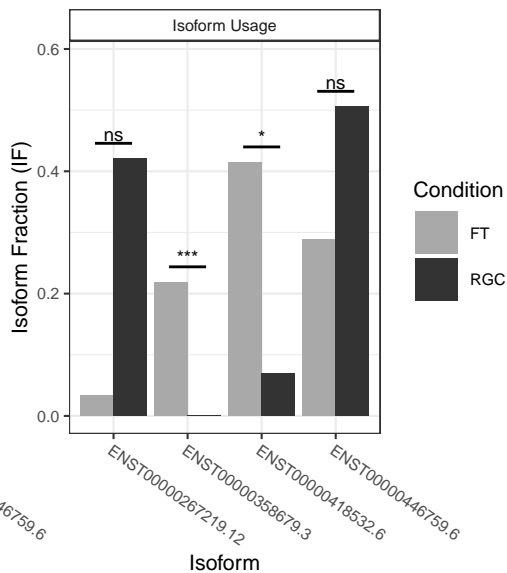
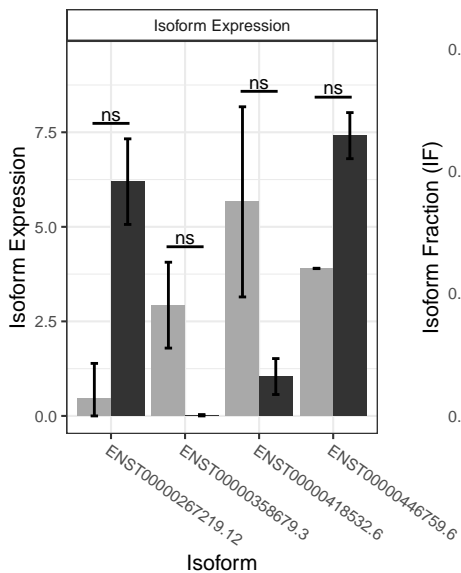
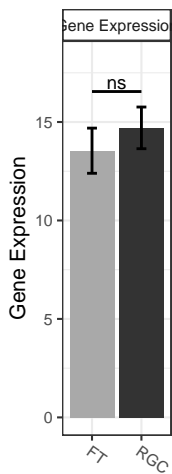
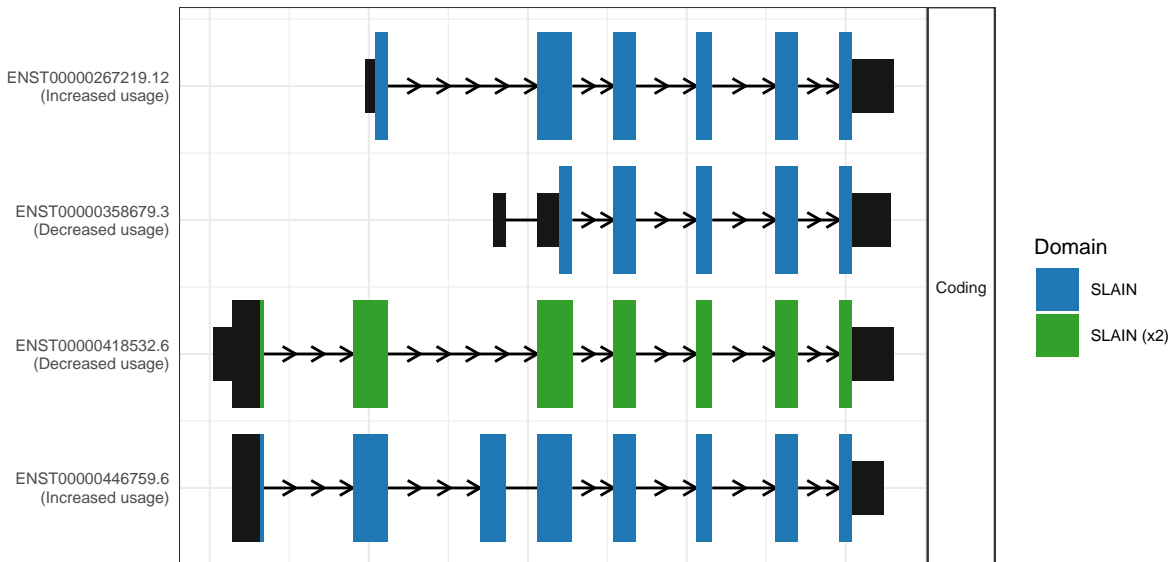
The isoform switch in PHF19 (FT vs RGC)





Domain



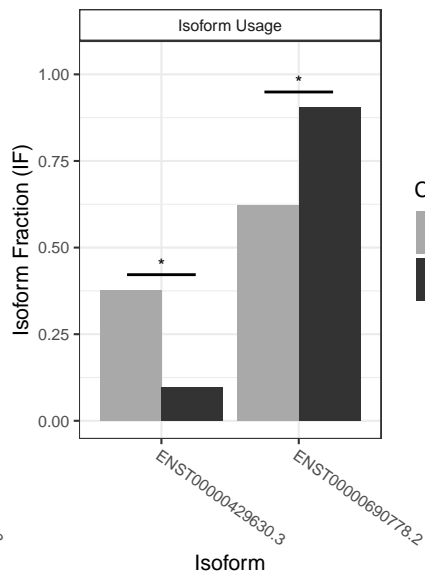
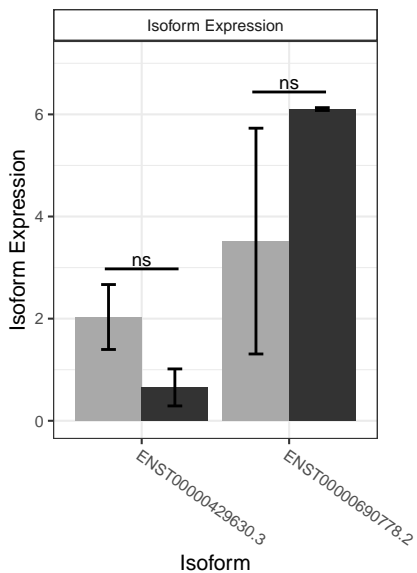
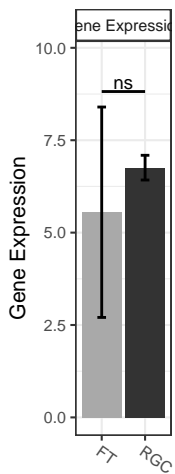
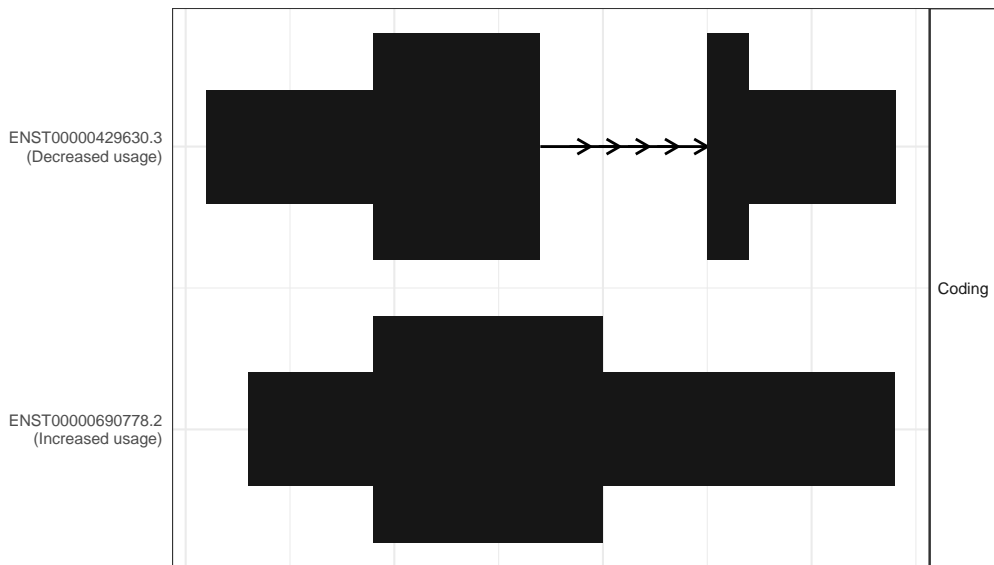
The isoform switch in SLAIN1 (FT vs RGC)



Domain

	SLAIN
	SLAIN (x2)

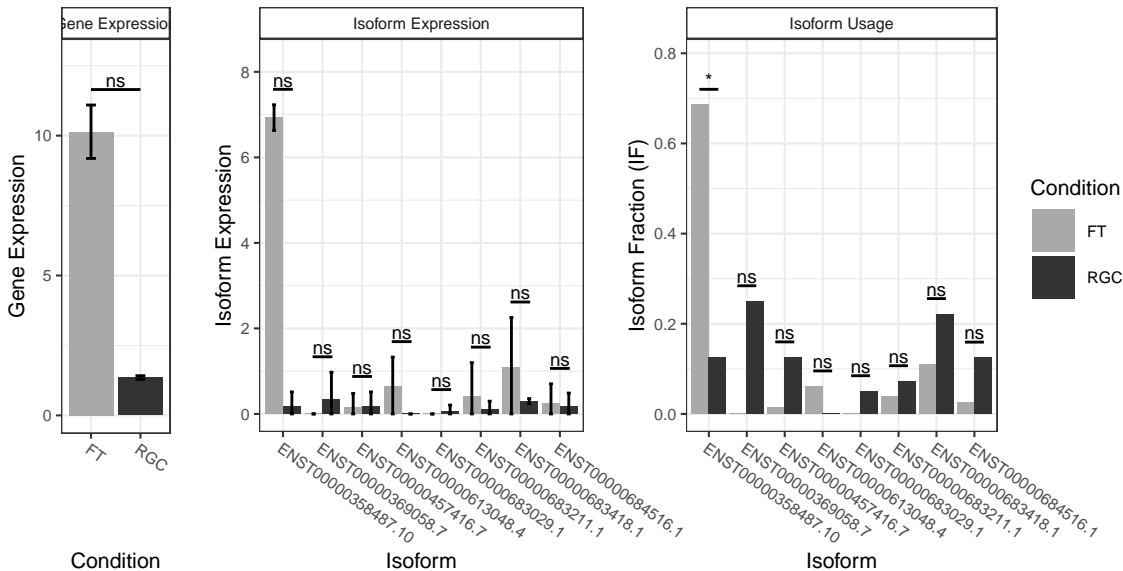
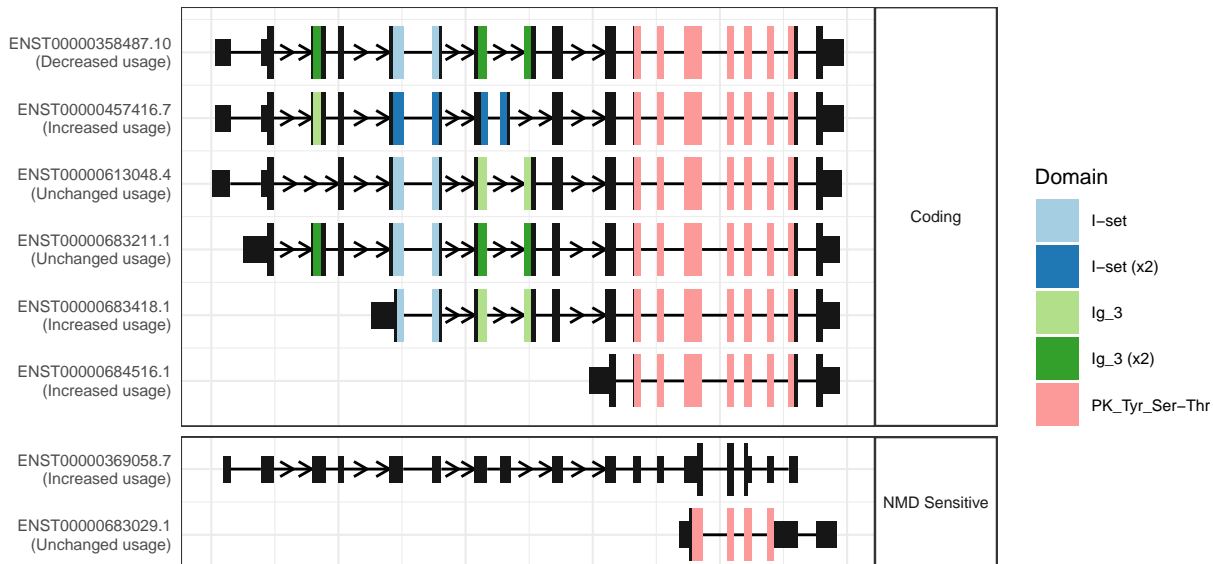
The isoform switch in FAM131B-AS2 (FT vs RGC)






Condition



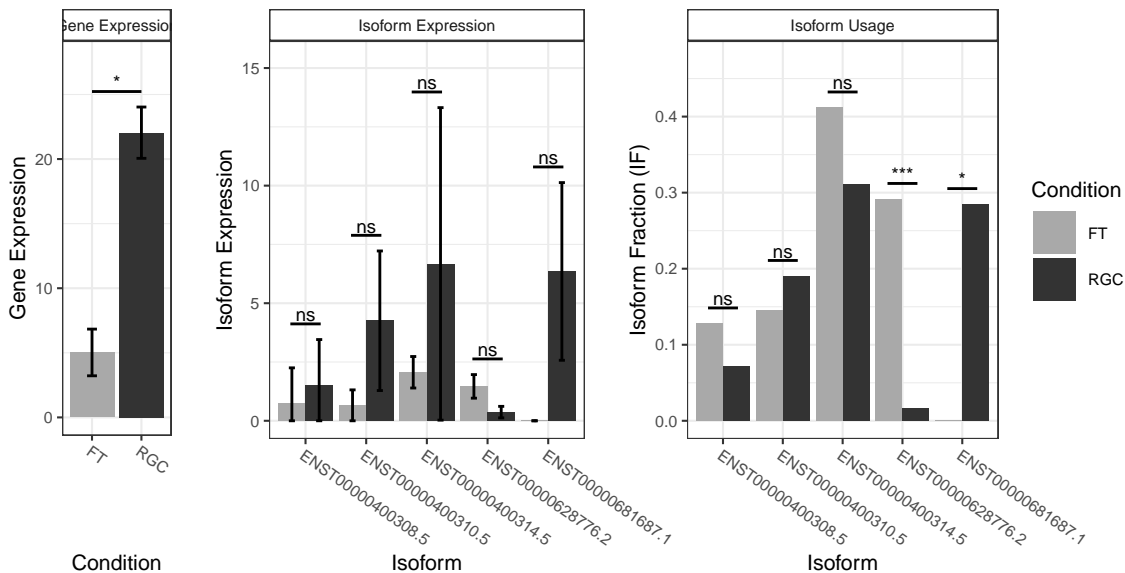
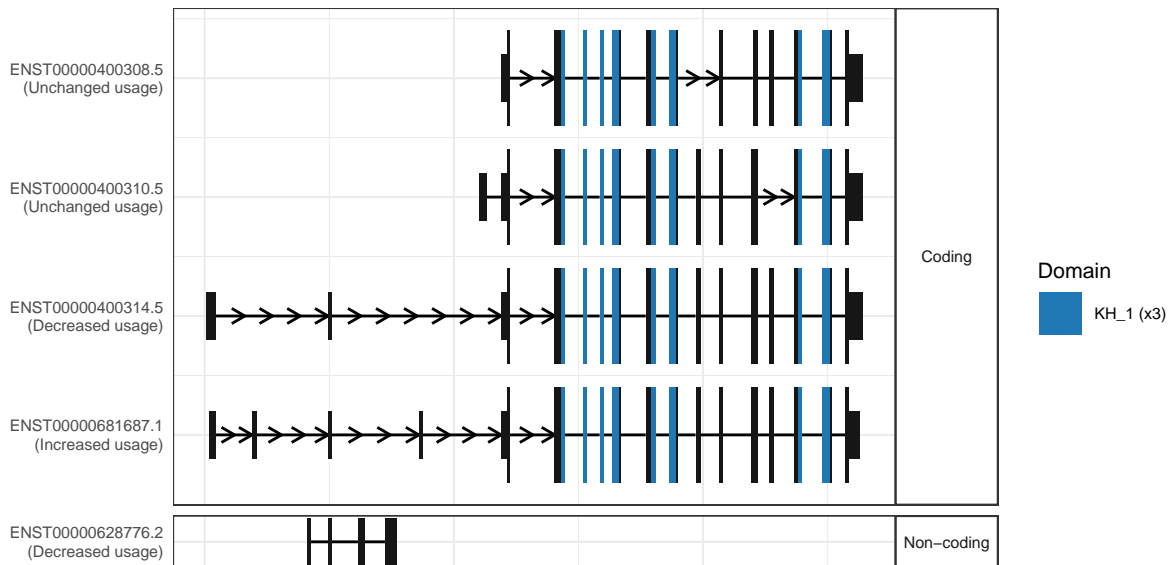
The isoform switch in FGFR2 (FT vs RGC)



Domain

	I-set
	I-set (x2)
	Ig_3
	Ig_3 (x2)
	PK_Tyr_Ser-Thr

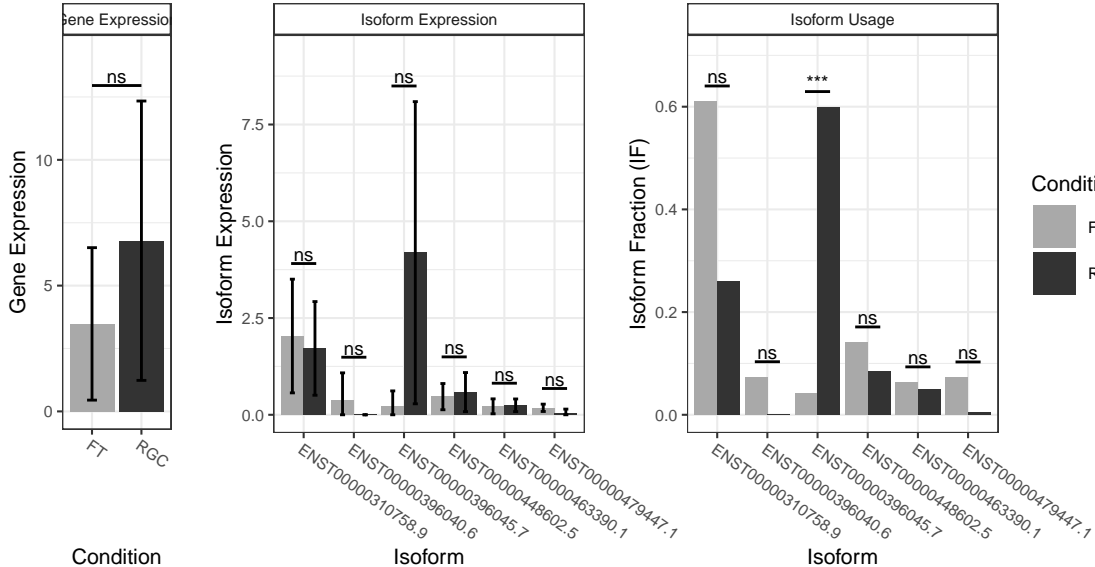
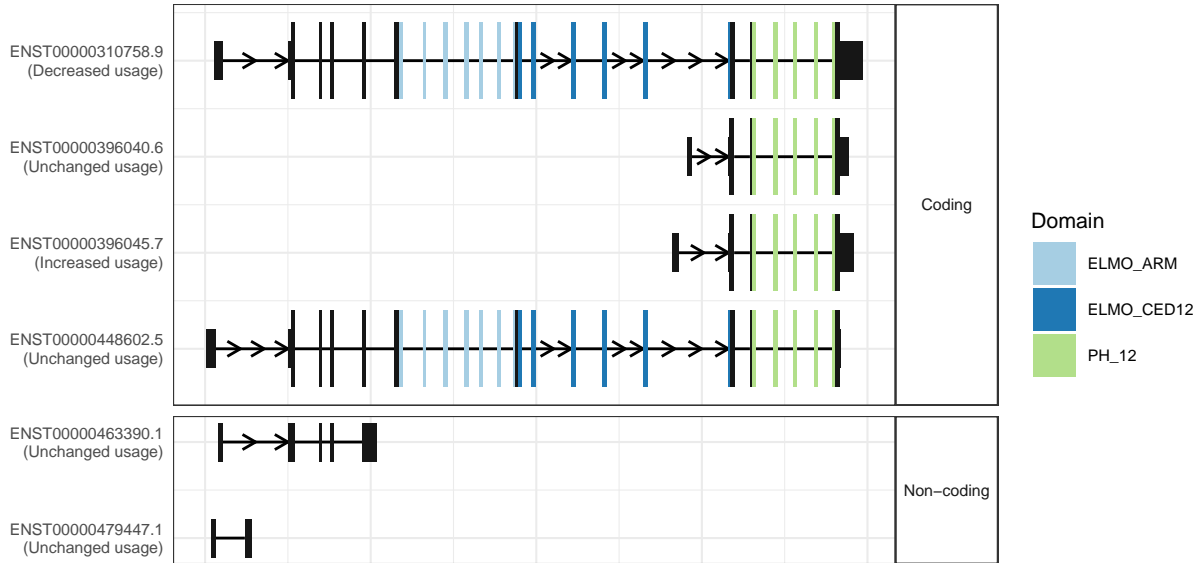
The isoform switch in PCBP3 (FT vs RGC)



Domain

 KH_1 (x3)

The isoform switch in ELMO1 (FT vs RGC)



Domain



ELMO_ARM

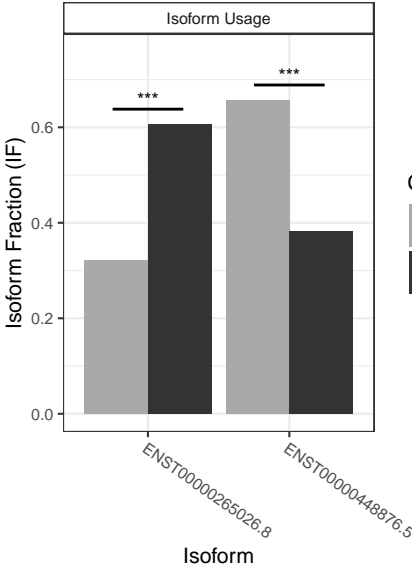
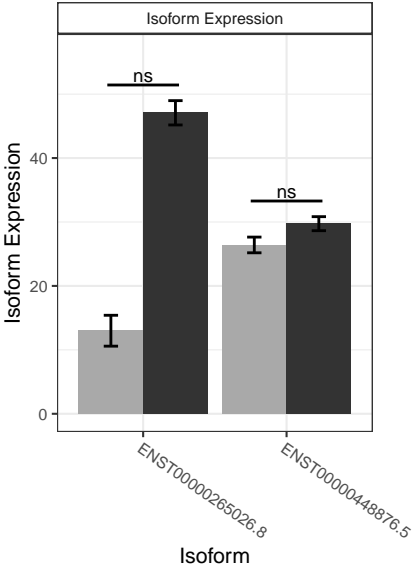
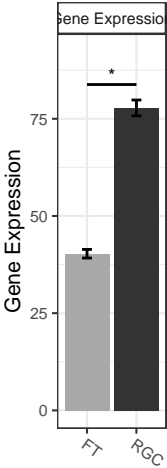
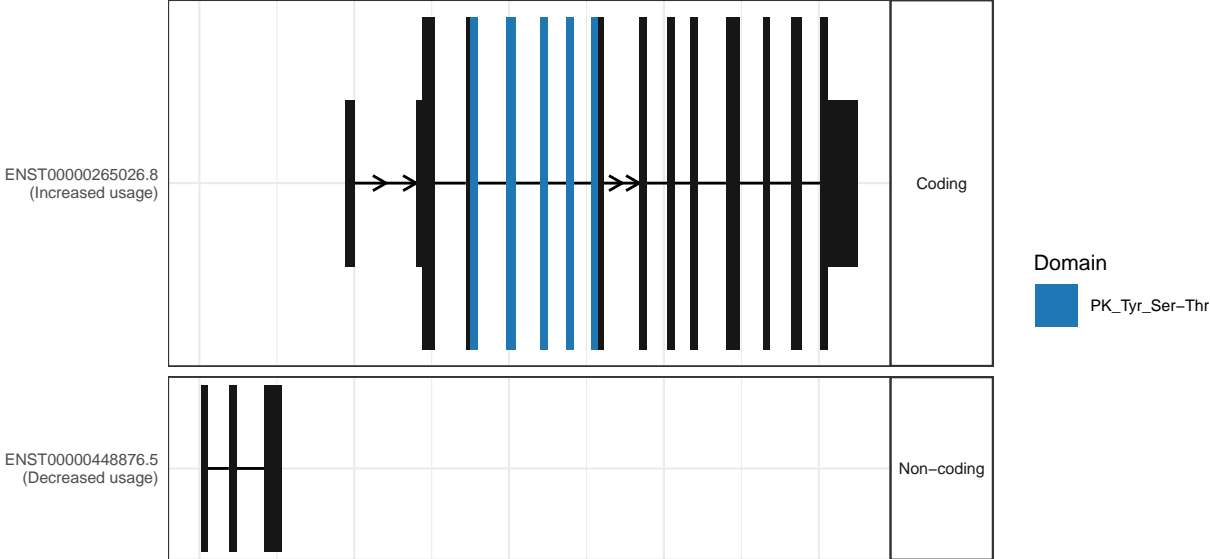


ELMO_CED12



PH_12

The isoform switch in MAP3K13 (FT vs RGC)

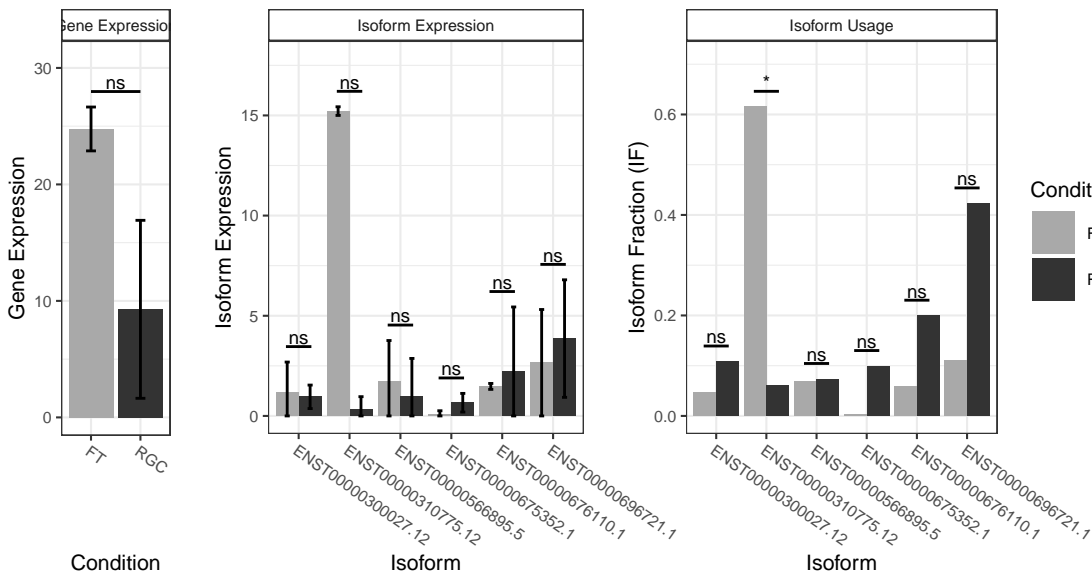
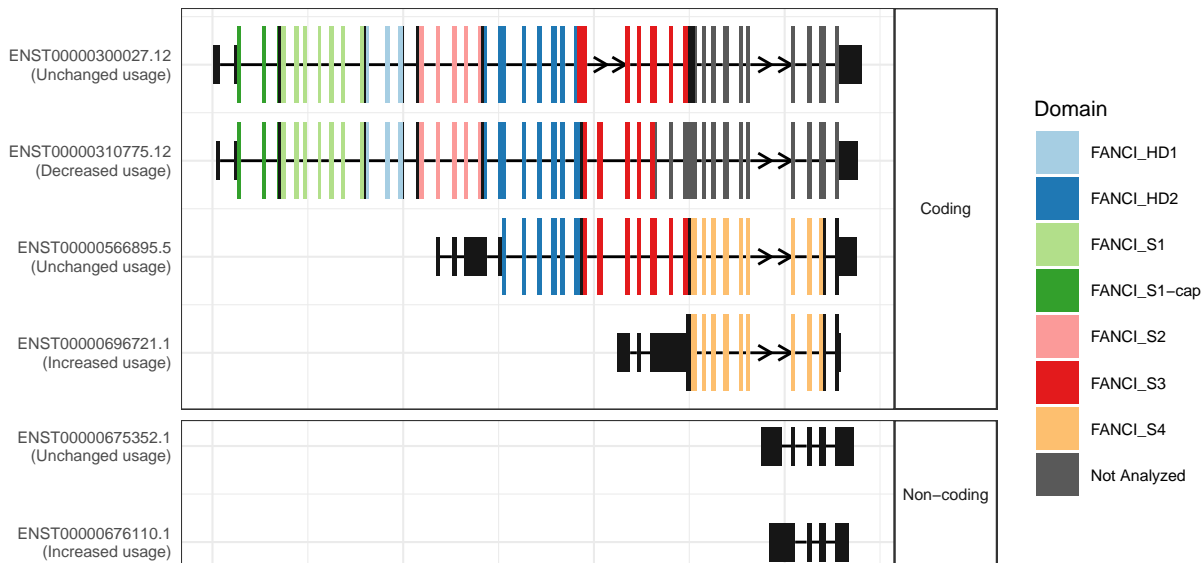


Domain



PK_Tyr_Ser-Thr

The isoform switch in FANCI (FT vs RGC)



Domain



FANCI_HD1



FANCI_HD2



FANCI_S1



FANCI_S1-cap



FANCI_S2



FANCI_S3

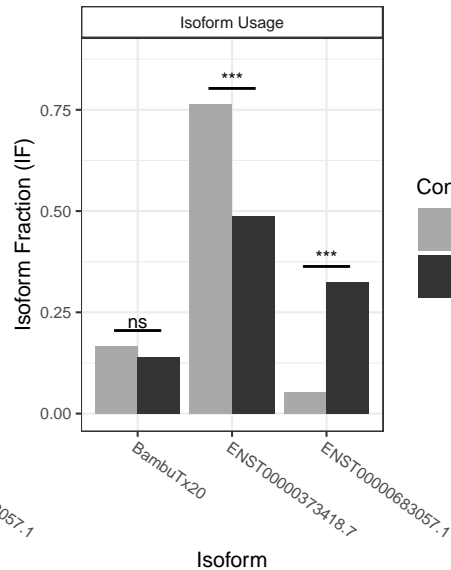
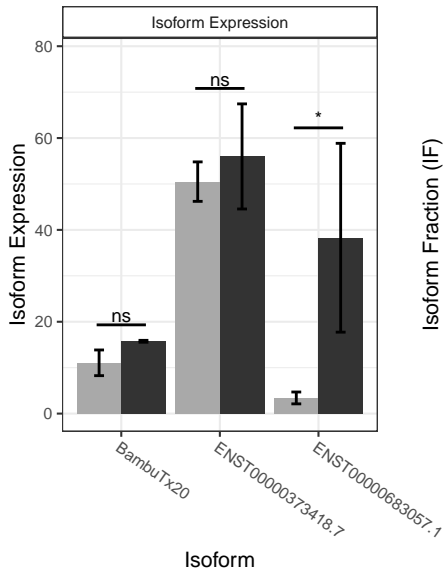
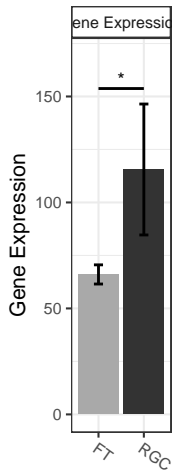
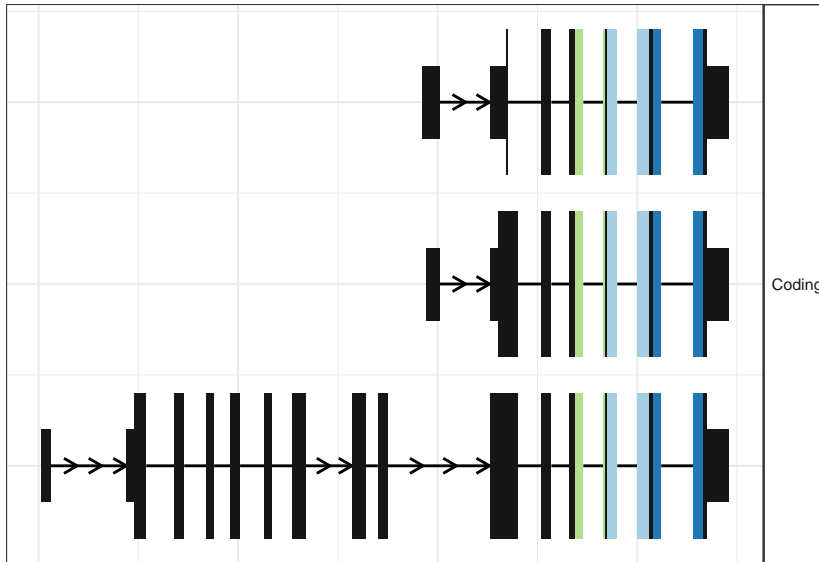


FANCI_S4






Not Analyzed

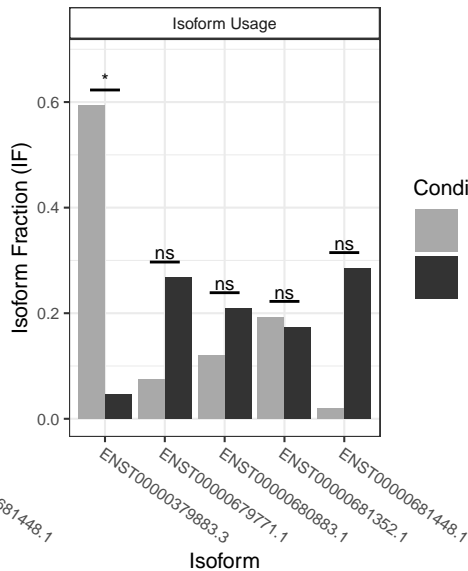
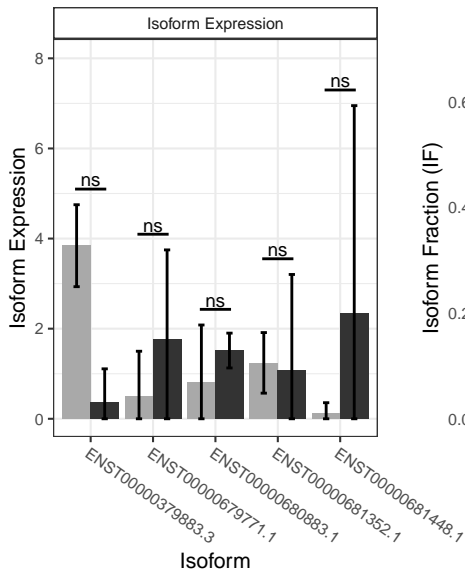
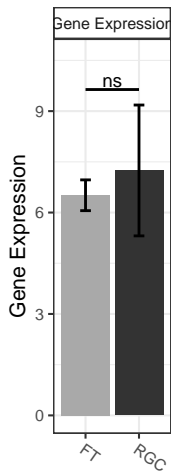
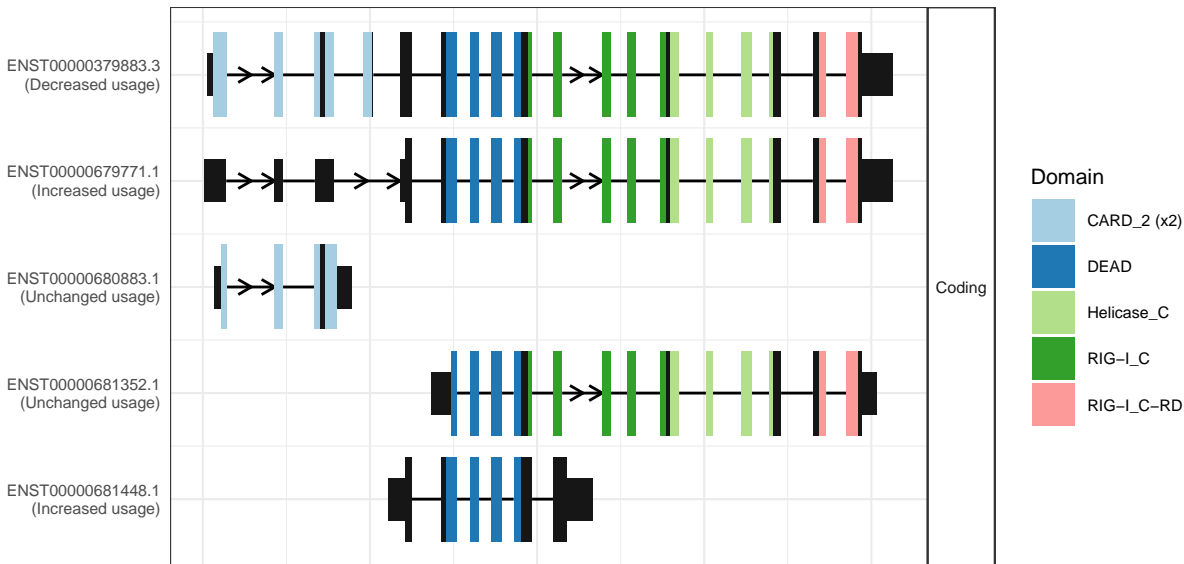
The isoform switch in PHC2 (FT vs RGC)



Domain

	PHC2_SAM_assoc
	SAM_1
	zf-FCS_1

The isoform switch in RIGI (FT vs RGC)



Domain



CARD_2 (x2)



DEAD



Helicase_C

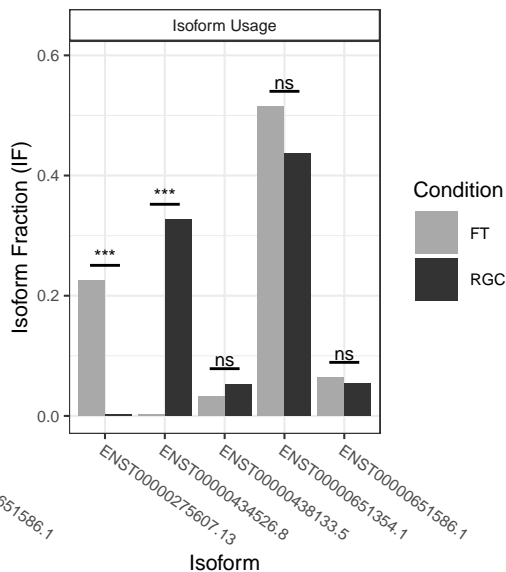
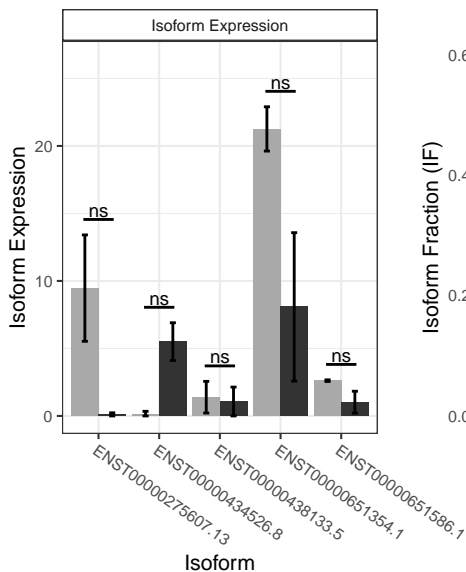
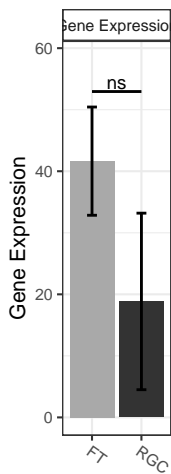
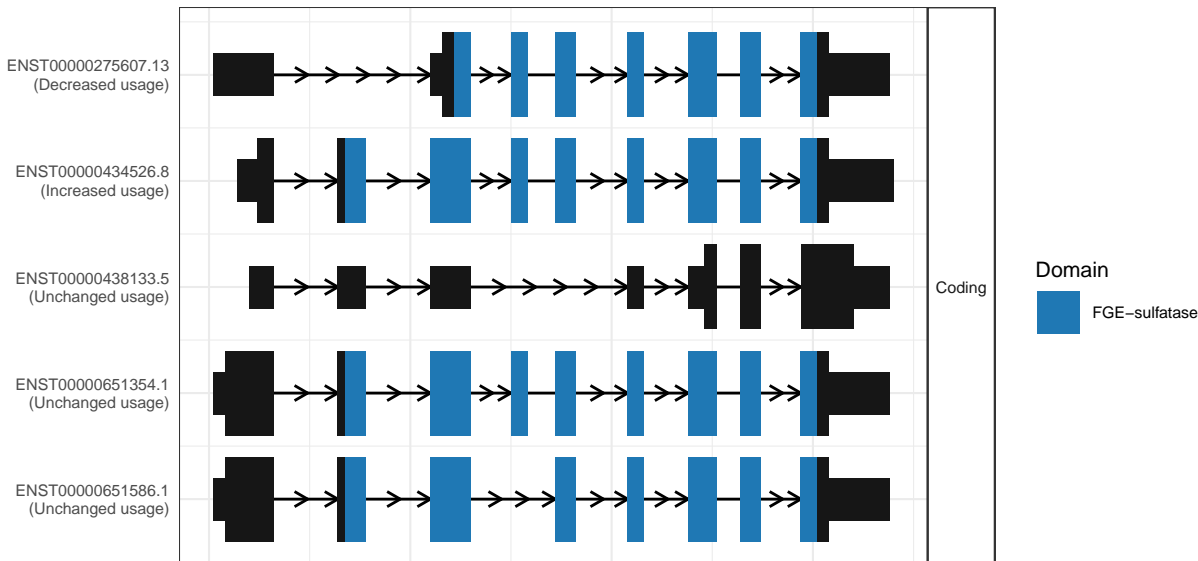


RIG-I_C



RIG-I_C-RD

The isoform switch in SUMF2 (FT vs RGC)

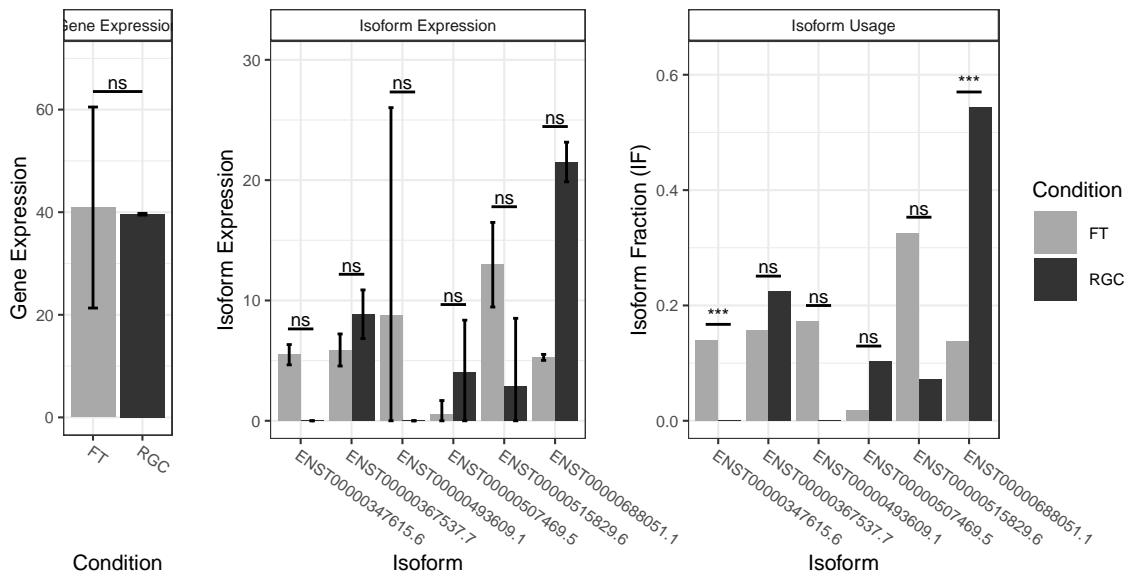
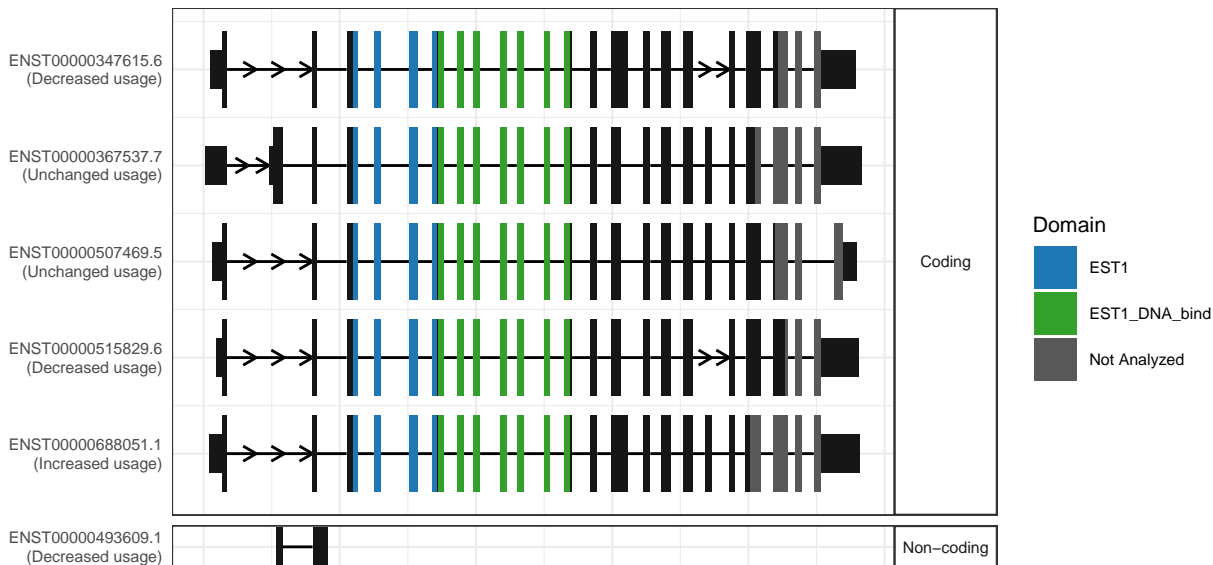


Domain






FGE-sulfatase

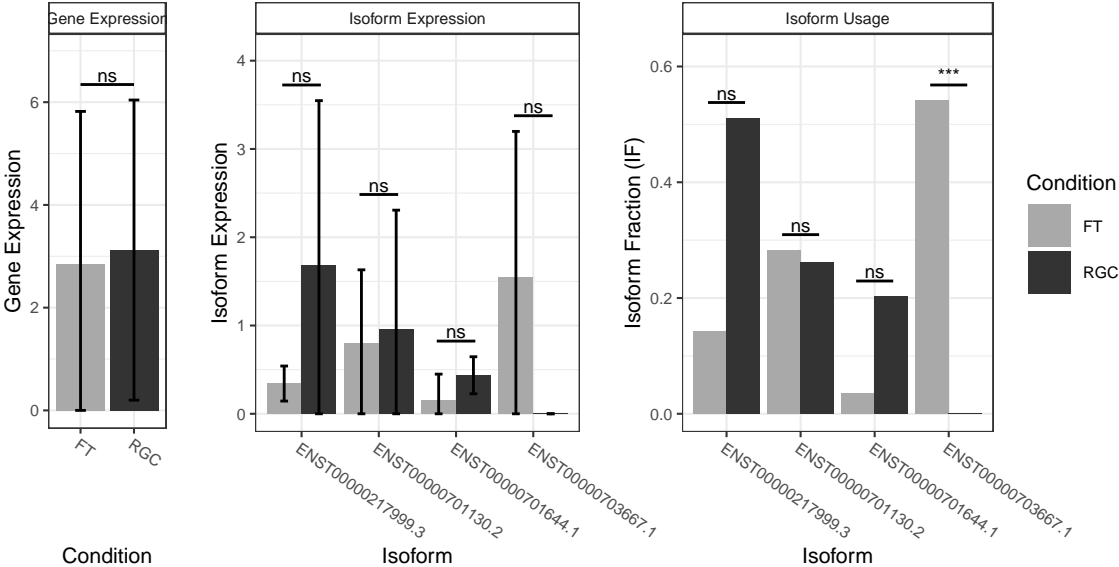
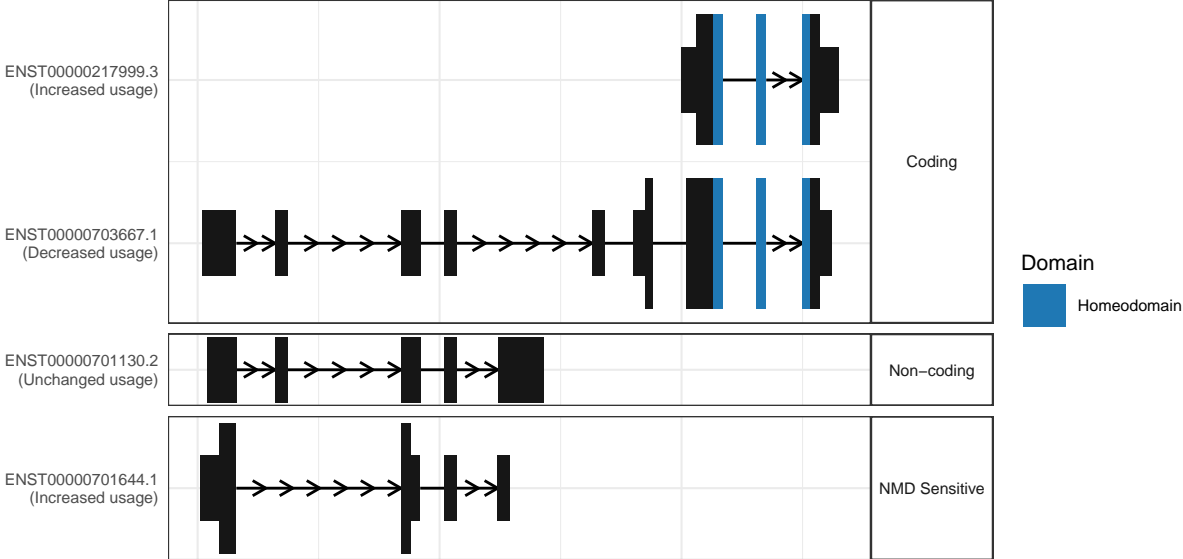
The isoform switch in SMG7 (FT vs RGC)



Domain

	EST1
	EST1_DNA_bind
	Not Analyzed

The isoform switch in RHOXF1(FT vs RGC)

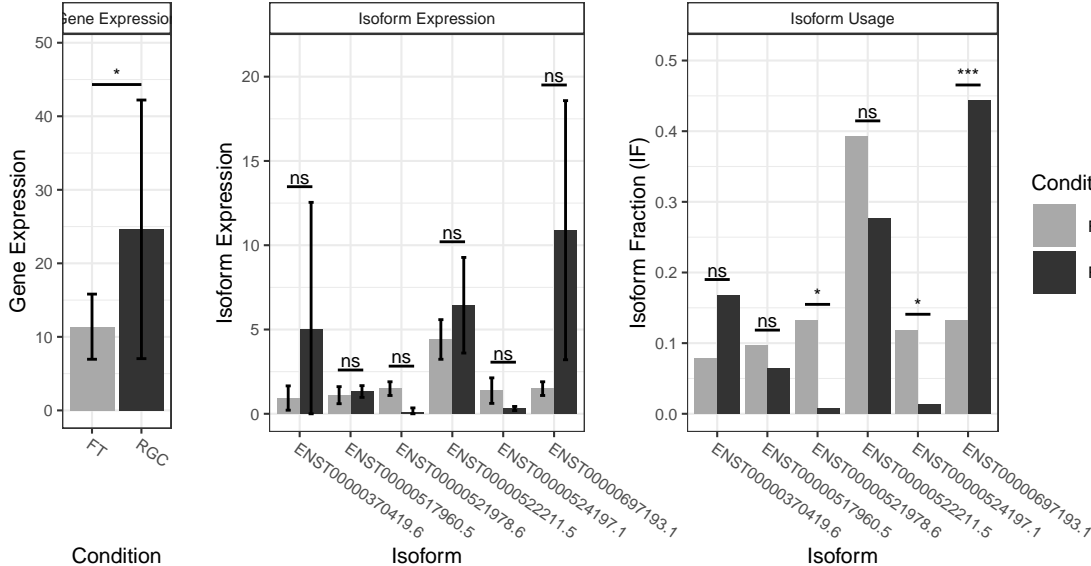
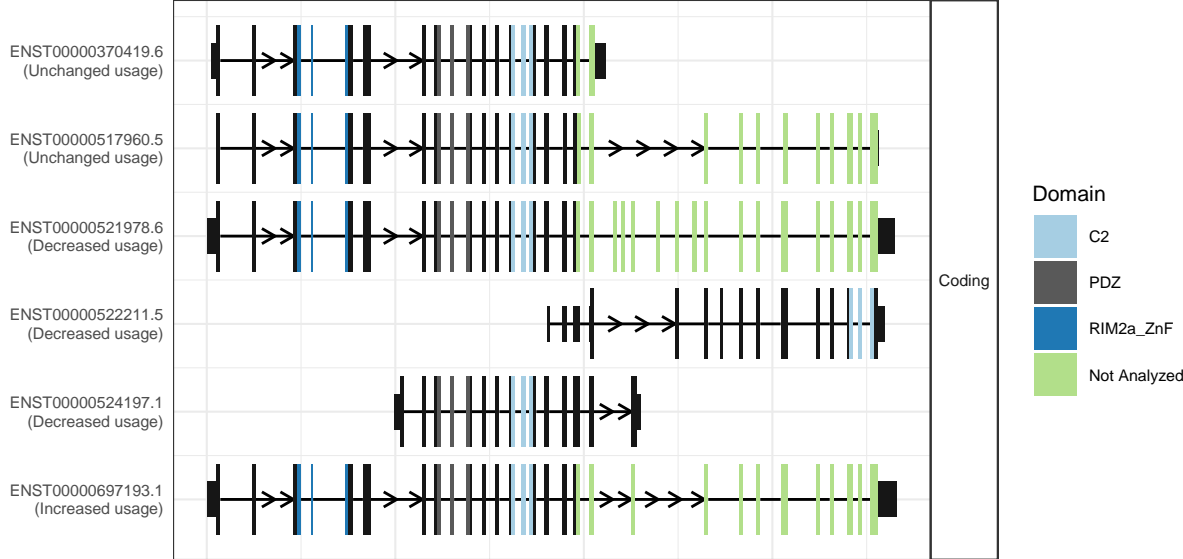


Domain



Homeodomain

The isoform switch in RIMS1 (FT vs RGC)



Domain



C2



PDZ

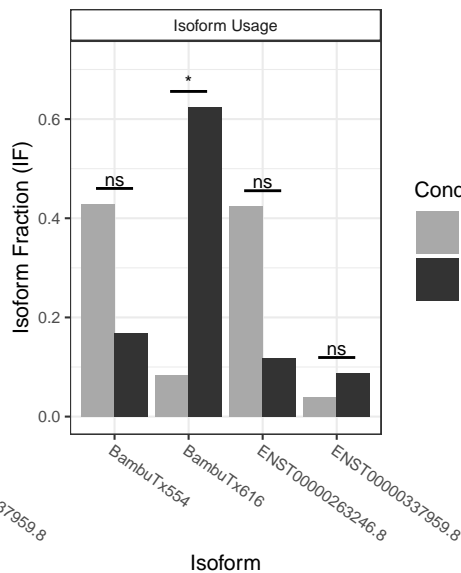
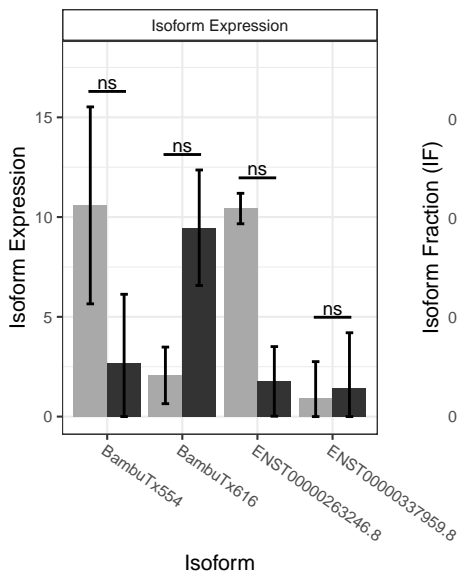
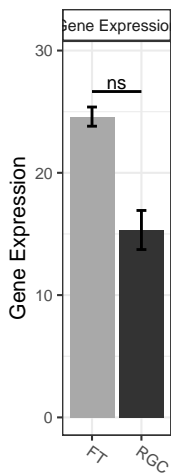
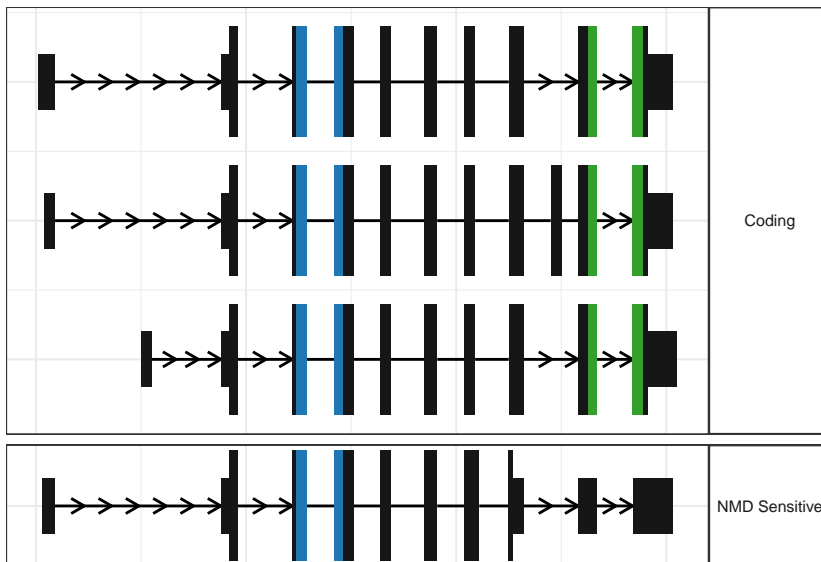


RIM2a_ZnF



Not Analyzed

The isoform switch in PACSIN2 (FT vs RGC)



Domain

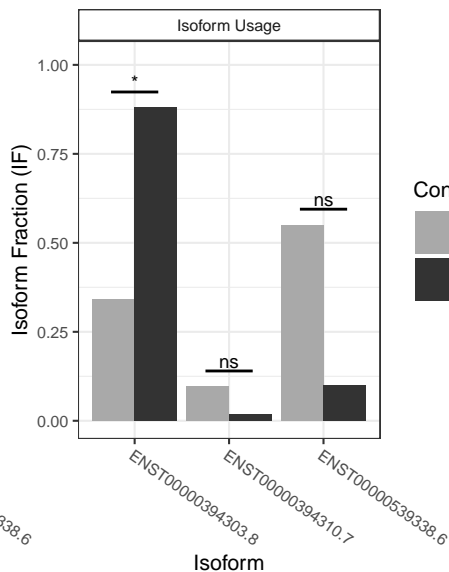
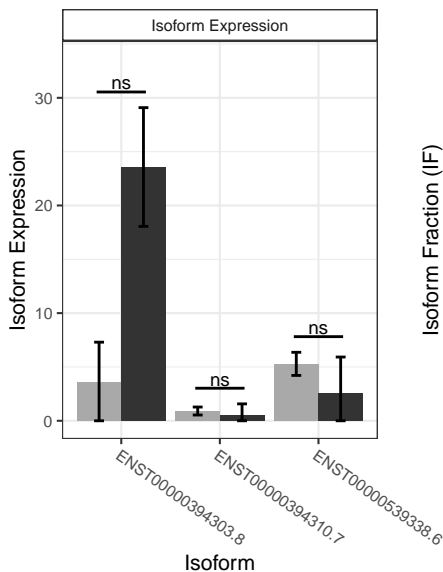
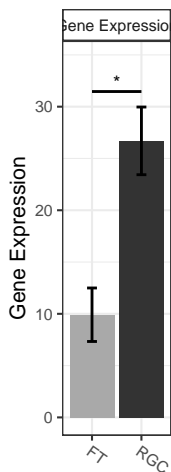
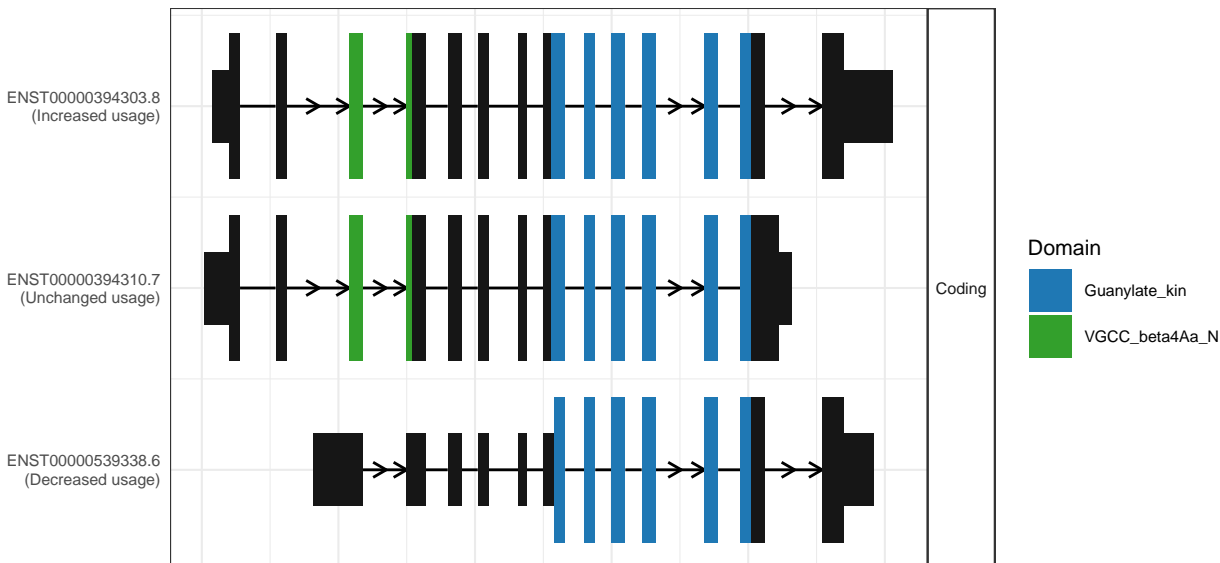


FCH



SH3_9

The isoform switch in CACNB1 (FT vs RGC)



Domain

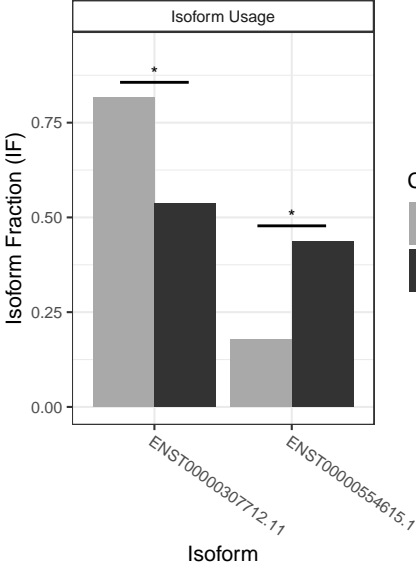
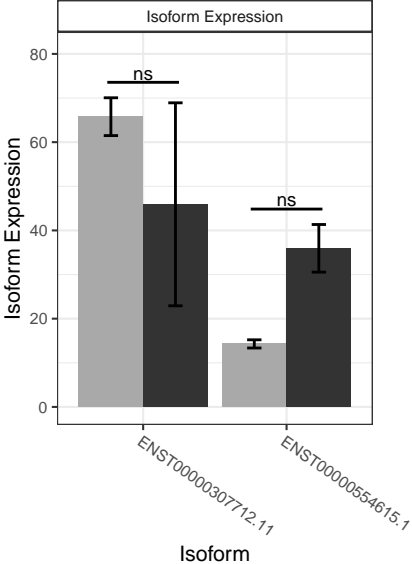
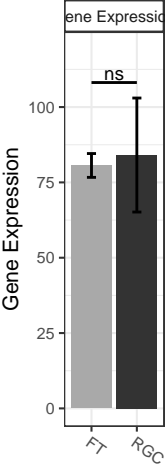
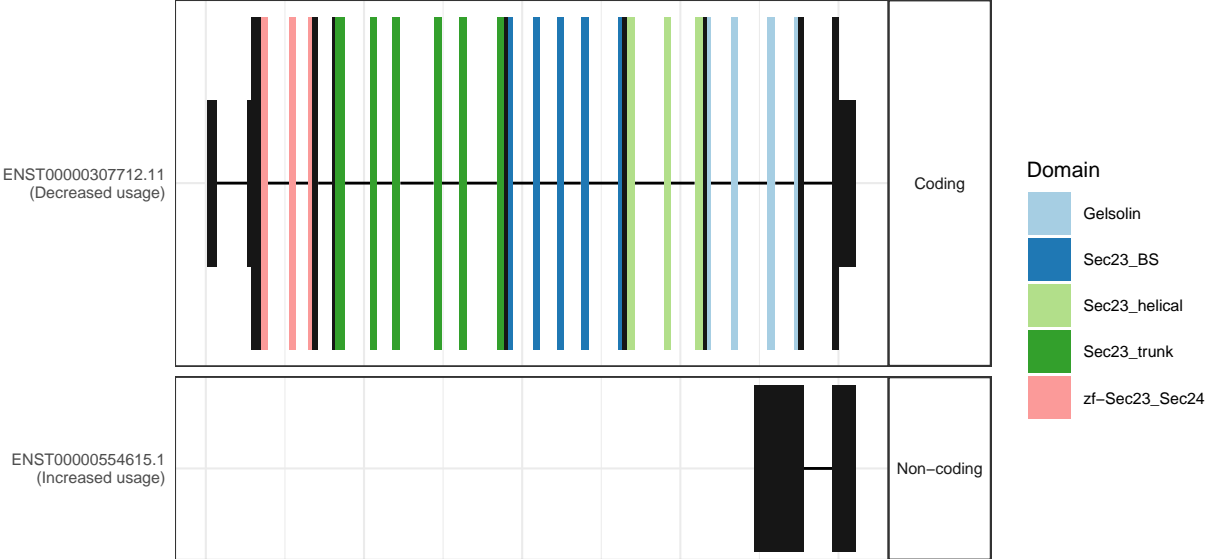


Guanylate_kin








VGCC_beta4Aa_N

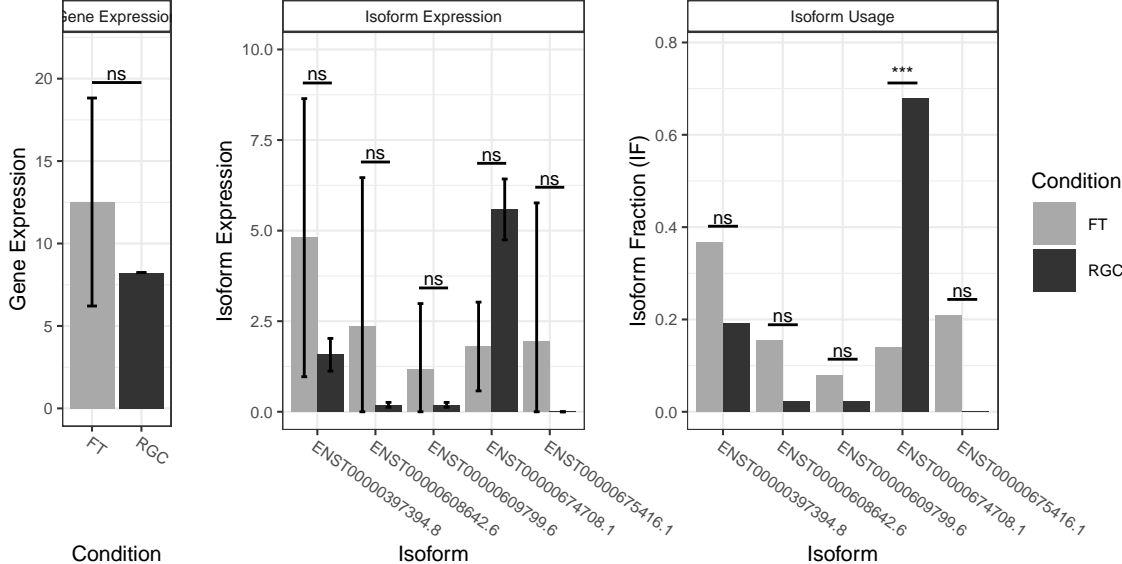
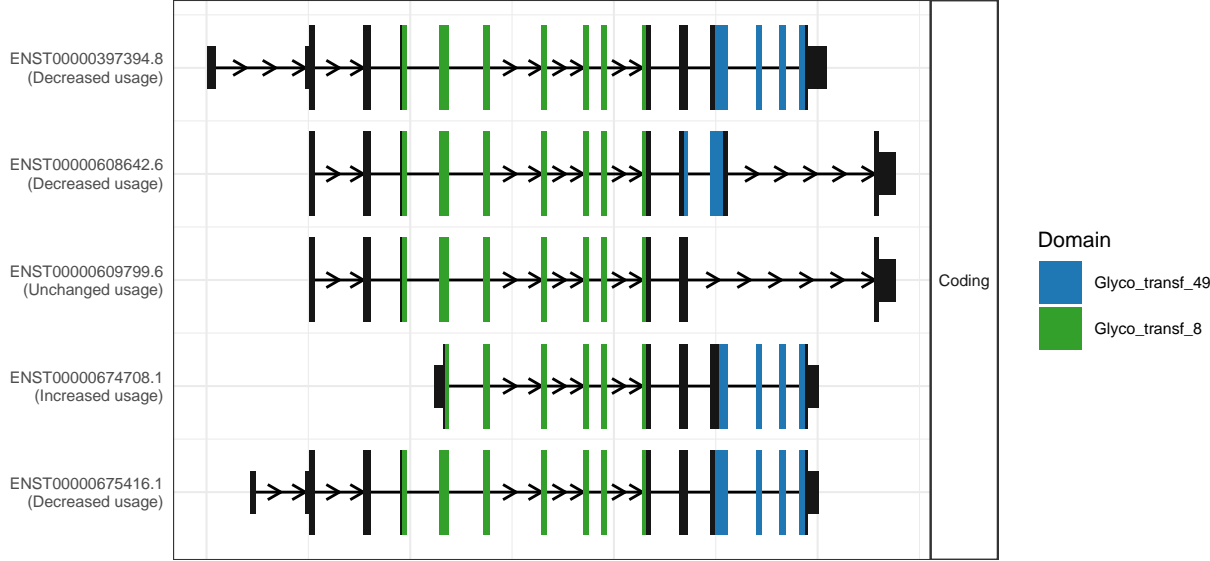
The isoform switch in SEC23A (FT vs RGC)



Domain

	Gelsolin
	Sec23_BS
	Sec23_helical
	Sec23_trunk
	zf-Sec23_Sec24

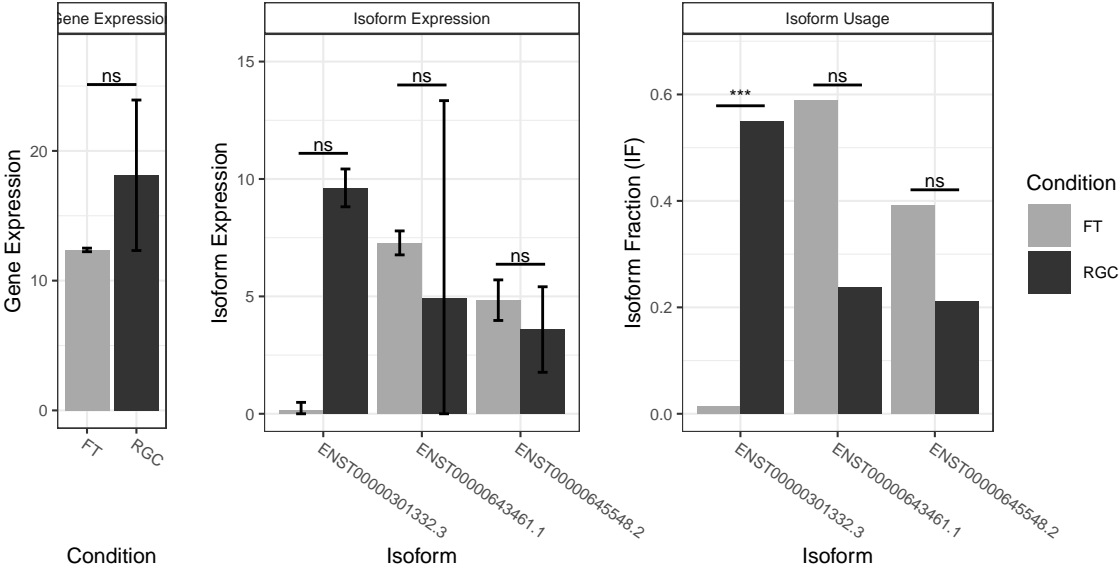
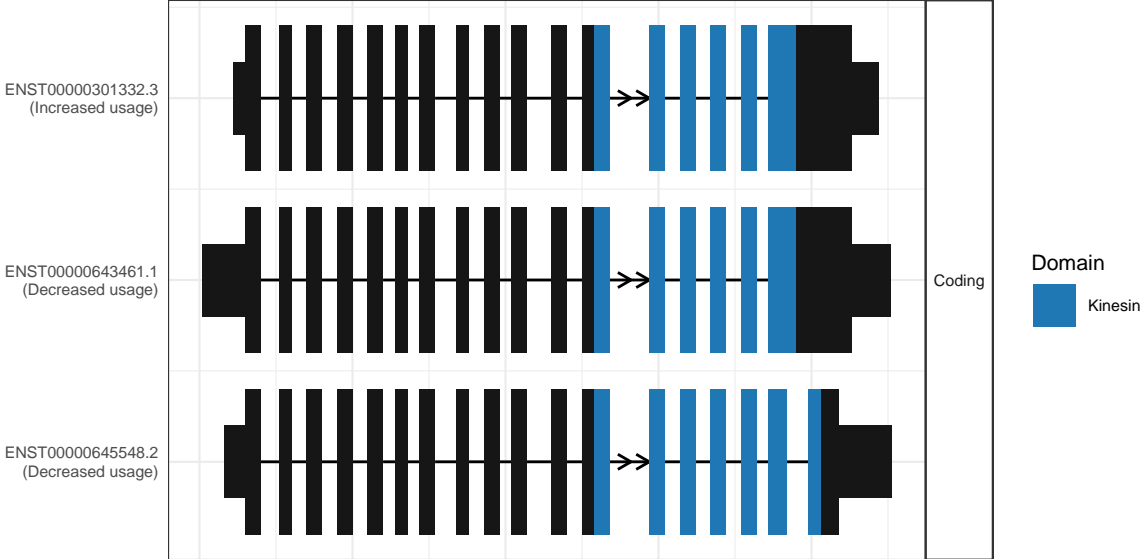
The isoform switch in LARGE1 (FT vs RGC)



Domain



The isoform switch in KIFC2 (FT vs RGC)

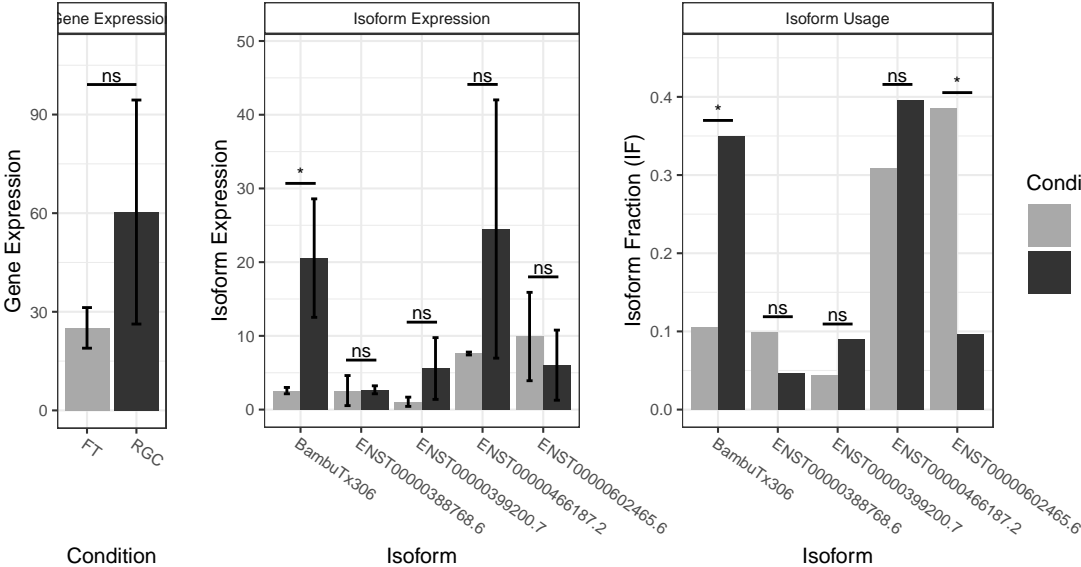
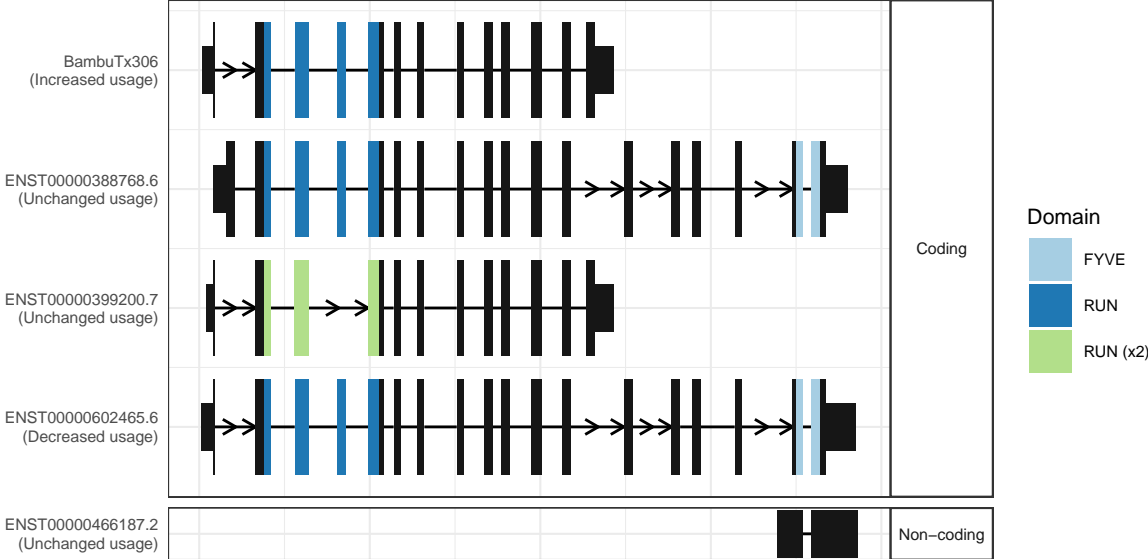


Domain



Kinesin

The isoform switch in RUFY2 (FT vs RGC)



Domain



FYVE

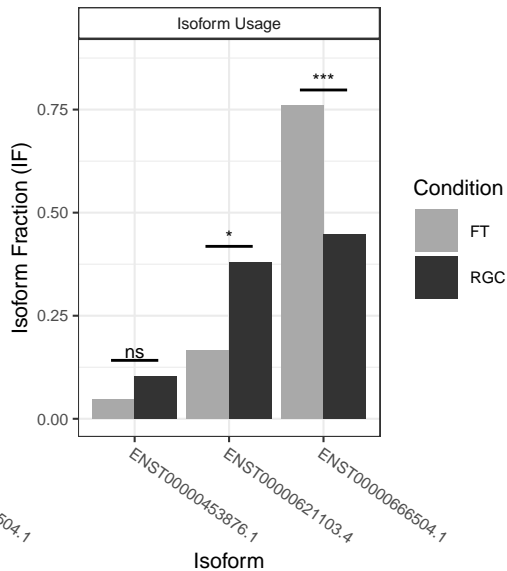
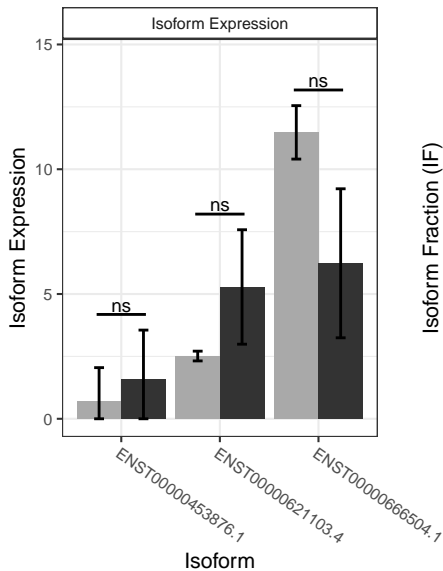
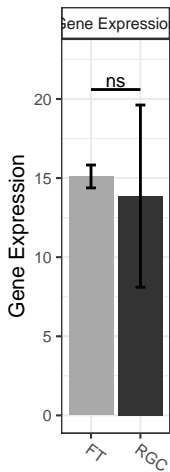
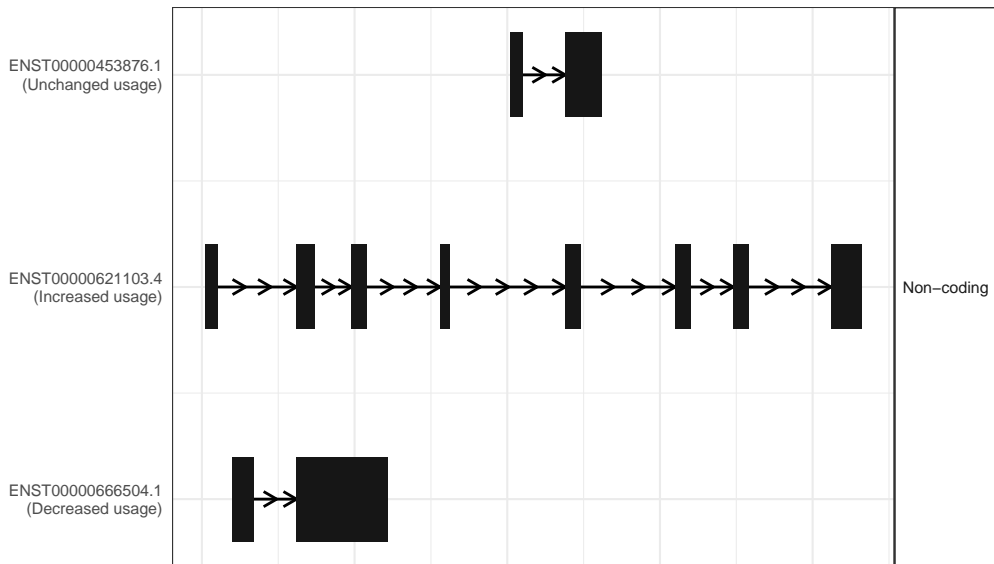


RUN

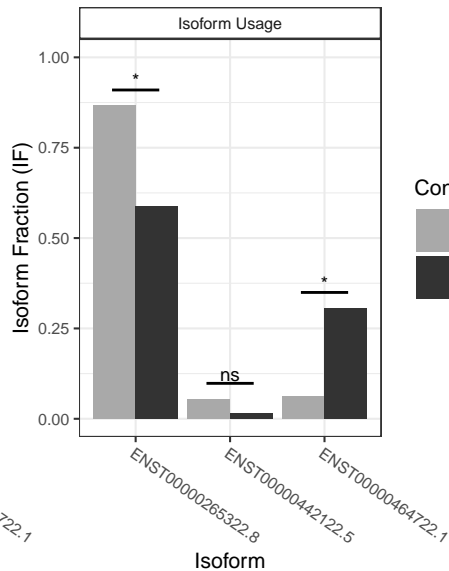
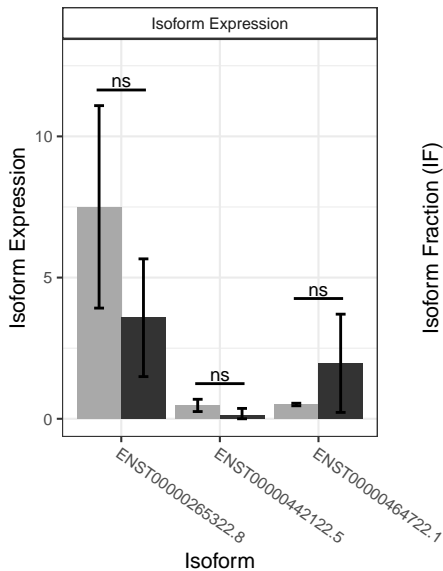
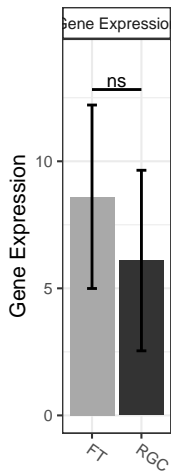
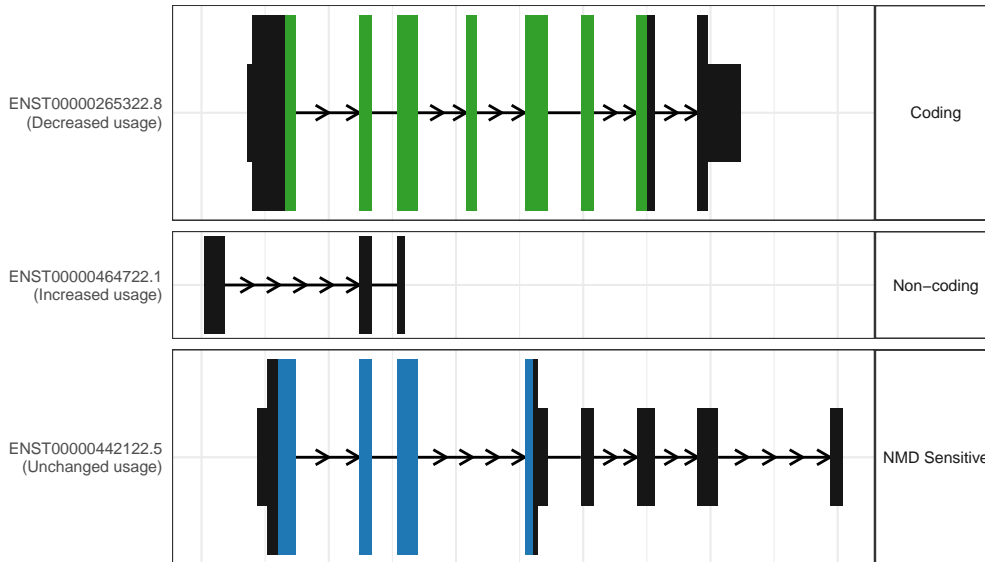


RUN (x2)

The isoform switch in MIR3936HG (FT vs RGC)



The isoform switch in PECCR (FT vs RGC)



Domain

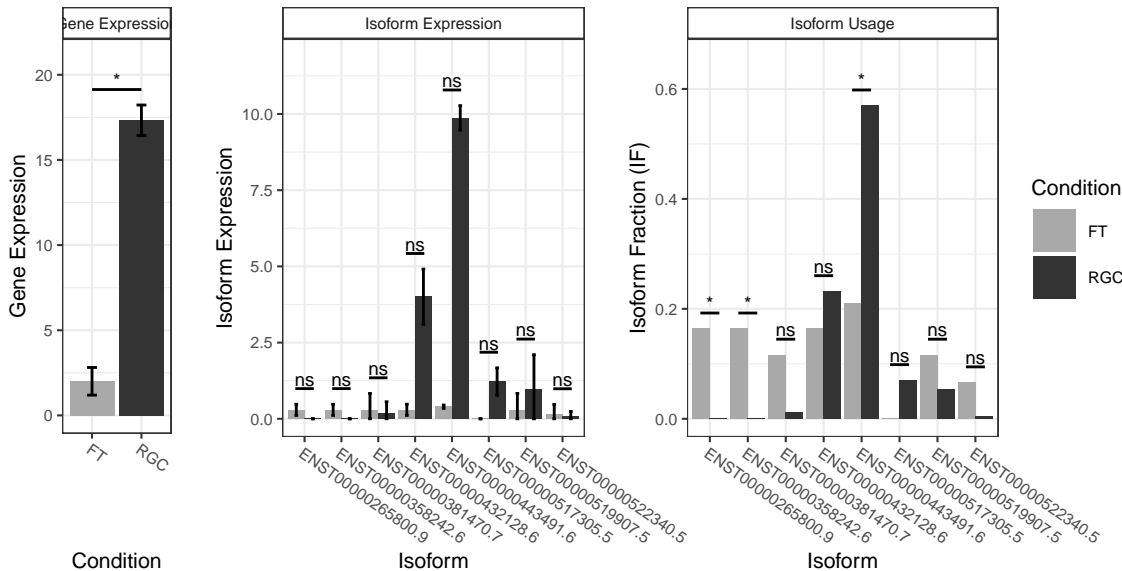
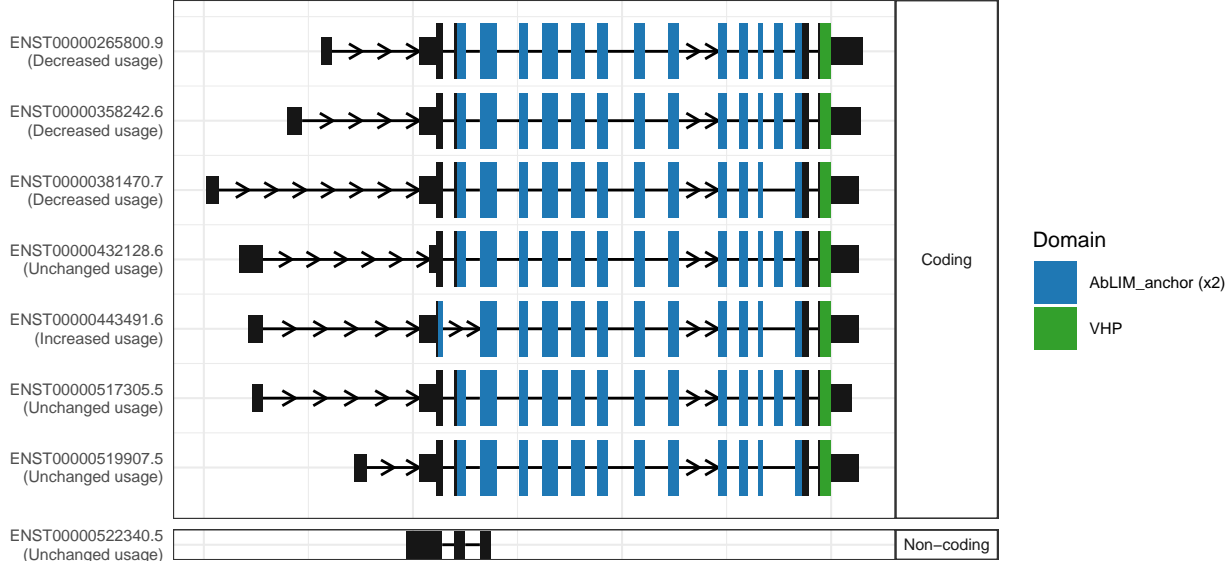


adh_short



adh_short_C2

The isoform switch in DMTN (FT vs RGC)



Domain

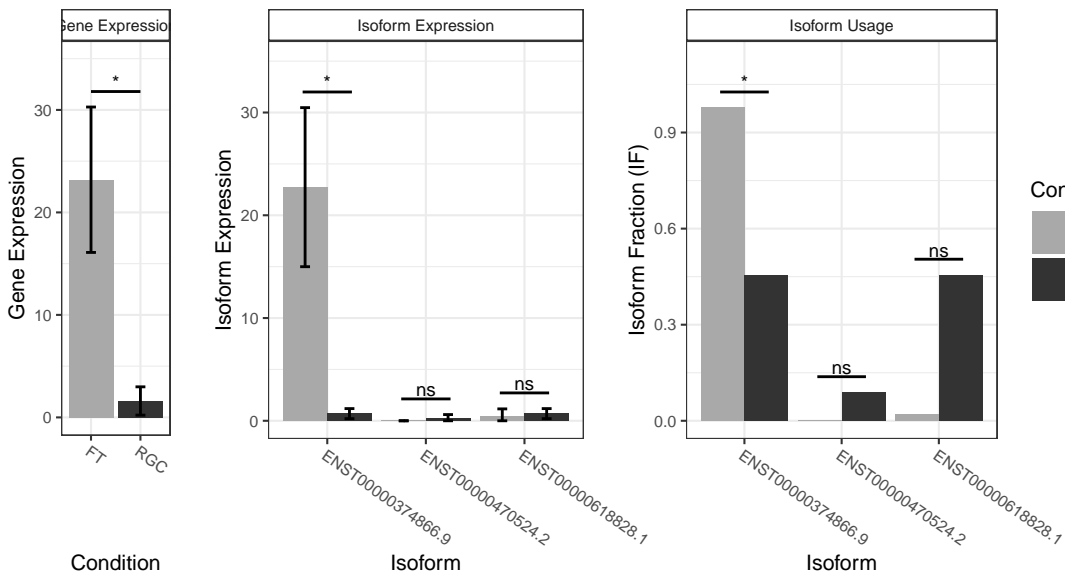
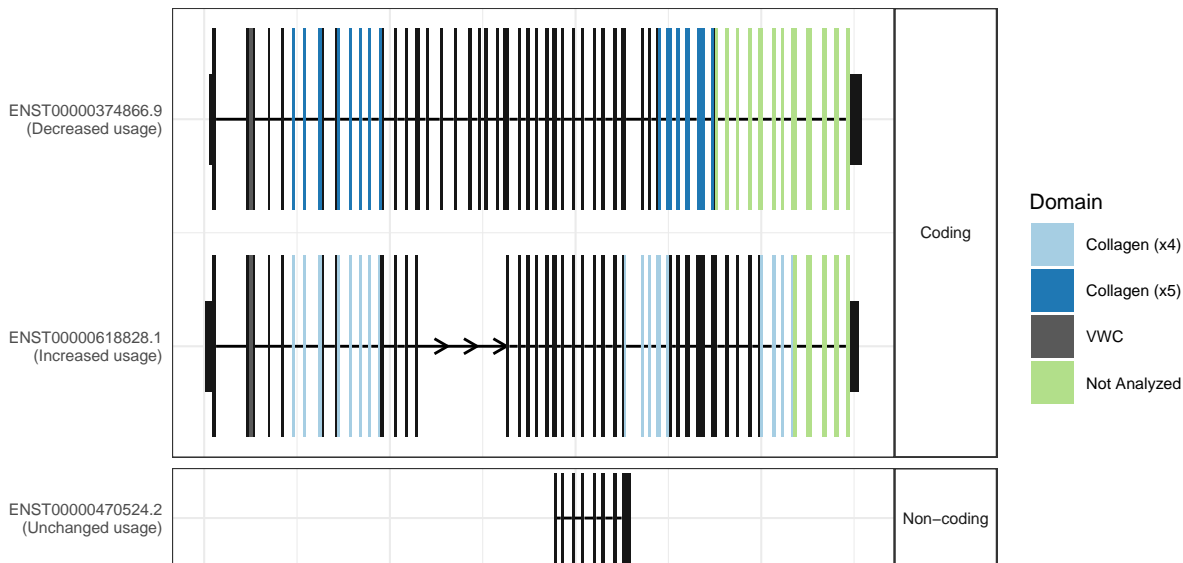


AbLIM_anchor (x2)

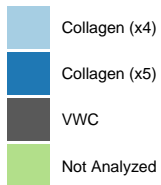


VHP

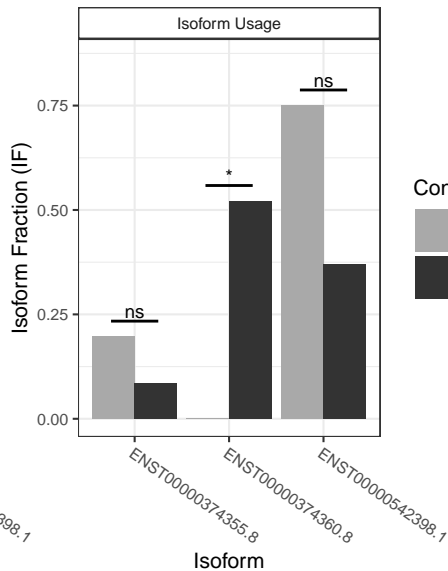
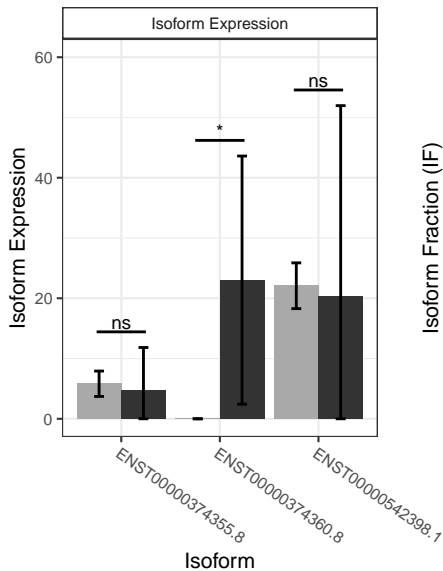
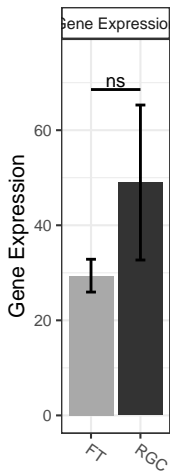
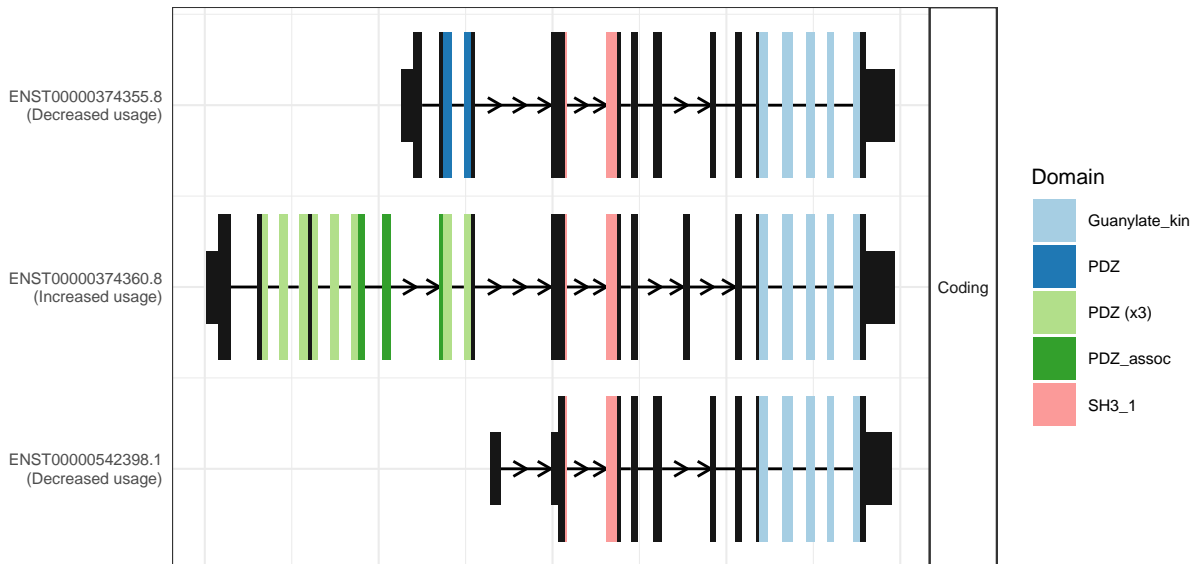
The isoform switch in COL5A2 (FT vs RGC)








Domain



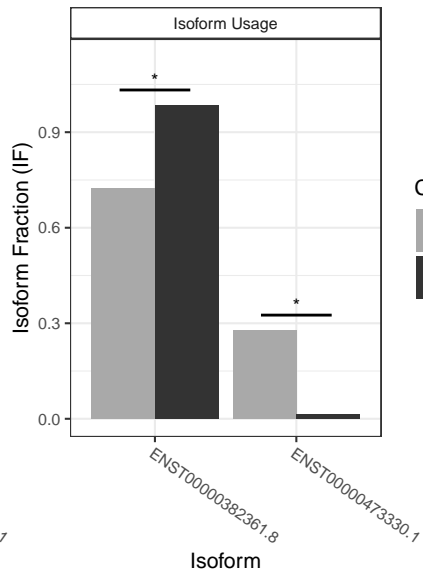
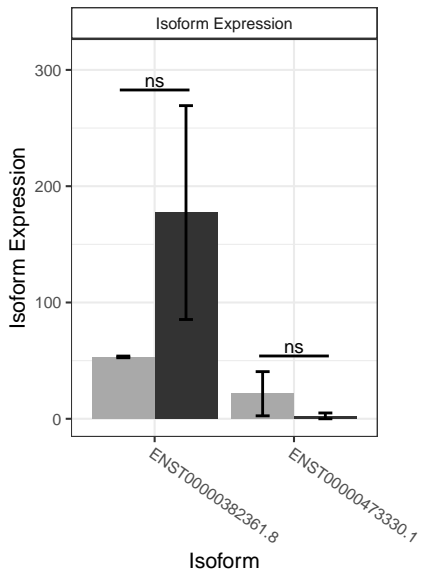
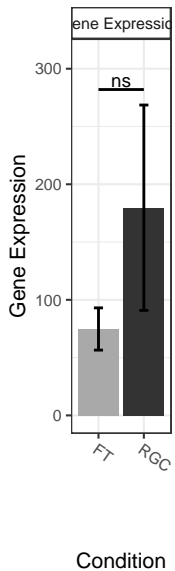
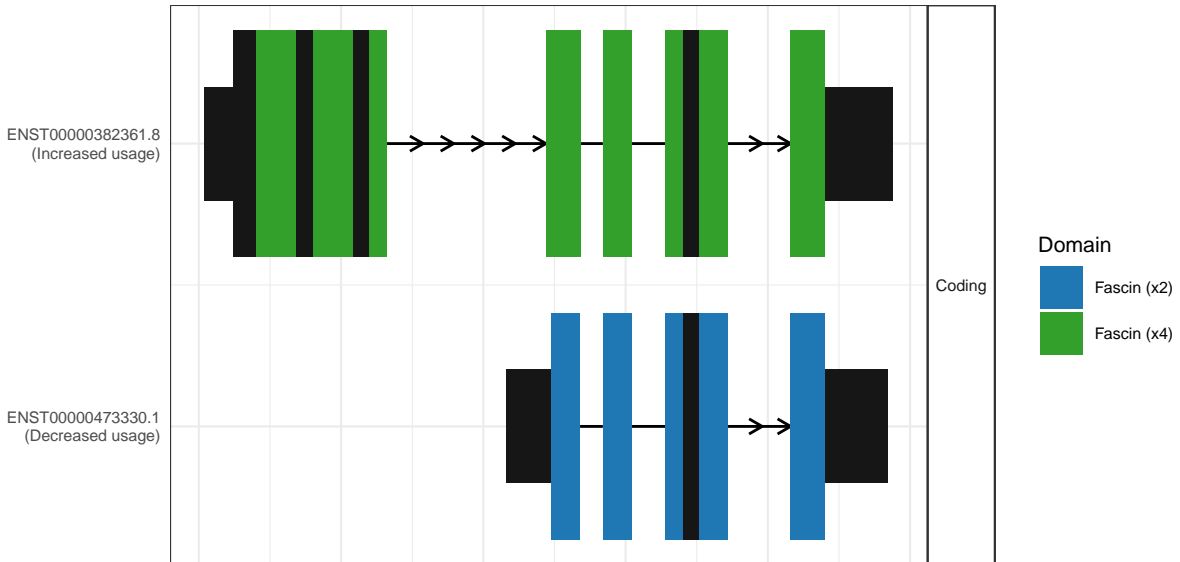
The isoform switch in DLG3 (FT vs RGC)



Domain

	Guanylate_kin
	PDZ
	PDZ (x3)
	PDZ_assoc
	SH3_1

The isoform switch in FSCN1 (FT vs RGC)



Domain

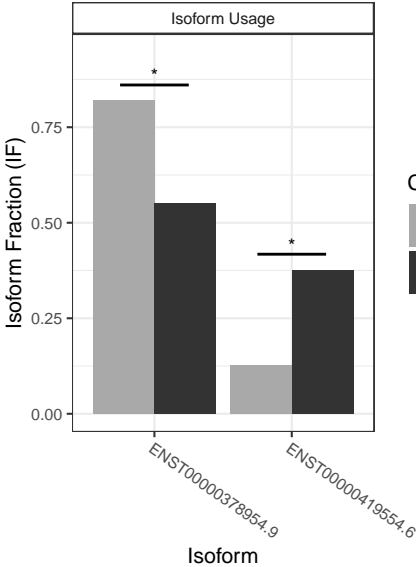
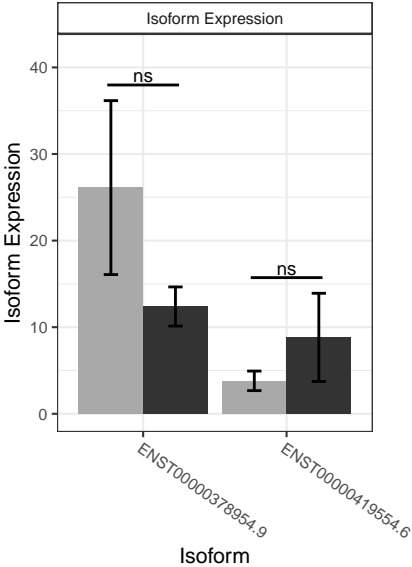
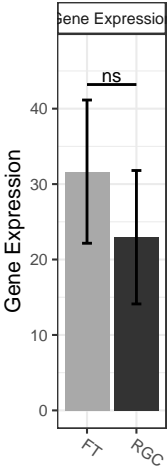
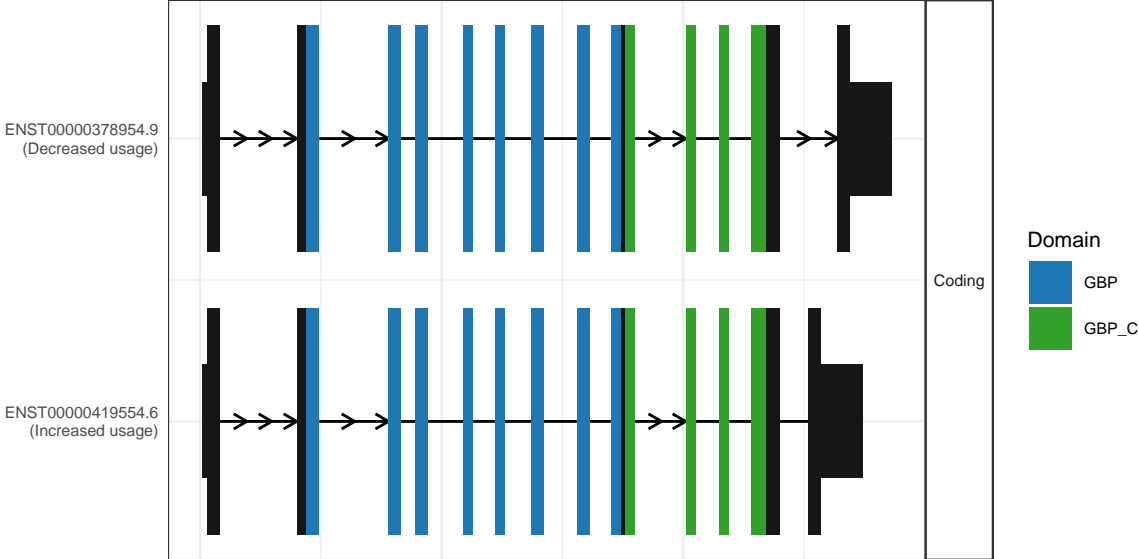


Fascin (x2)



Fascin (x4)

The isoform switch in ATL2 (FT vs RGC)



Domain

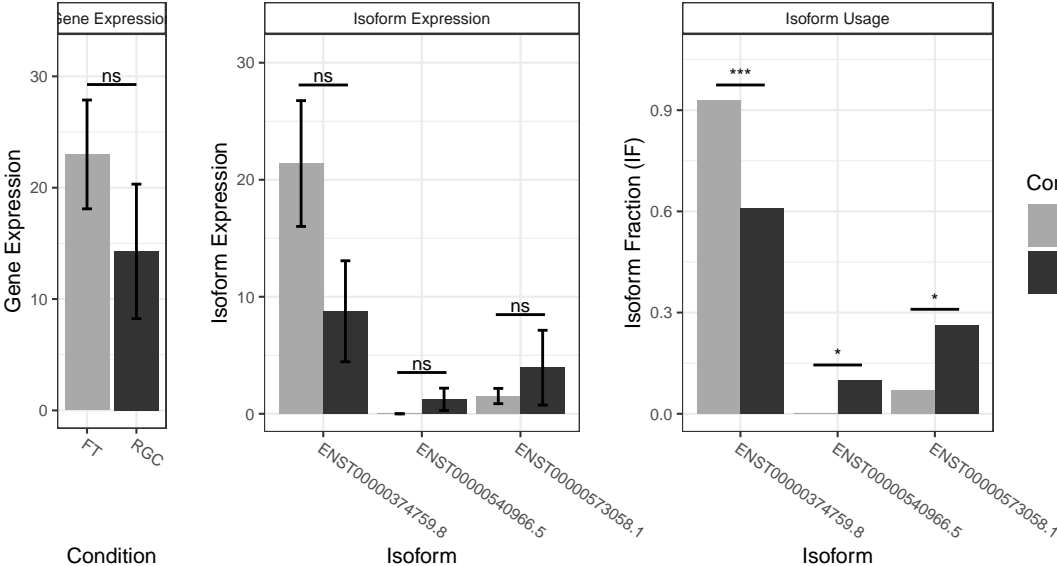
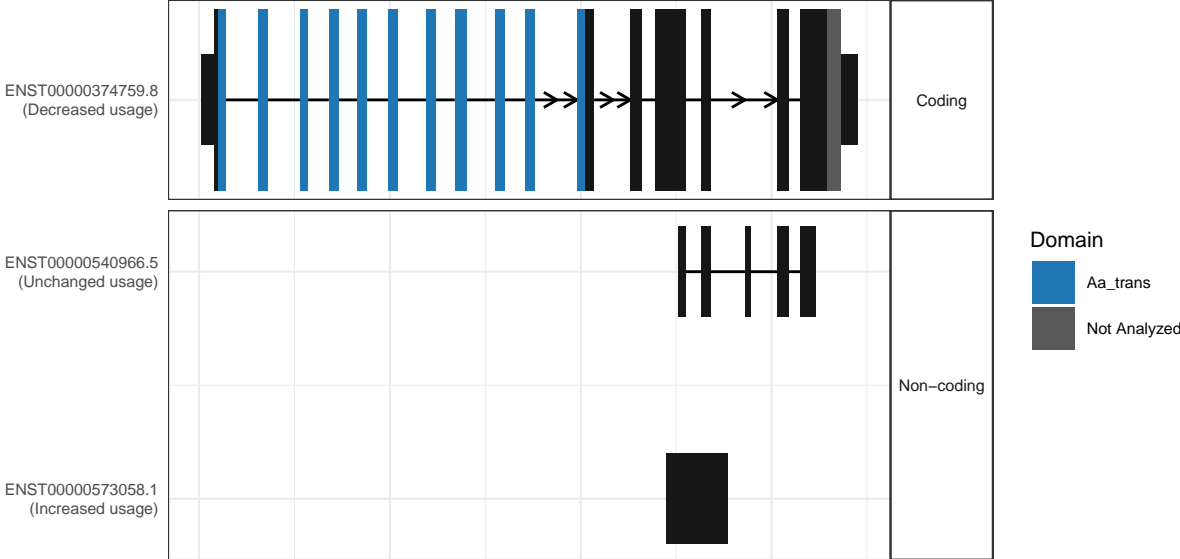


GBP



GBP_C

The isoform switch in SLC38A10 (FT vs RGC)



Domain

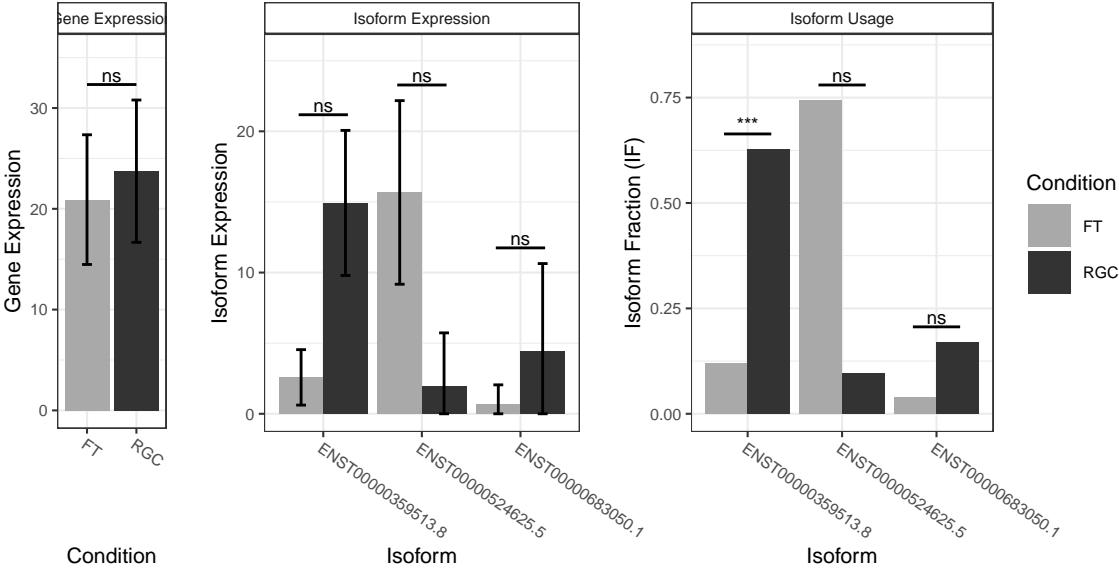
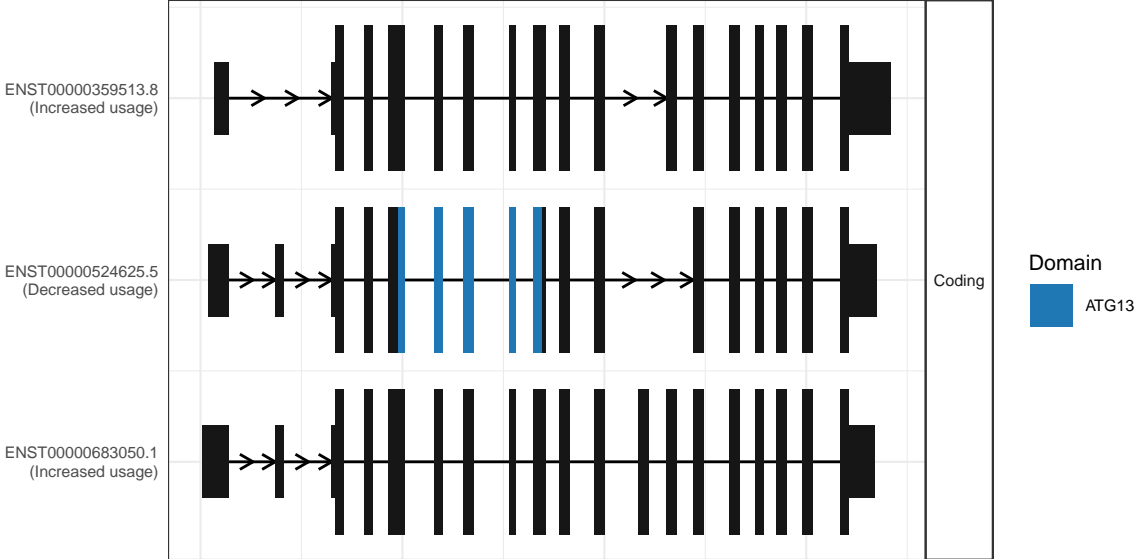


Aa_trans



Not Analyzed

The isoform switch in ATG13 (FT vs RGC)

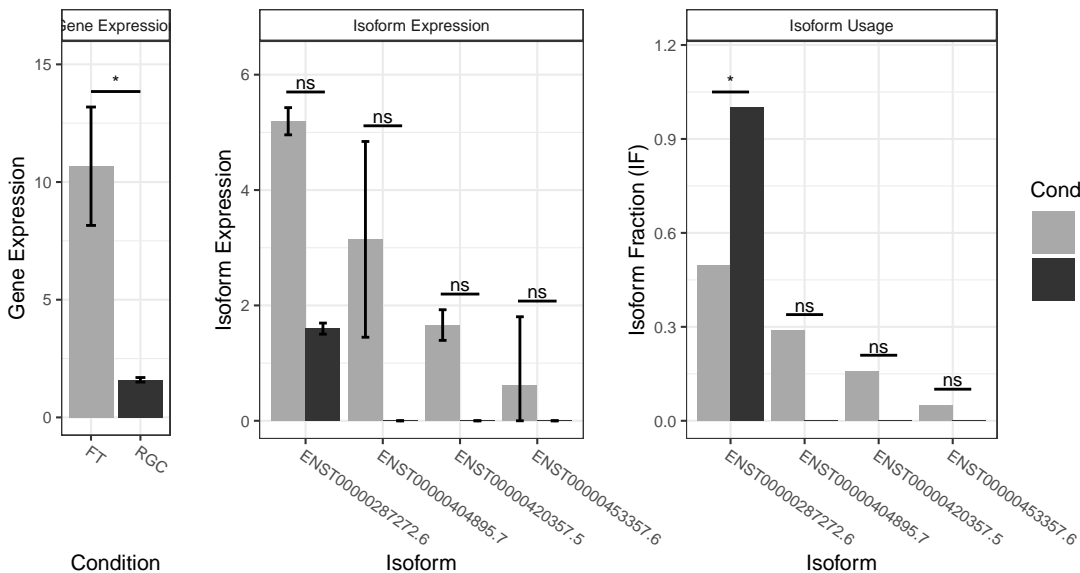
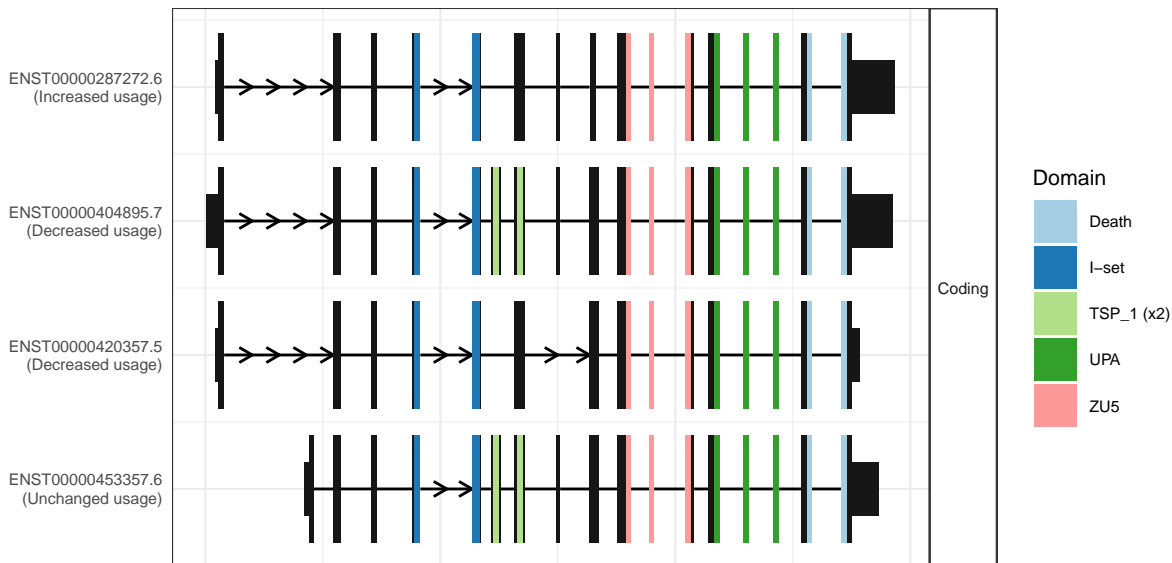


Domain



ATG13

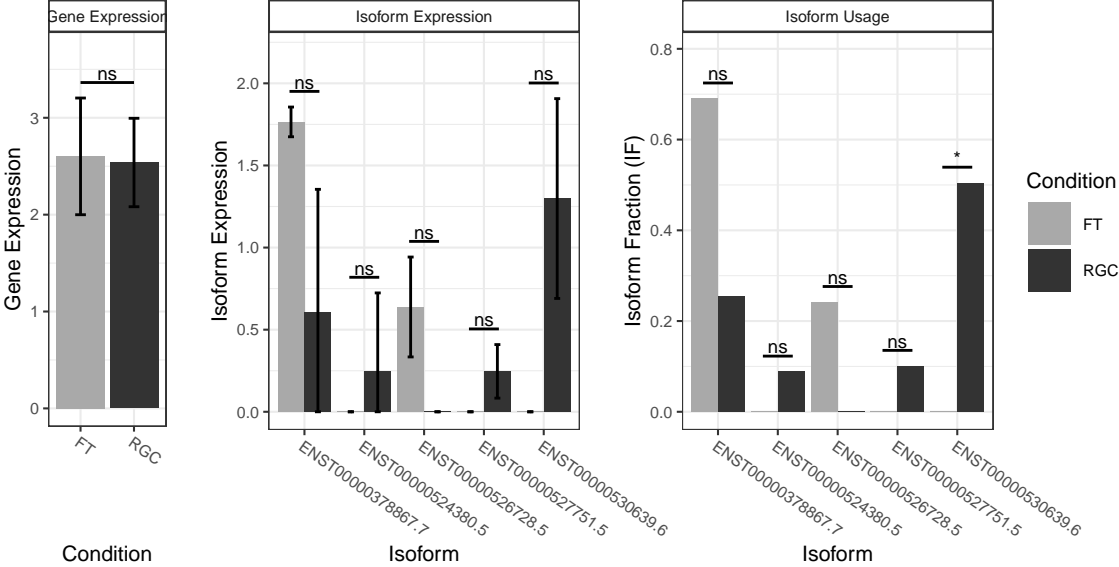
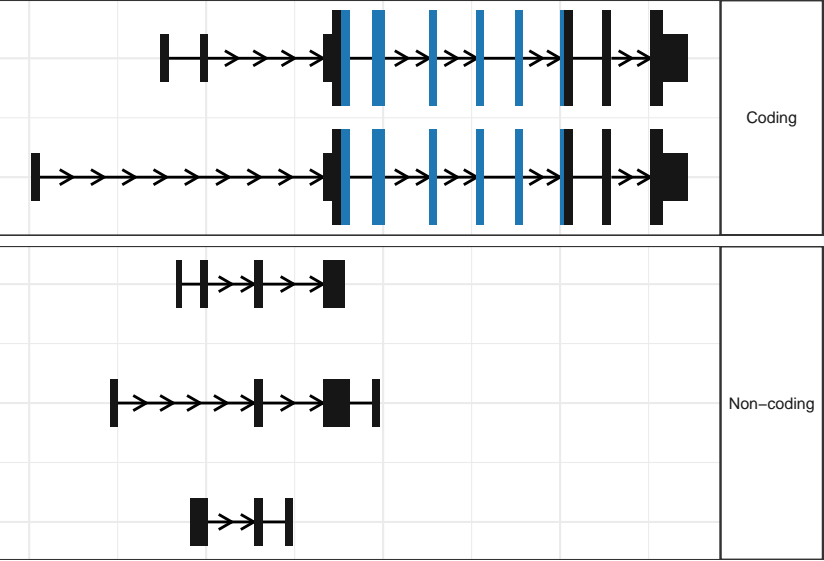
The isoform switch in UNC5D (FT vs RGC)



Domain

	Death
	I-set
	TSP_1 (x2)
	UPA
	ZU5

The isoform switch in PRR5L (FT vs RGC)

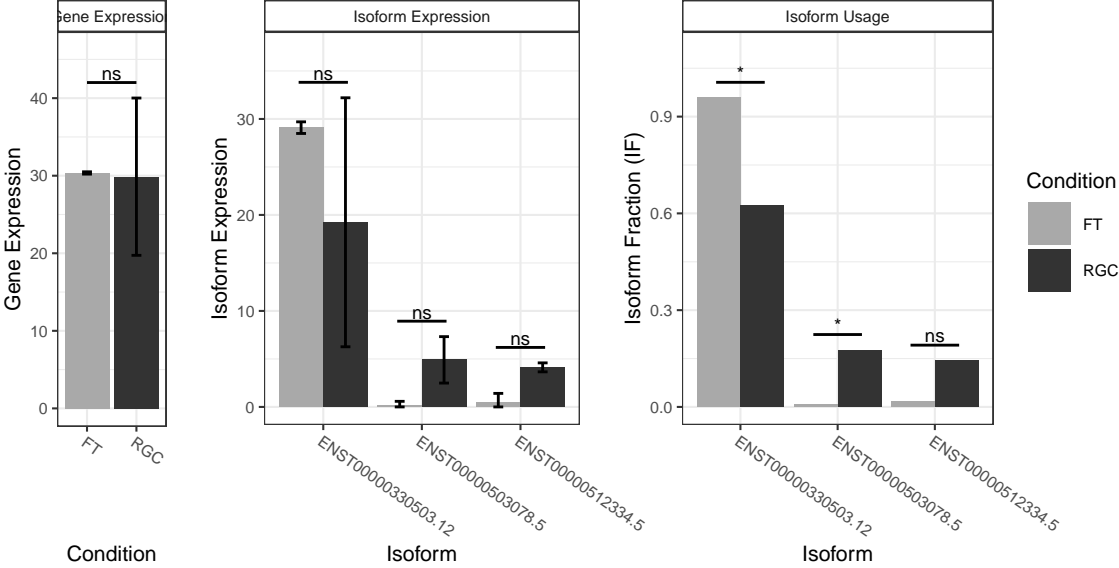
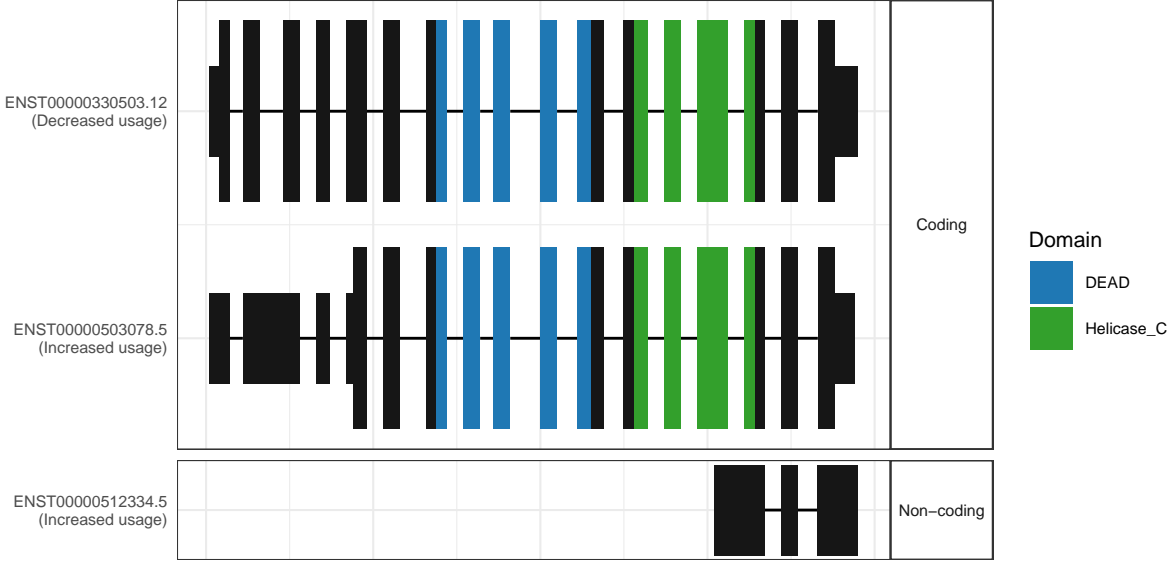


Domain



HbrB

The isoform switch in DDX41 (FT vs RGC)



Domain

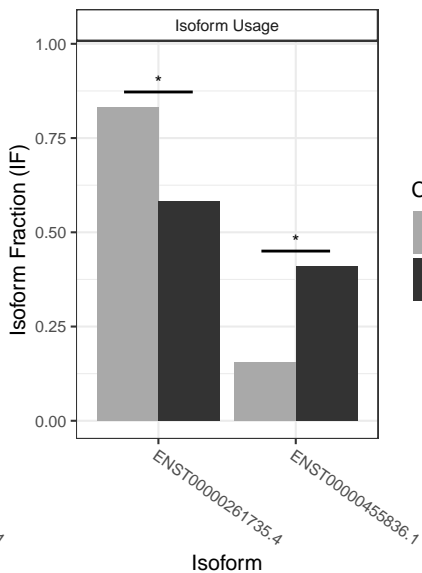
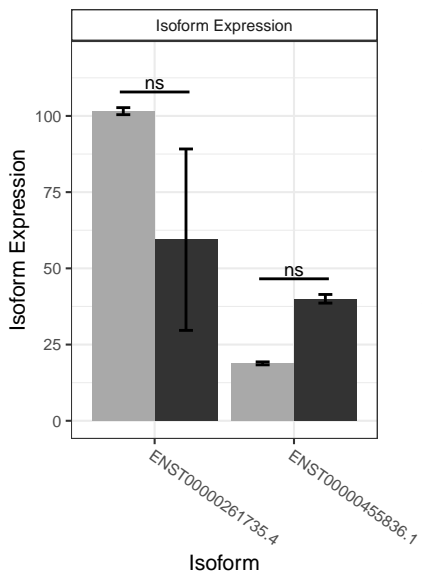
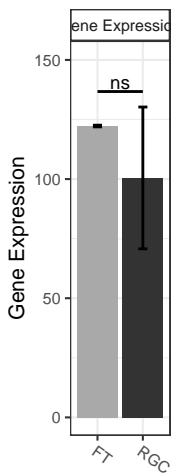
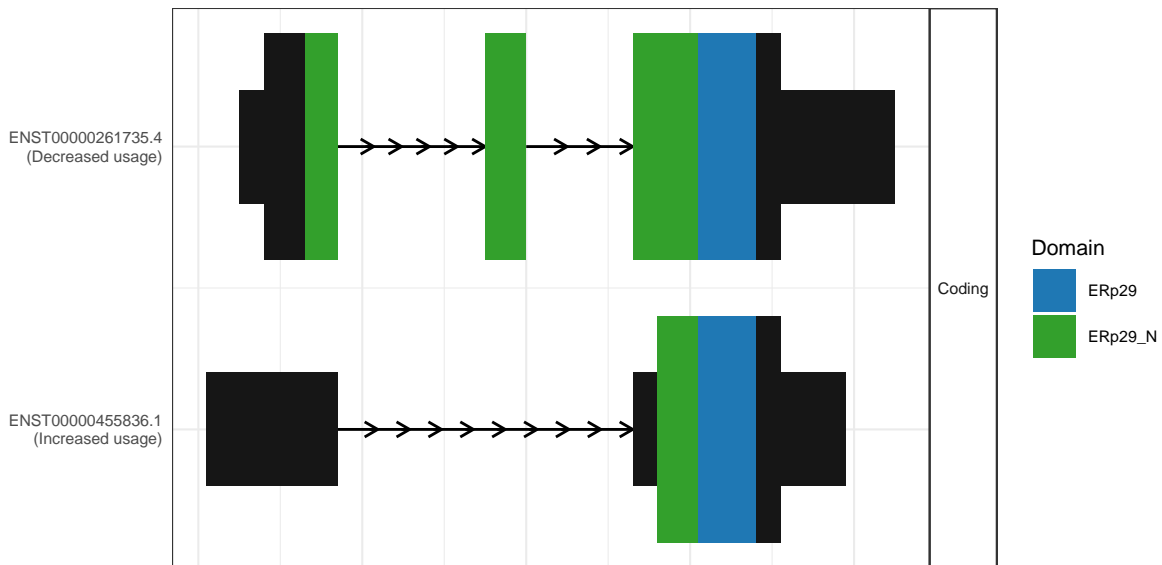


DEAD



Helicase_C

The isoform switch in ERP29 (FT vs RGC)



Domain

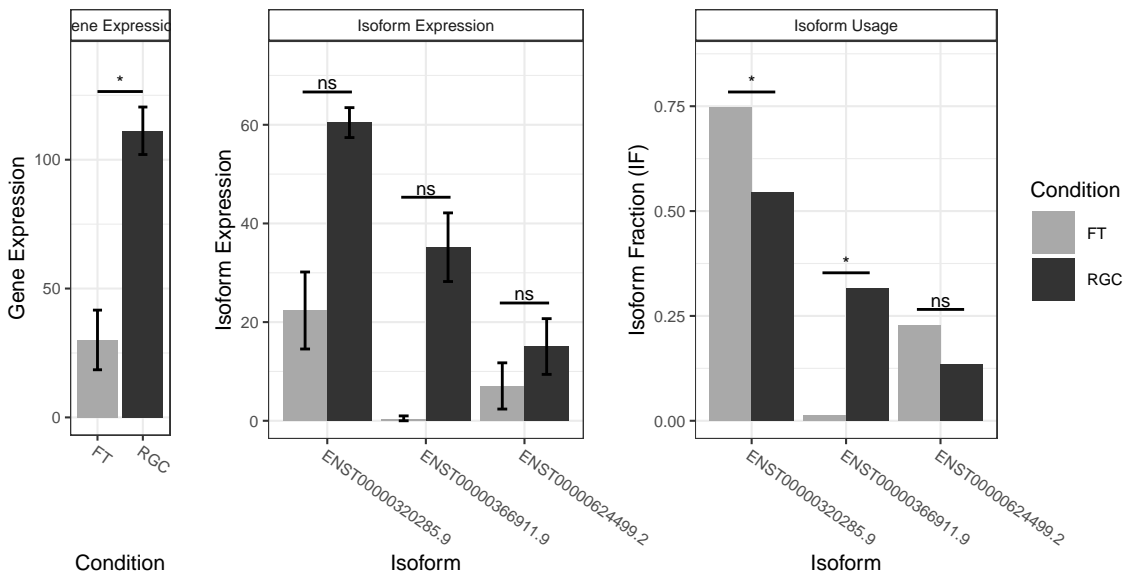
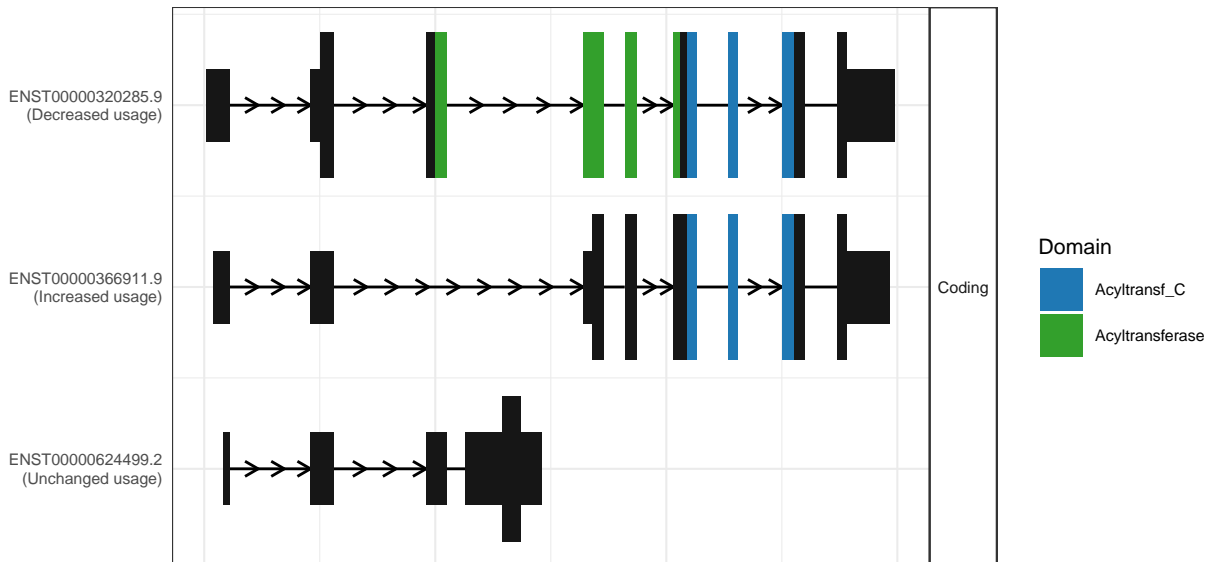


ERp29



ERp29_N

The isoform switch in AGPAT4 (FT vs RGC)



Domain

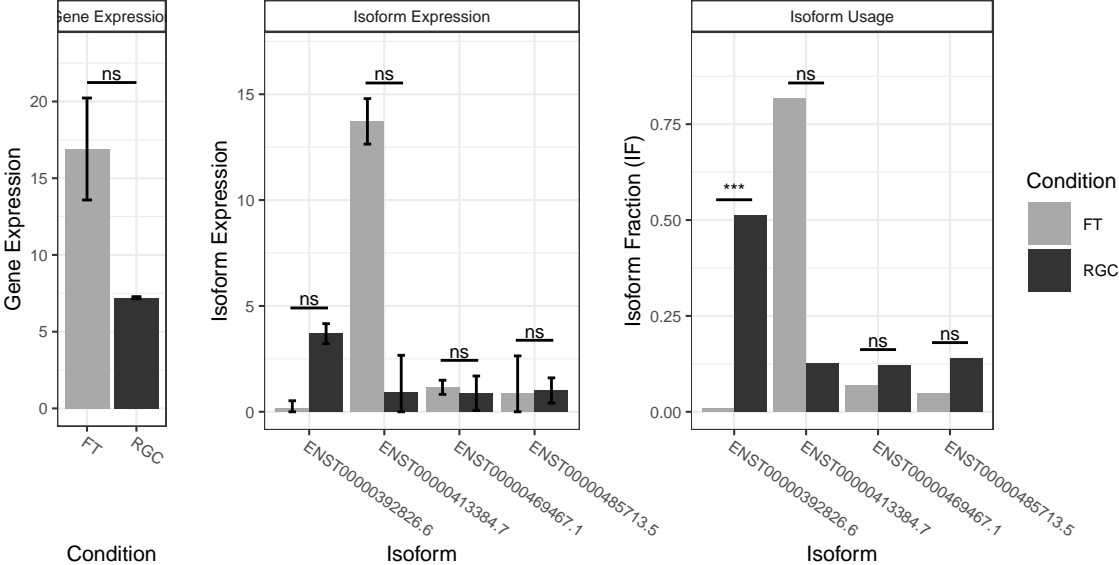
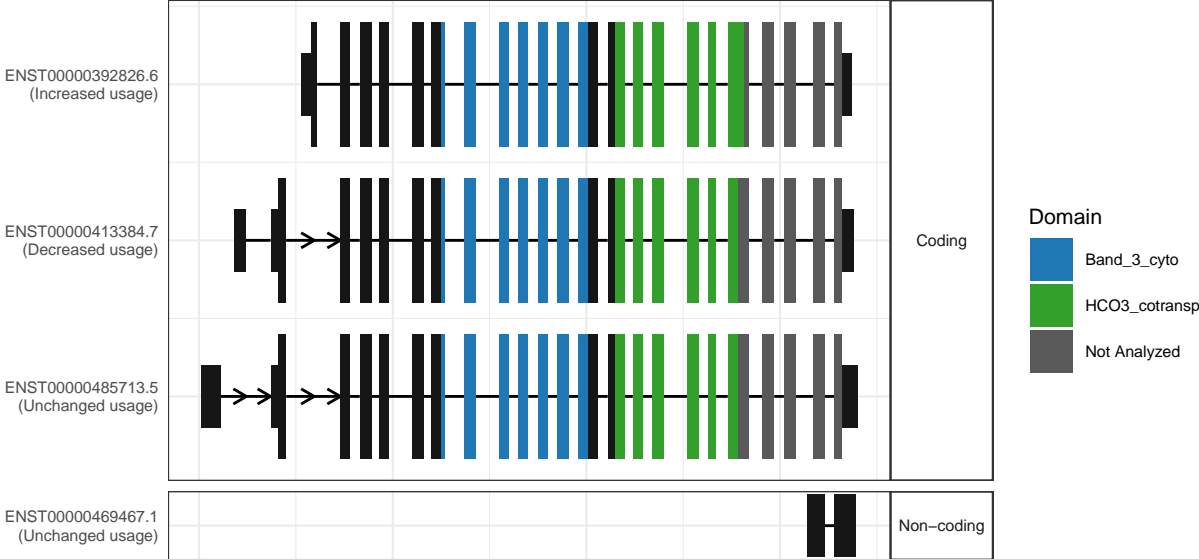


Acyltransf_C

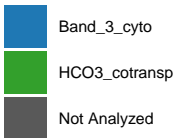


Acyltransferase

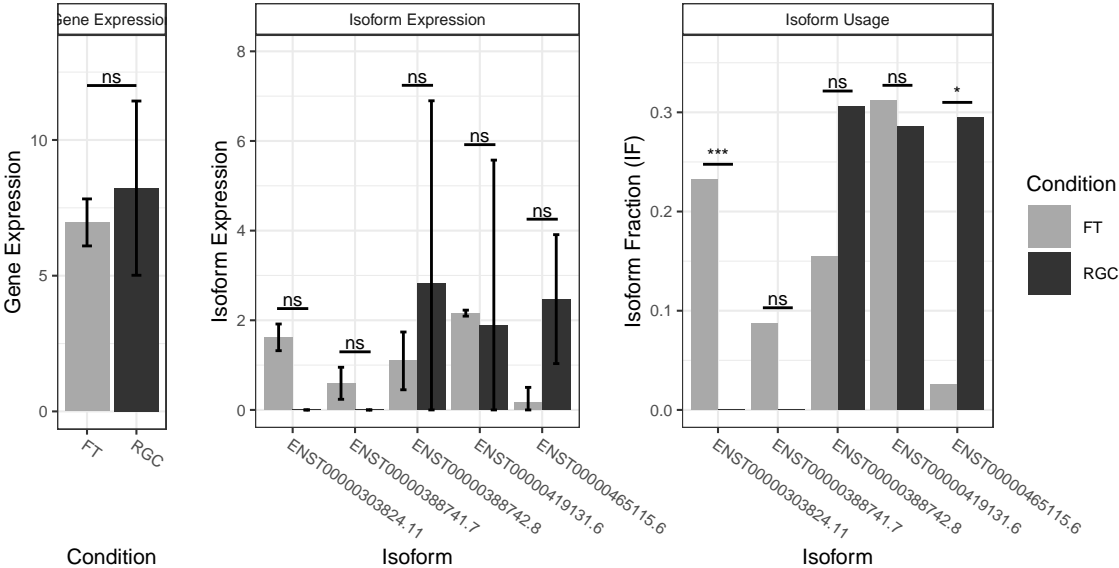
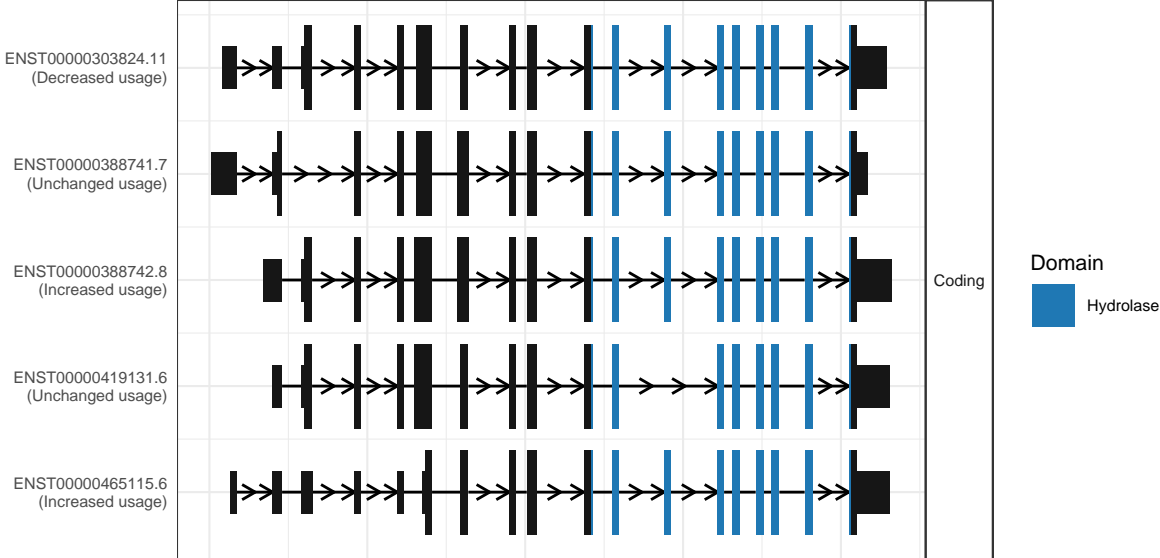
The isoform switch in SLC4A2 (FT vs RGC)



Domain



The isoform switch in EYA1 (FT vs RGC)

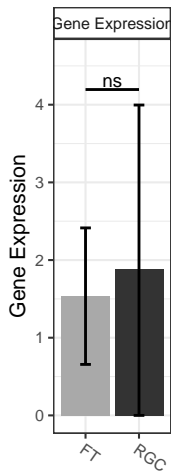
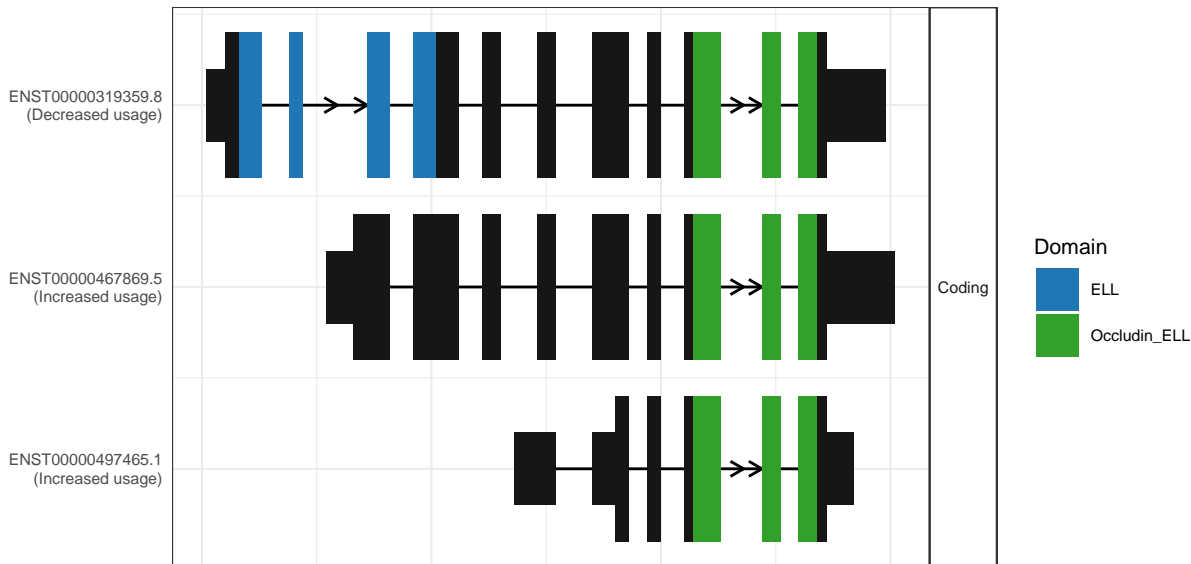


Domain

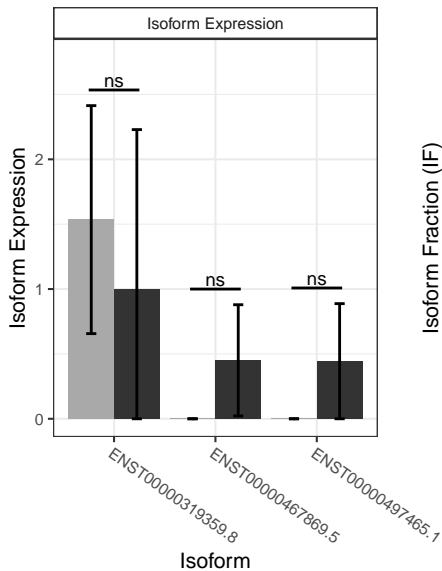


Hydrolase

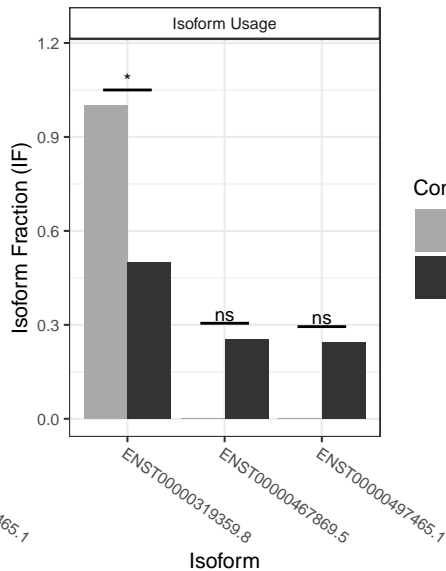
The isoform switch in ELL3 (FT vs RGC)



Condition



Isoform



Isoform

Condition



Domain

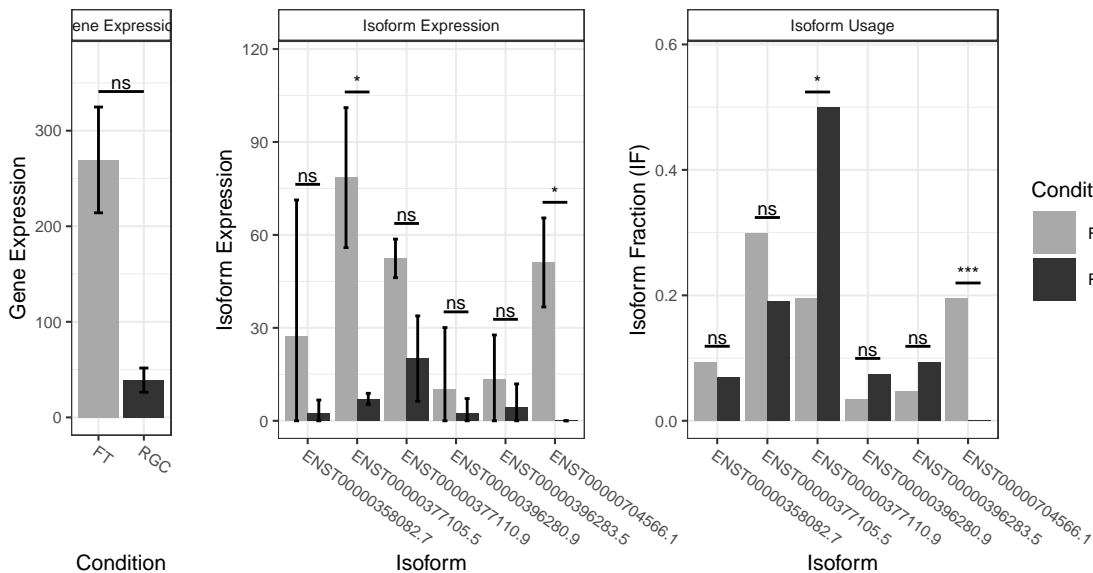
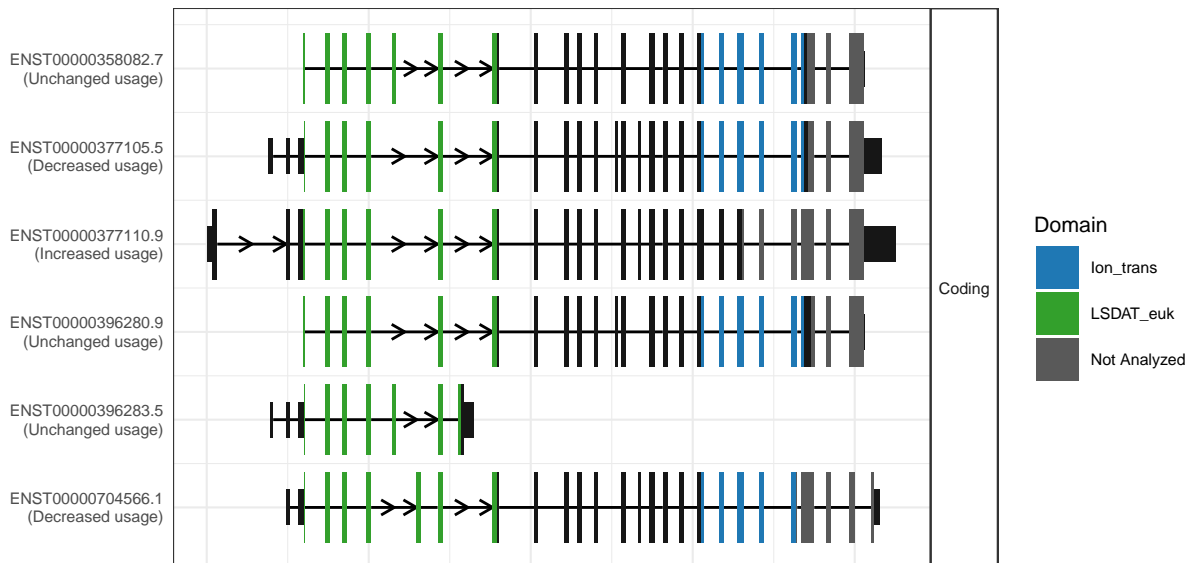


ELL

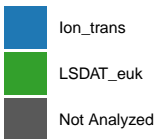


Occludin_ELL

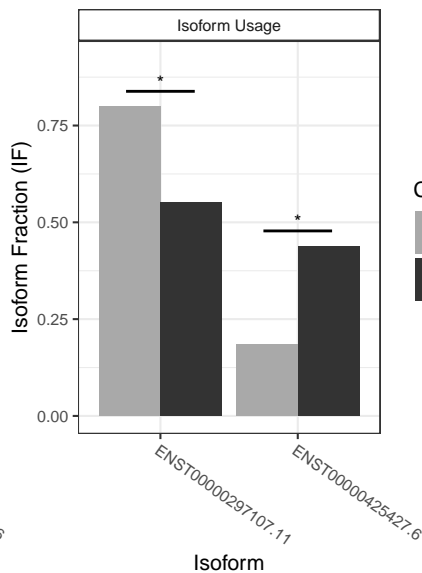
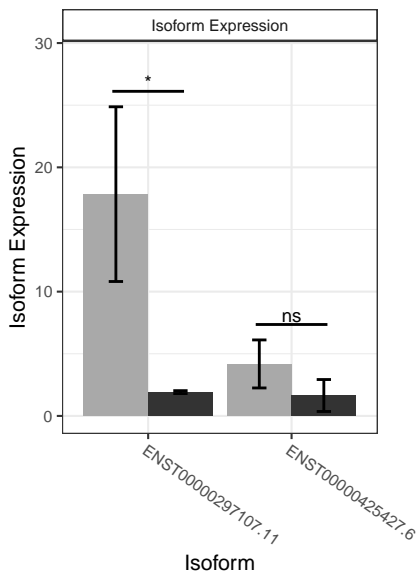
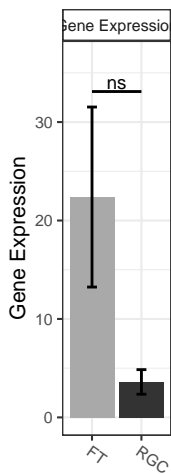
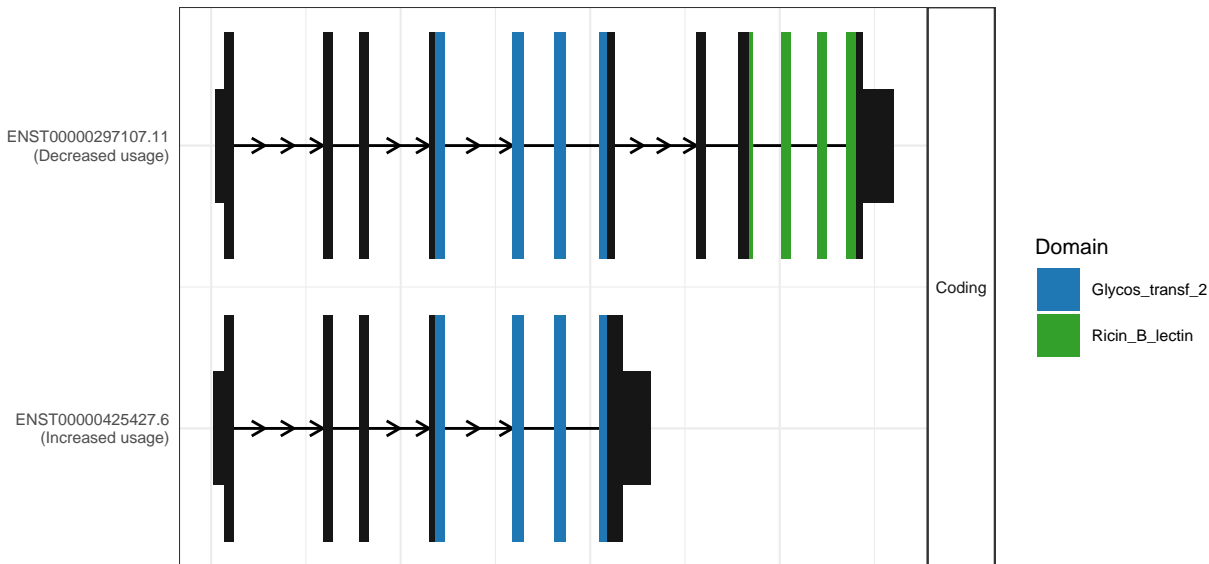
The isoform switch in TRPM3 (FT vs RGC)



Domain



The isoform switch in GALNT10 (FT vs RGC)



Domain

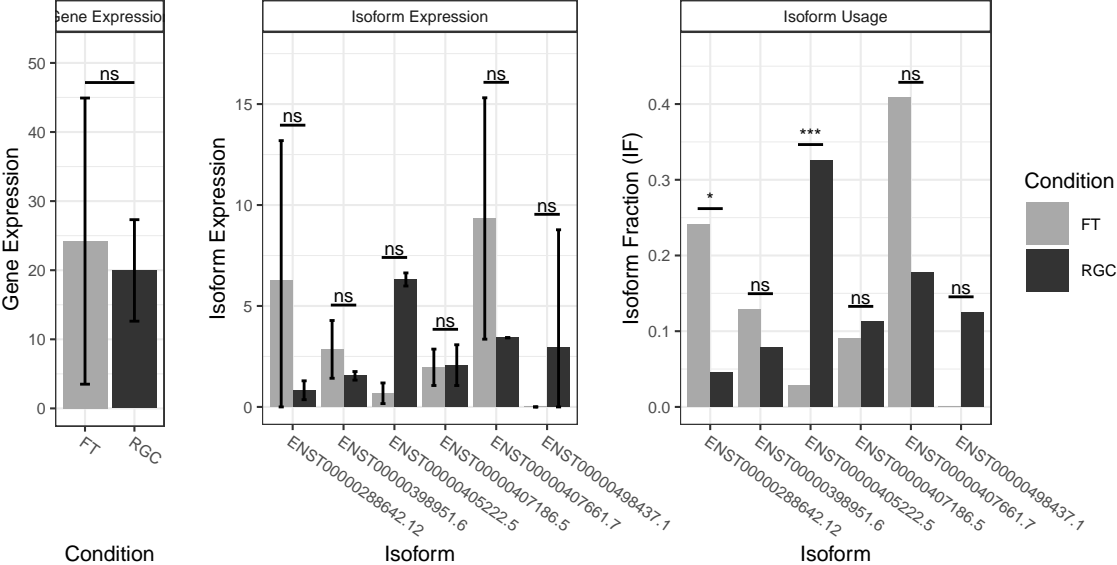
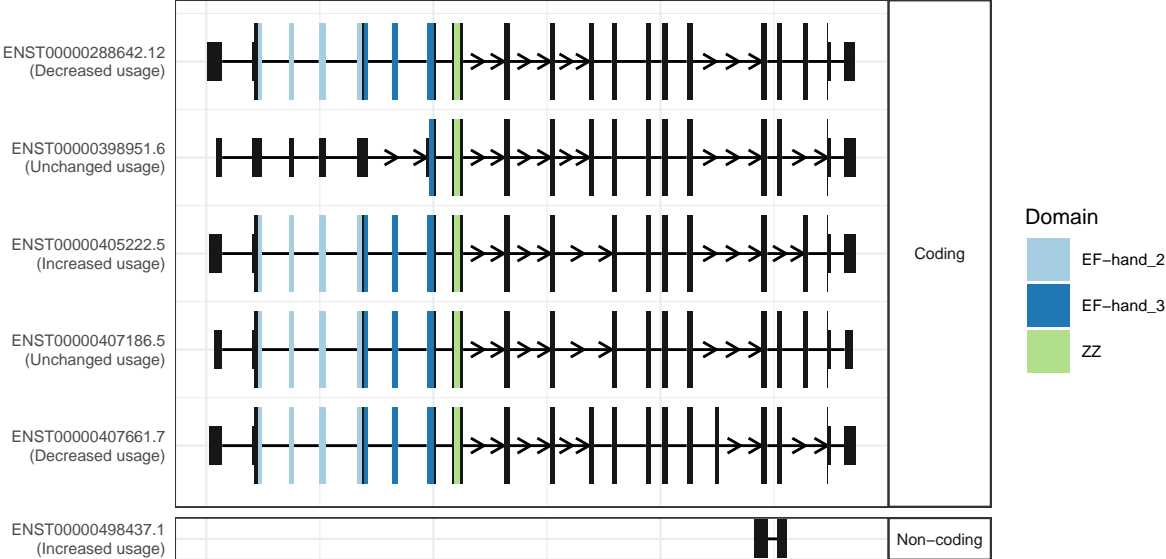


Glycos_transf_2



Ricin_B_lectin

The isoform switch in DTNB (FT vs RGC)



Domain



EF-hand_2

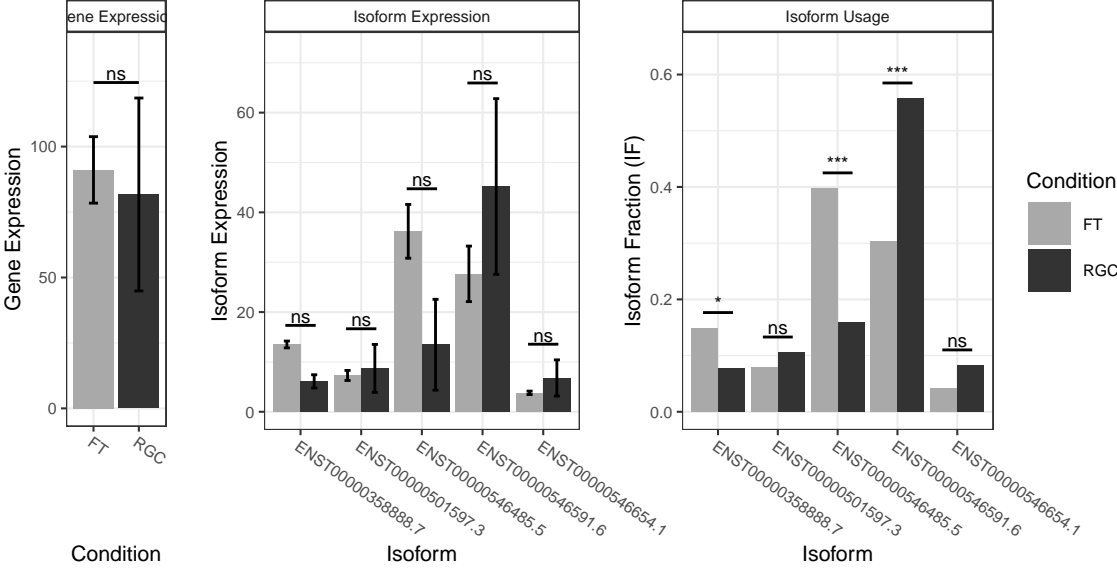
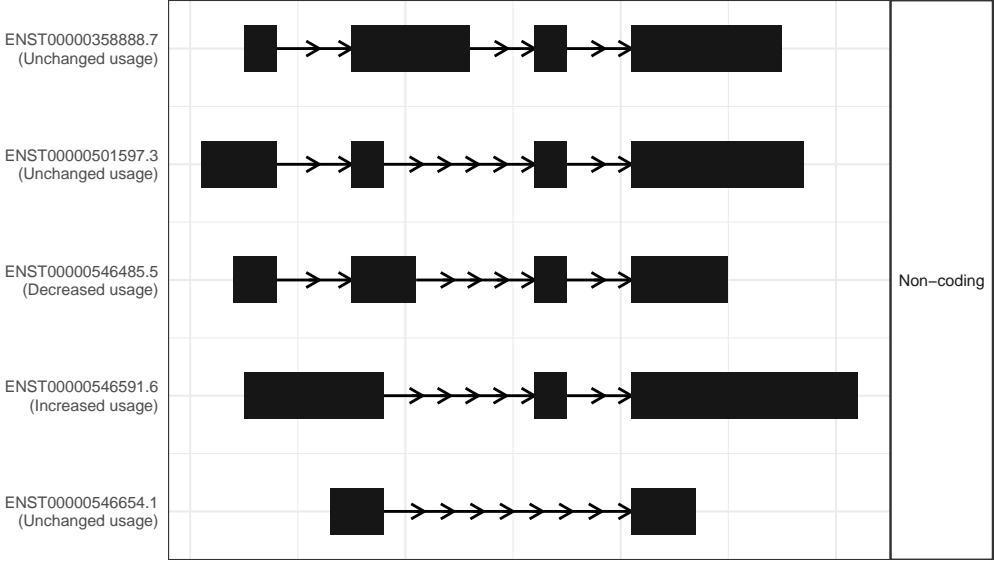


EF-hand_3

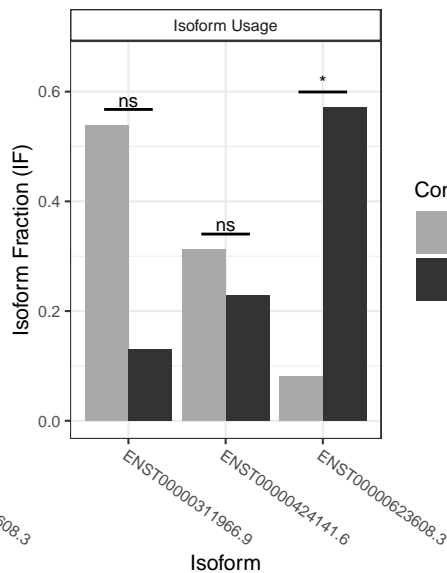
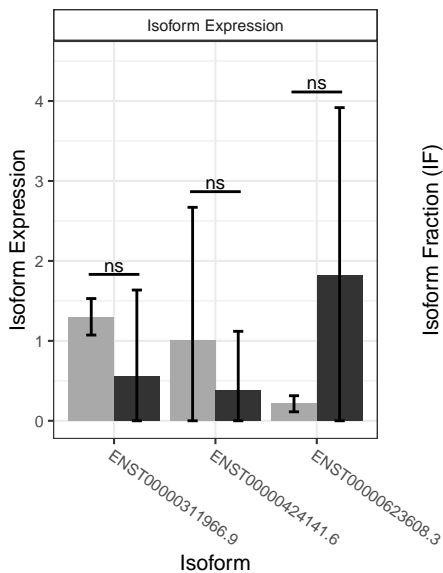
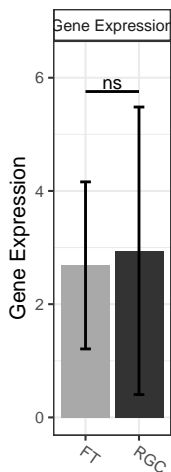
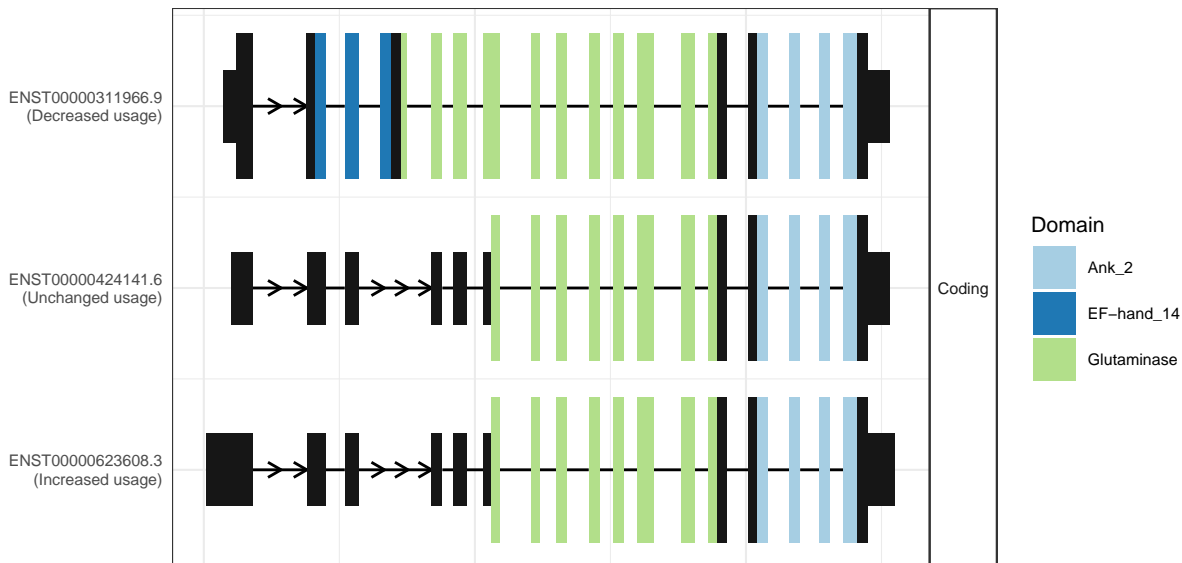


ZZ

The isoform switch in RPL41 (FT vs RGC)



The isoform switch in GLS2 (FT vs RGC)



Domain



Ank_2



EF-hand_14



Glutaminase

Domain

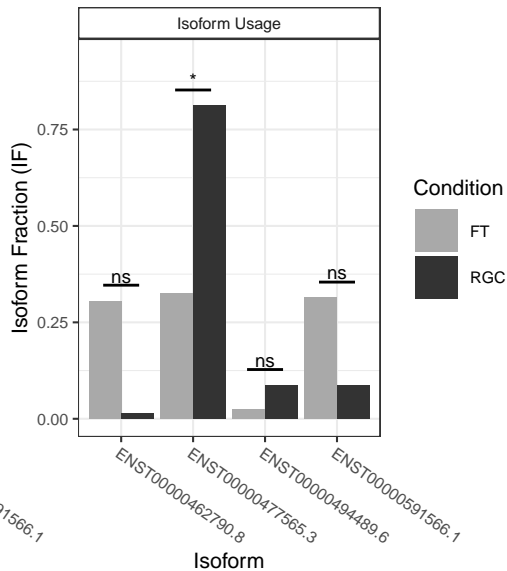
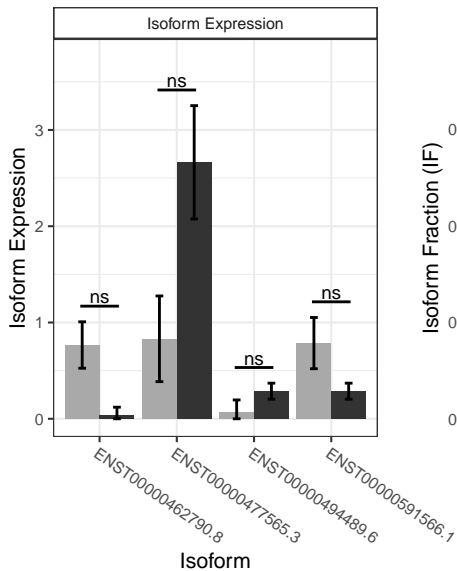
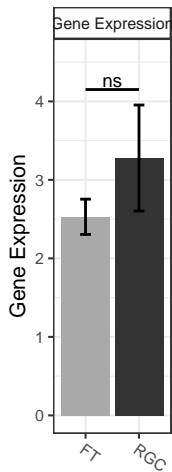
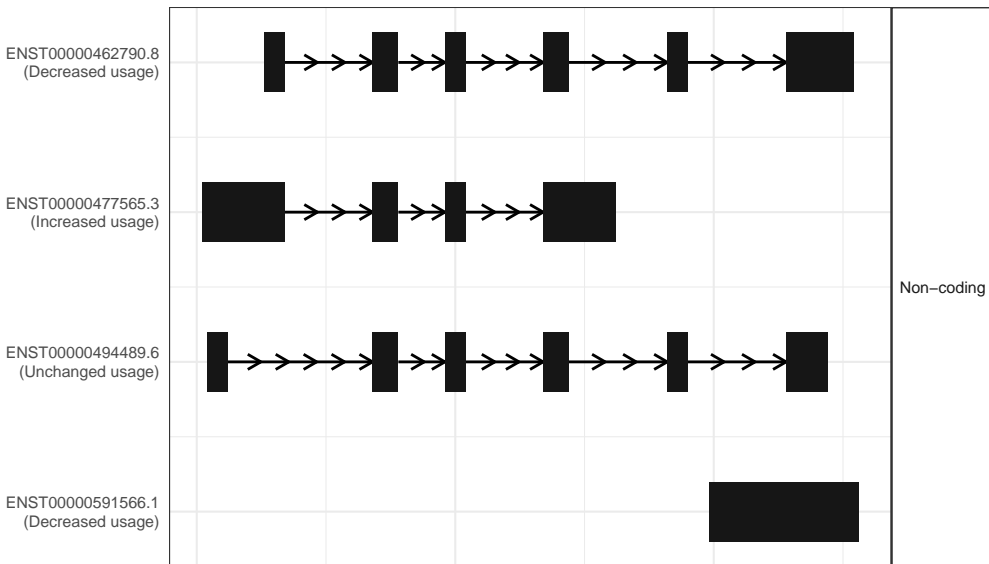


BAR

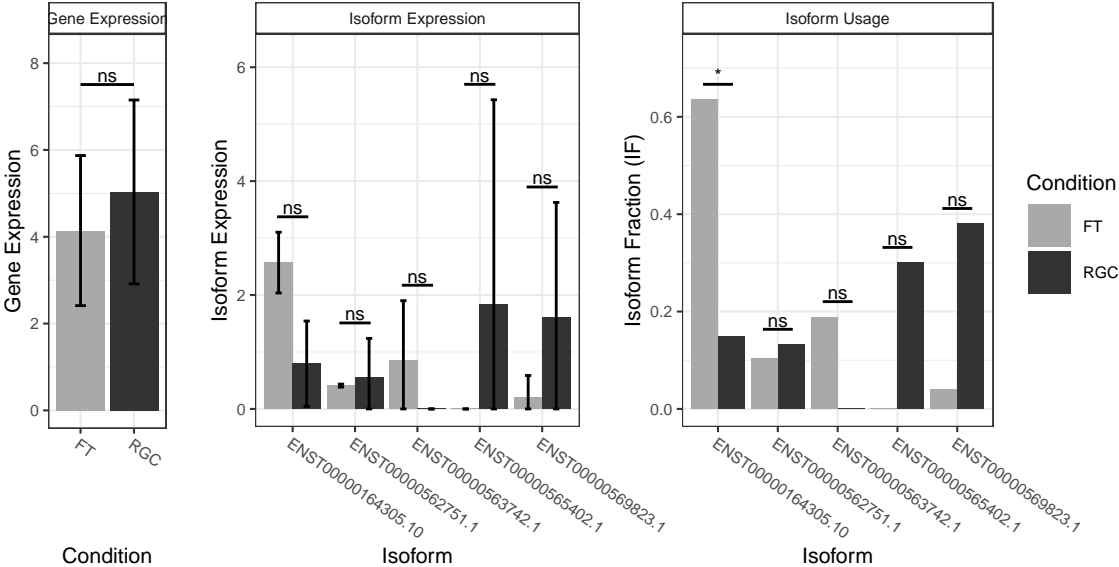
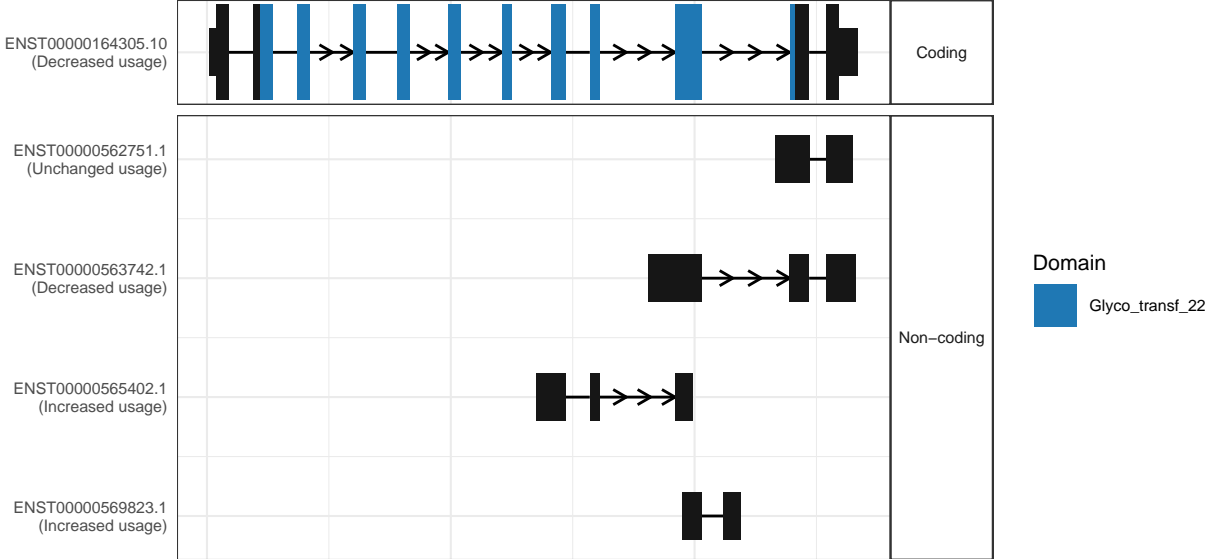


SH3_9

The isoform switch in BORCS8 (FT vs RGC)



The isoform switch in PIGB (FT vs RGC)

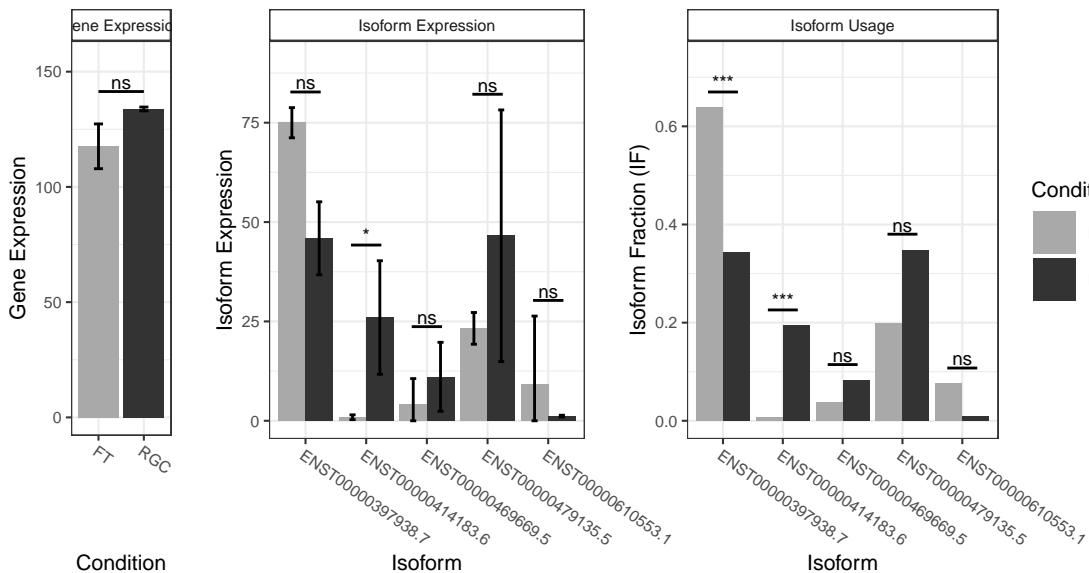
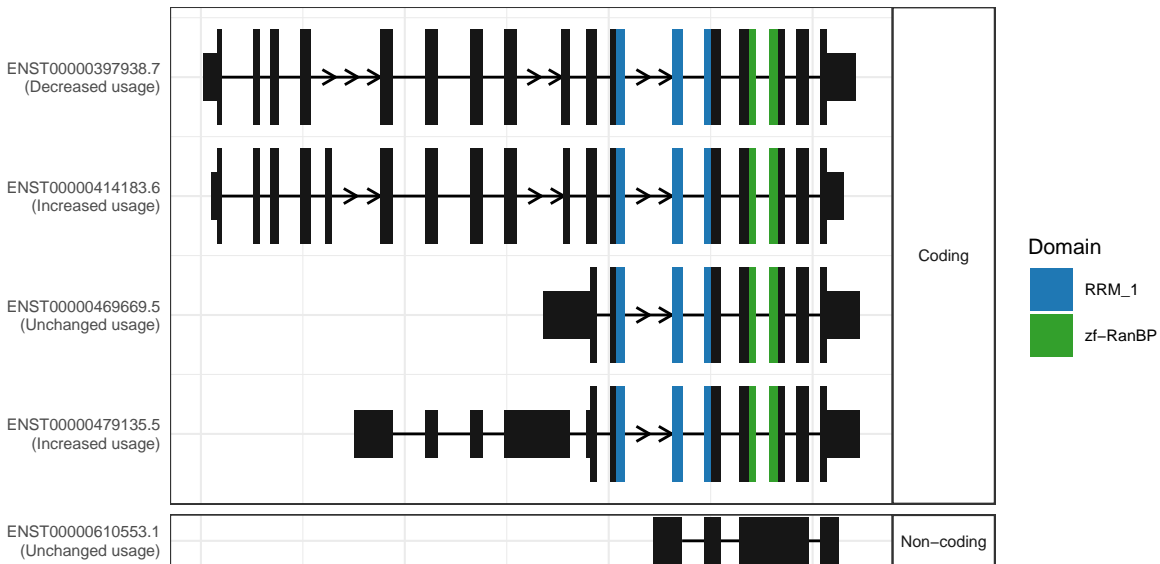


Domain



Glyco_transf_22

The isoform switch in EWSR1 (FT vs RGC)



Domain

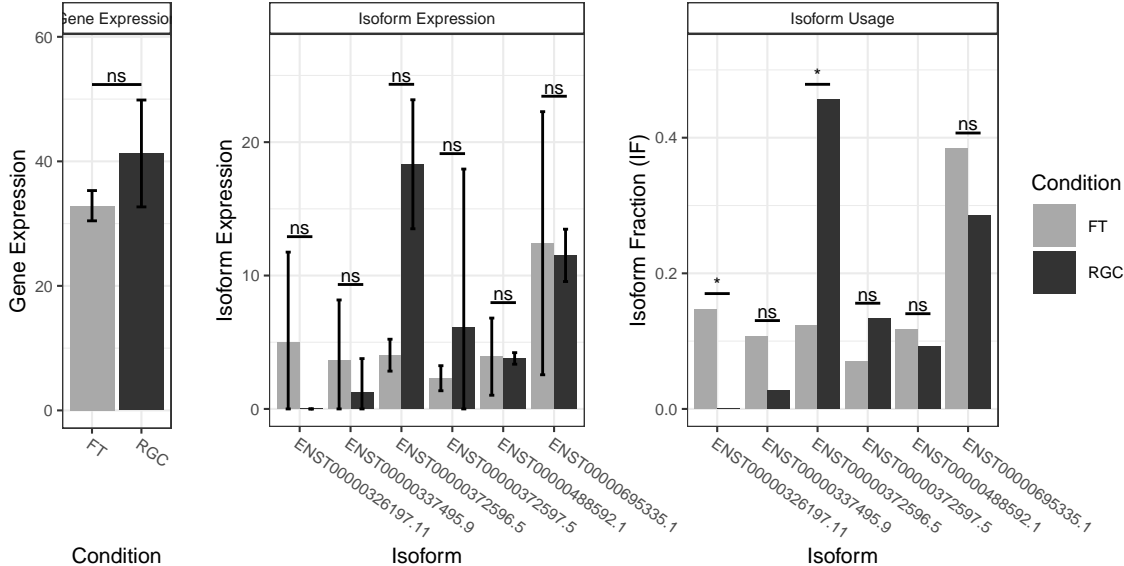
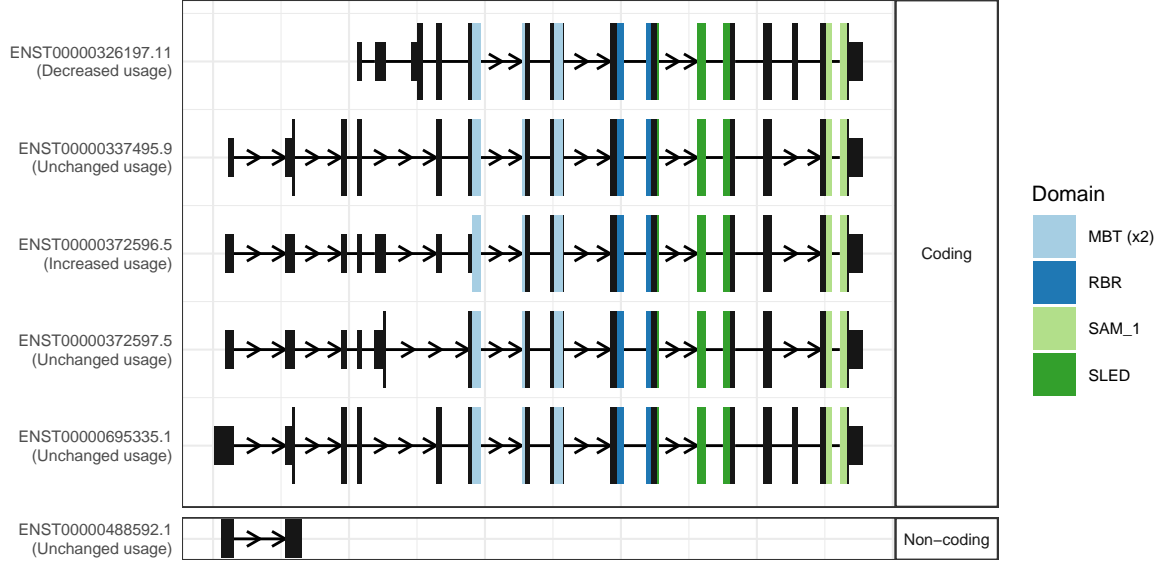


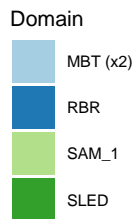
RRM_1



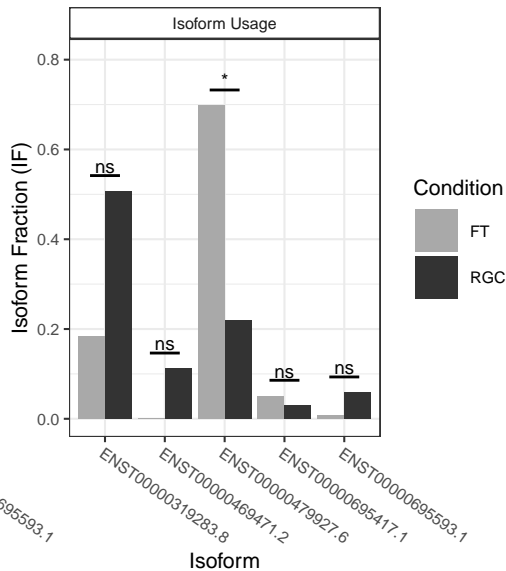
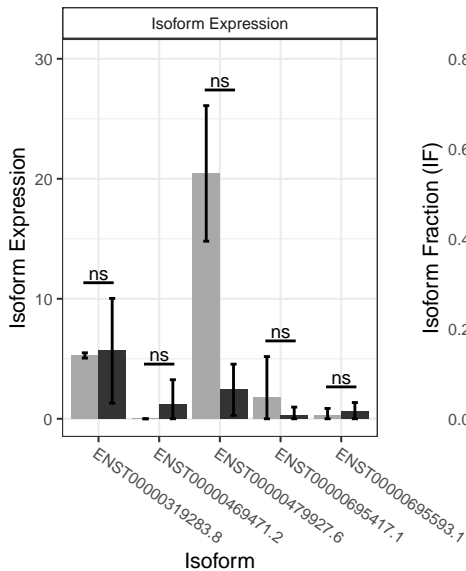
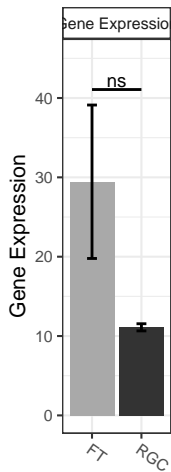
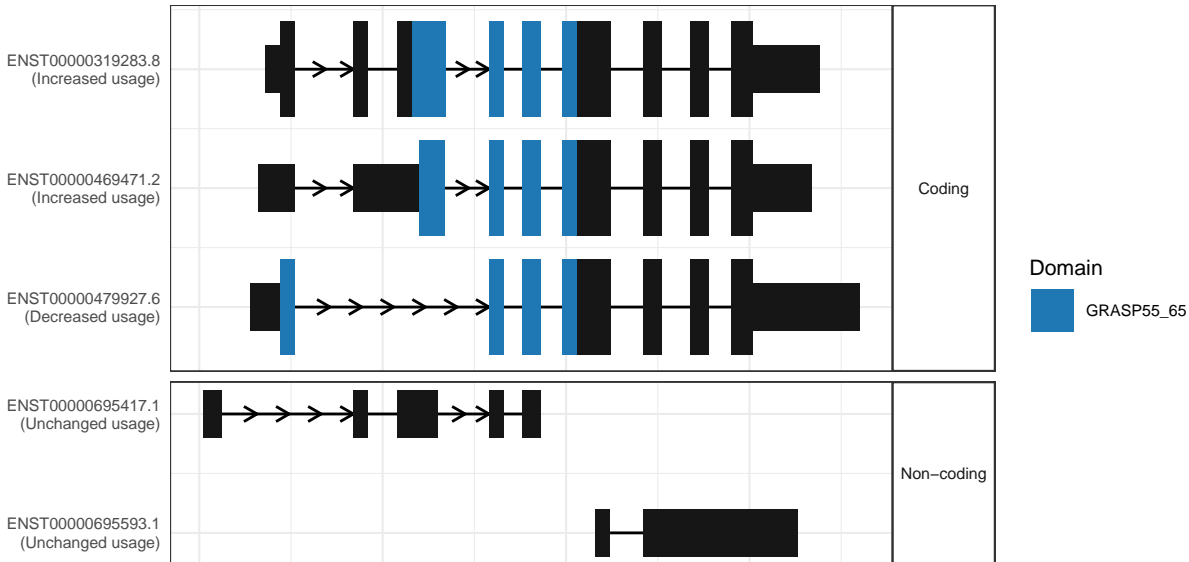
zf-RanBP

The isoform switch in SCMH1 (FT vs RGC)





The isoform switch in GORASP1 (FT vs RGC)

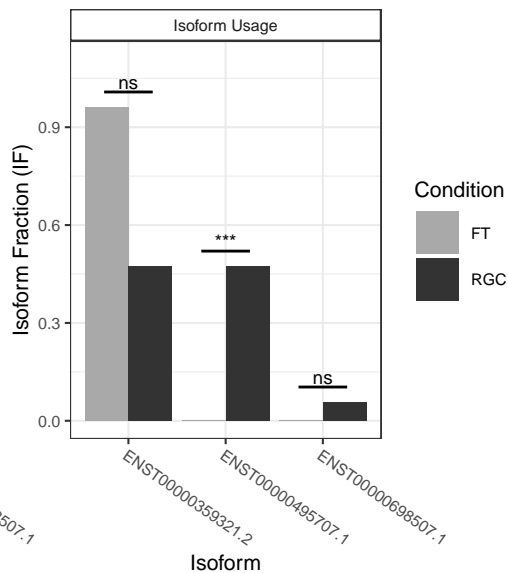
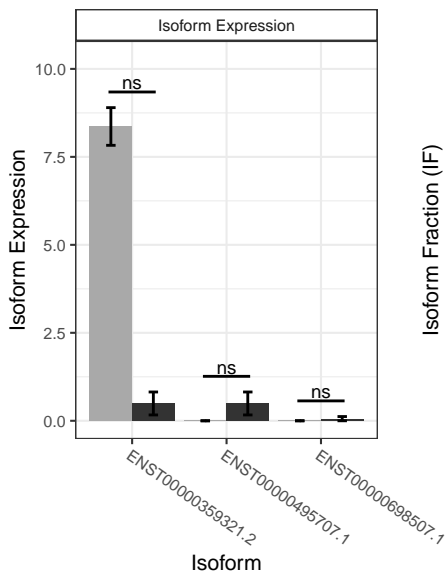
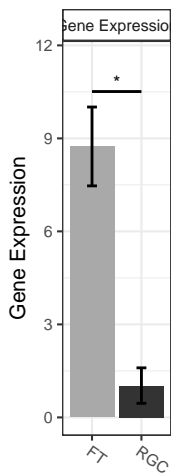
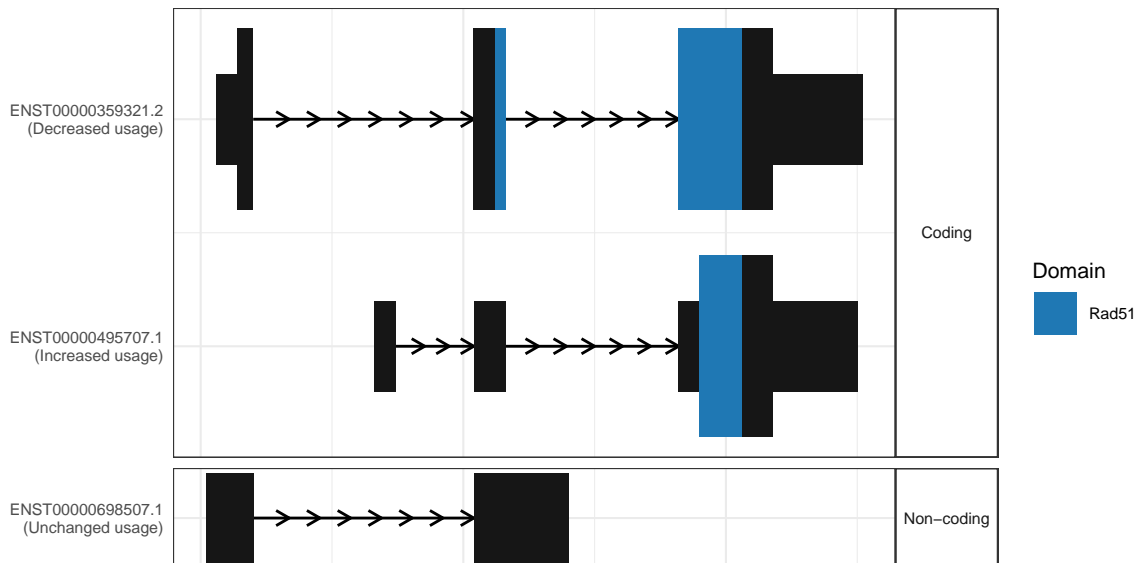


Domain



GRASP55_65

The isoform switch in XRCC2 (FT vs RGC)

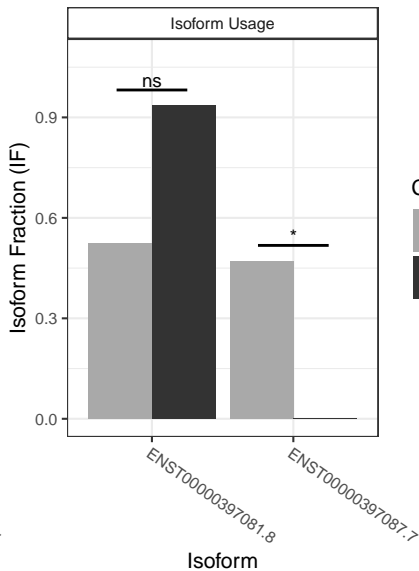
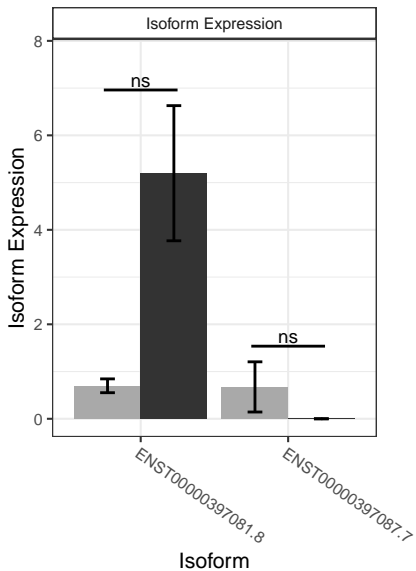
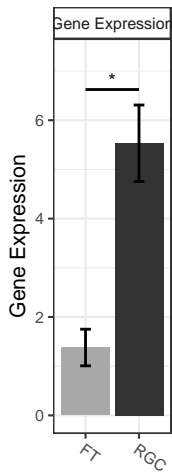
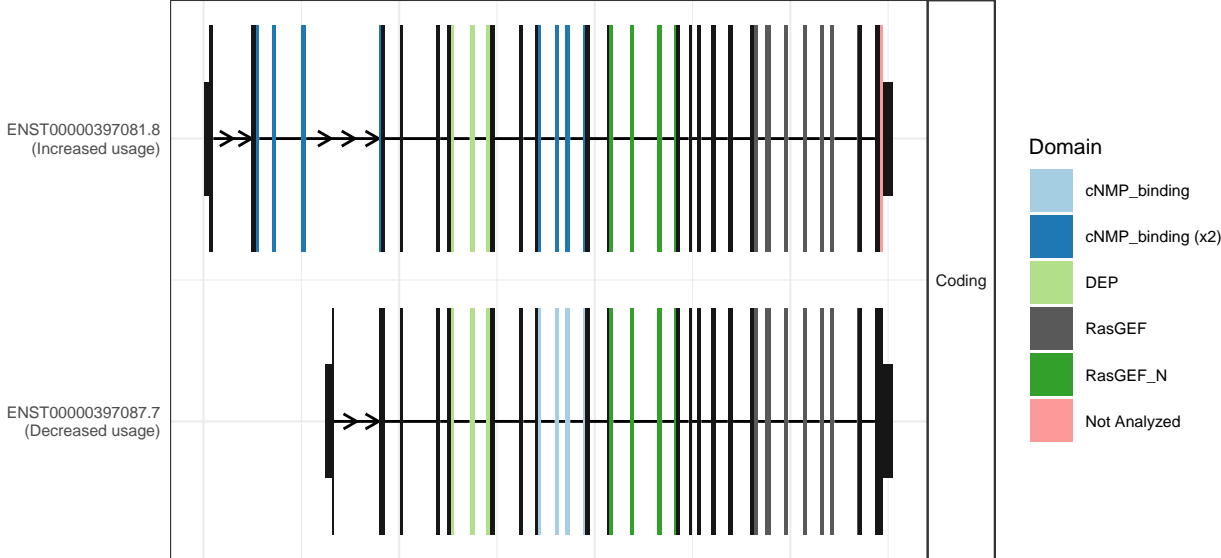


Domain









Rad51

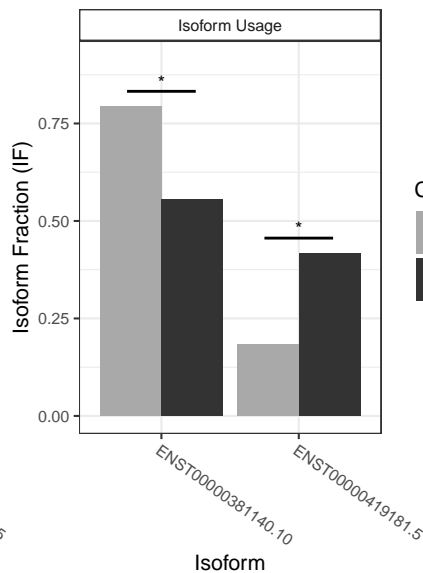
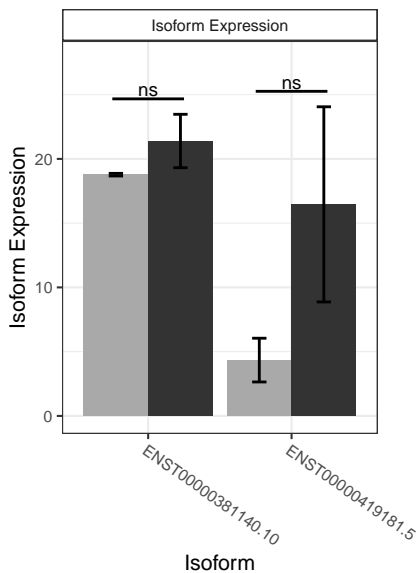
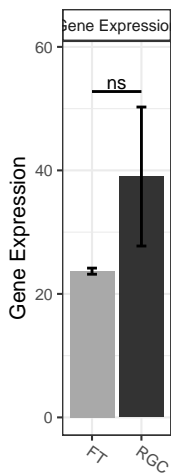
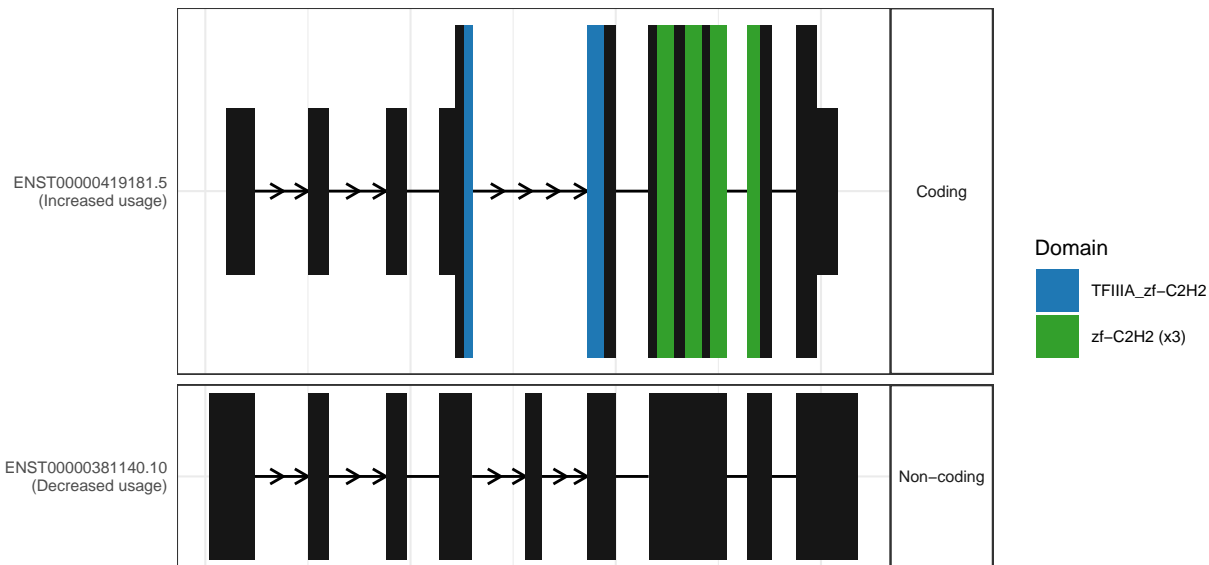
The isoform switch in RAPGEF4 (FT vs RGC)



Domain

	cNMP_binding
	cNMP_binding (x2)
	DEP
	RasGEF
	RasGEF_N
	Not Analyzed

The isoform switch in GTF3A (FT vs RGC)



Domain

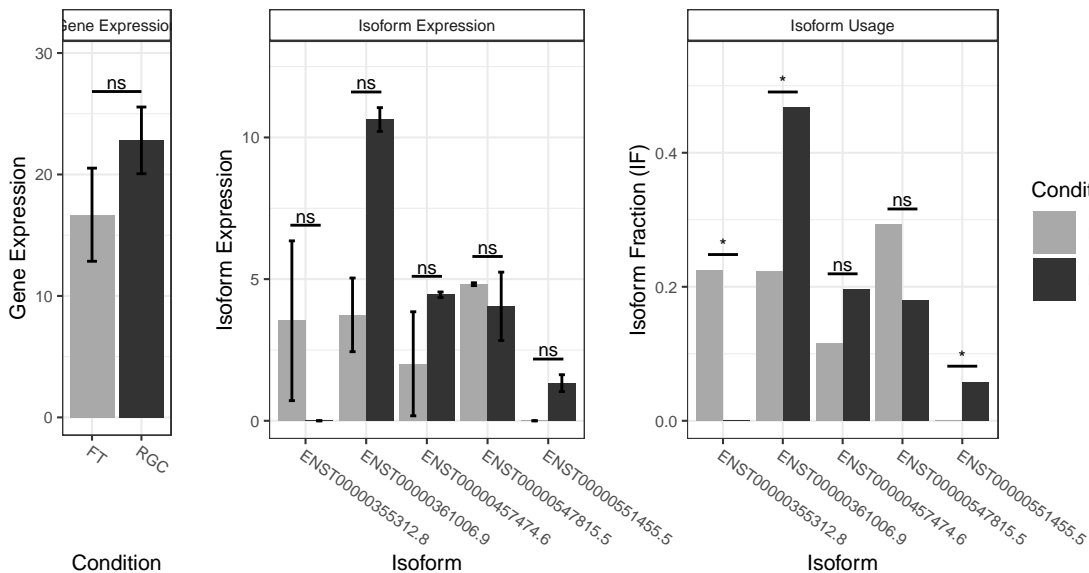
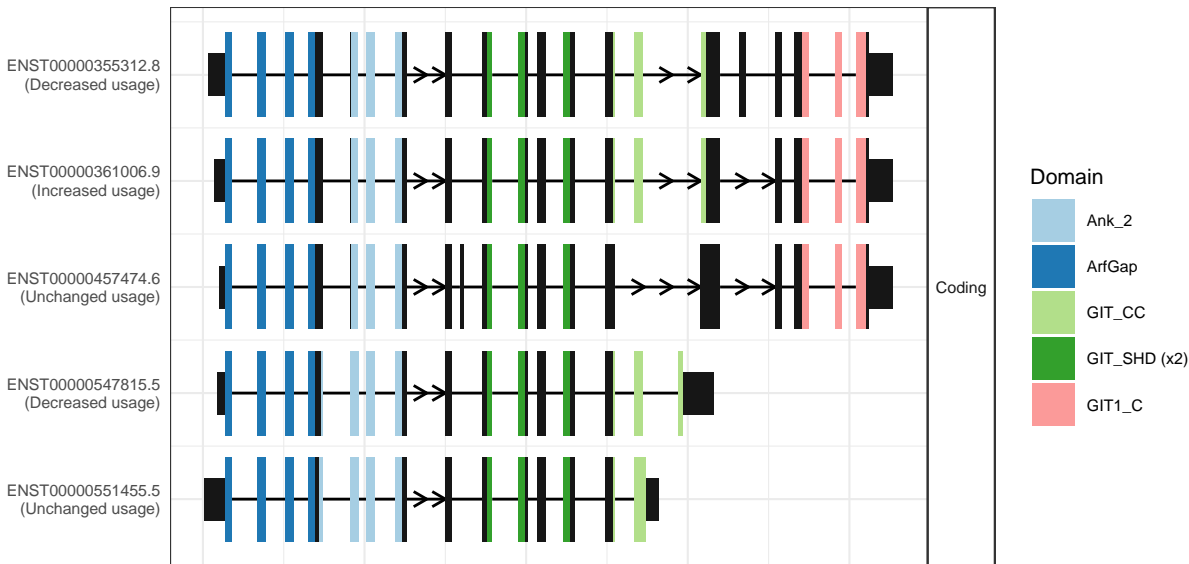


TFIIIA_zf-C2H2



zf-C2H2 (x3)

The isoform switch in GIT2 (FT vs RGC)



Domain



Ank_2



ArfGap



GIT_CC

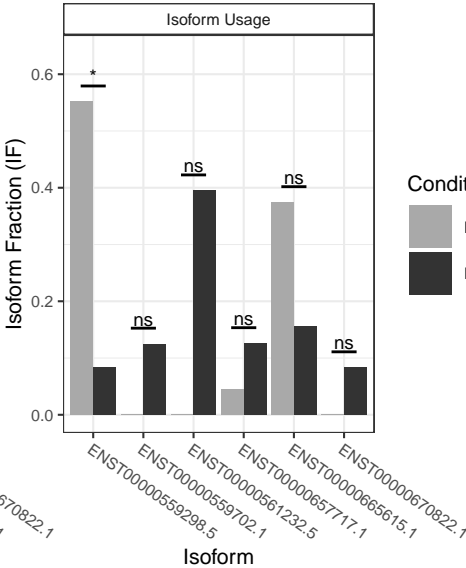
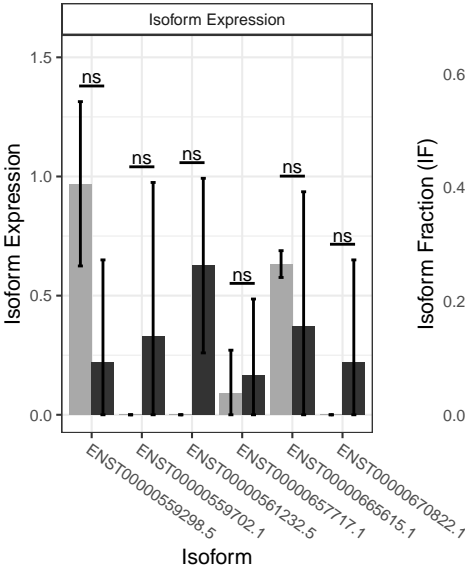
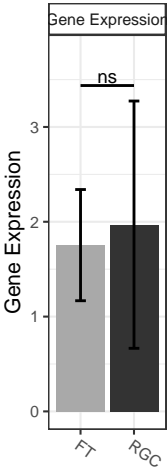
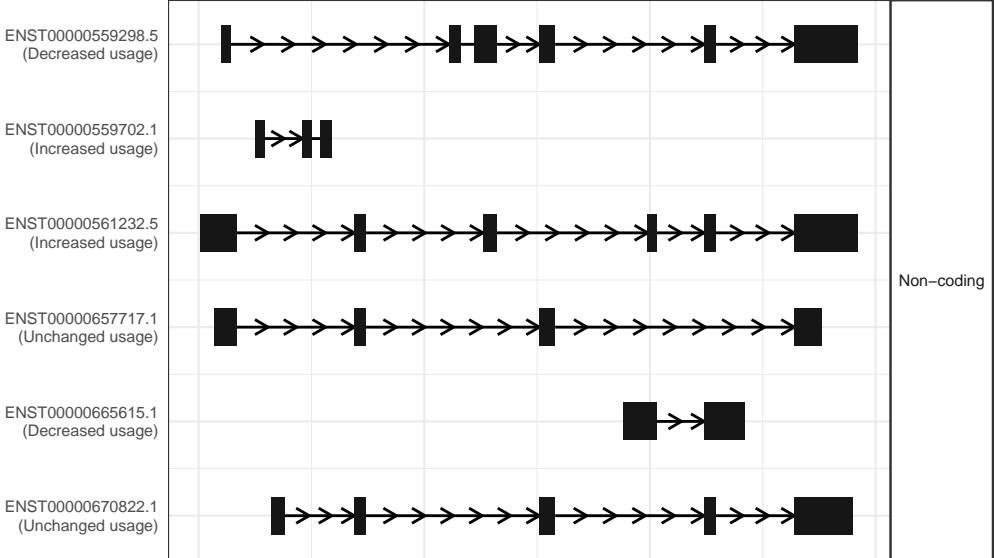


GIT_SHD (x2)



GIT1_C

The isoform switch in IQCH-AS1 (FT vs RGC)

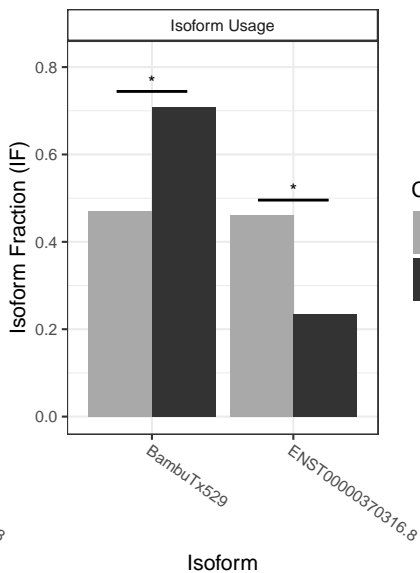
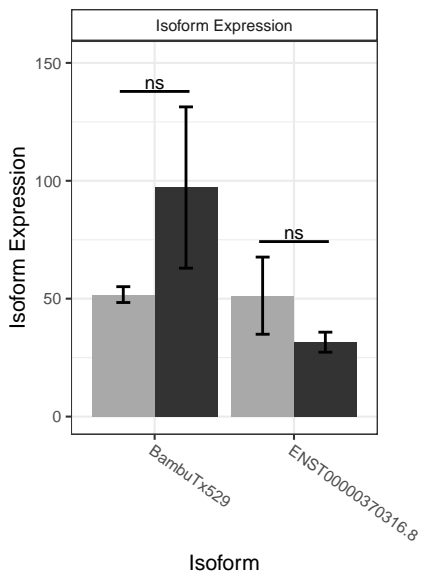
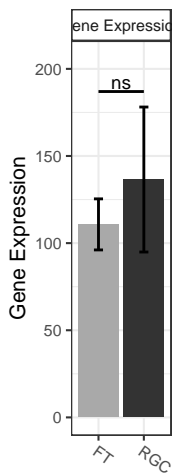
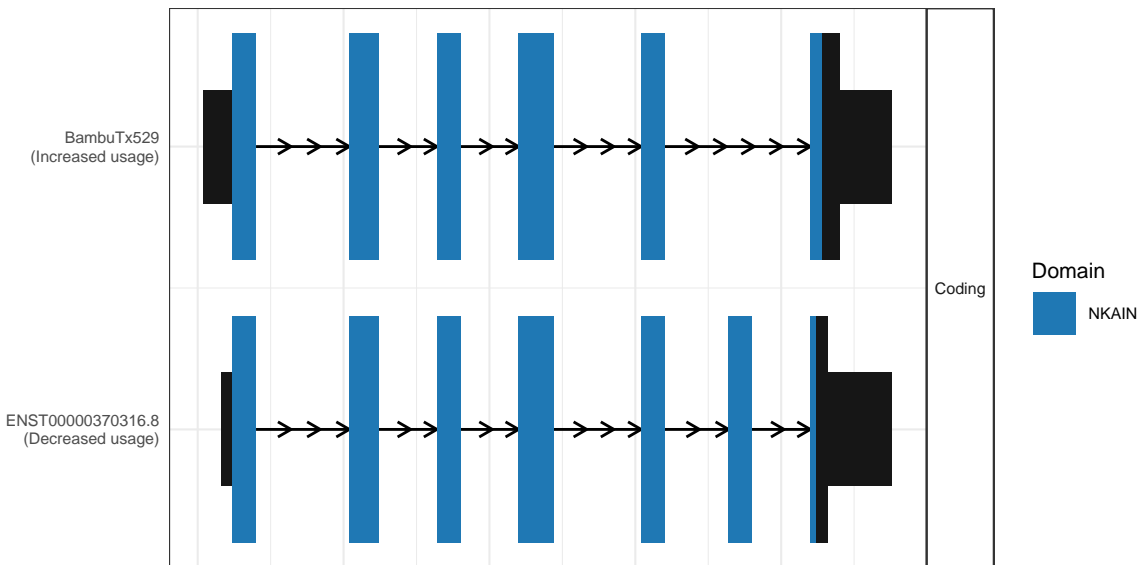


Condition

FT

RGC

The isoform switch in NKAIN4 (FT vs RGC)

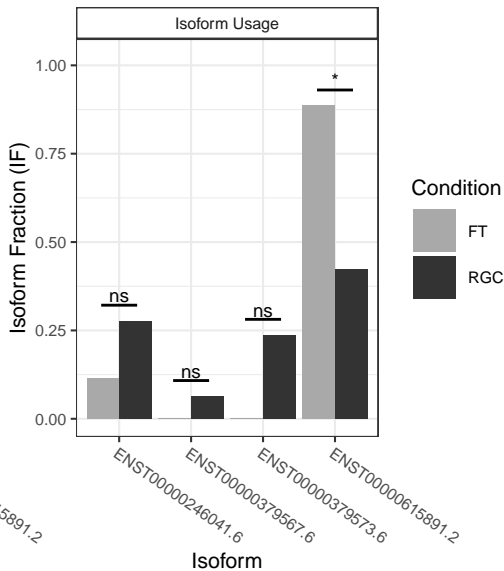
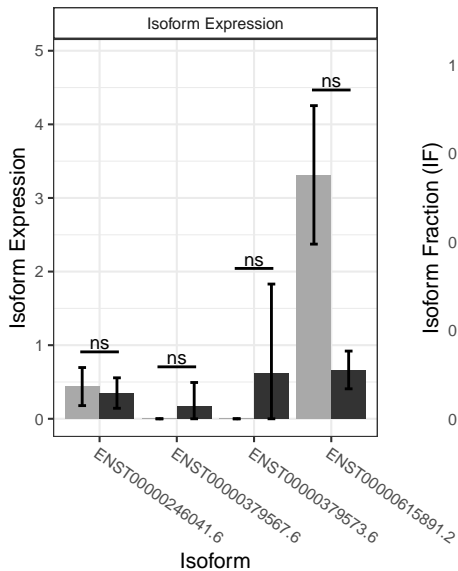
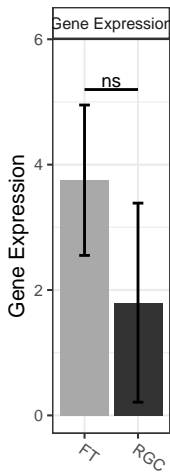
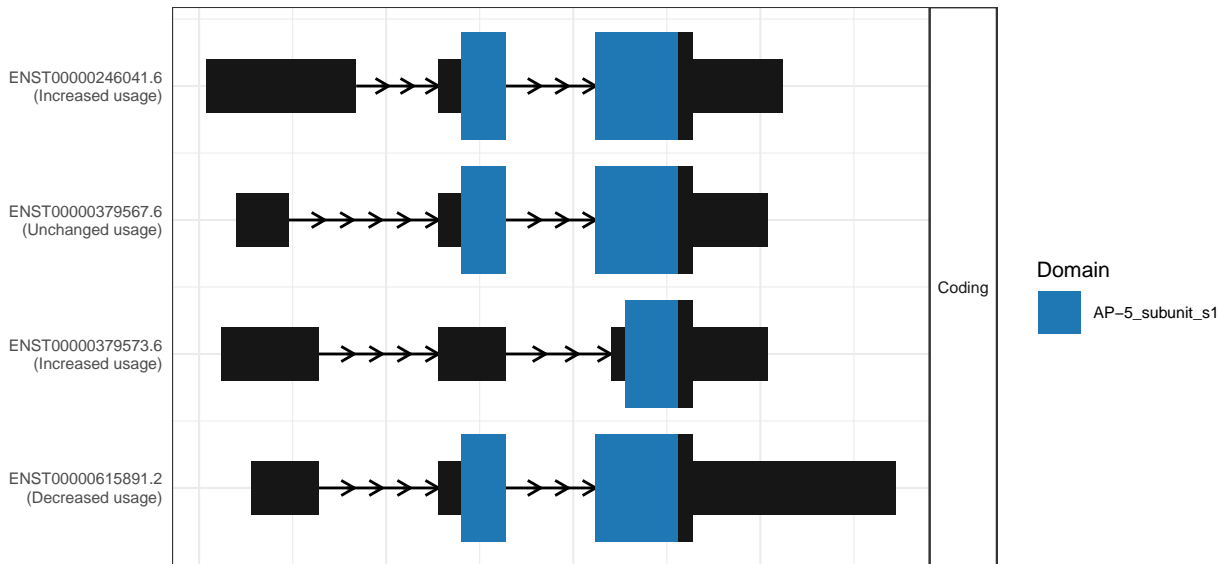


Domain




NKAIN

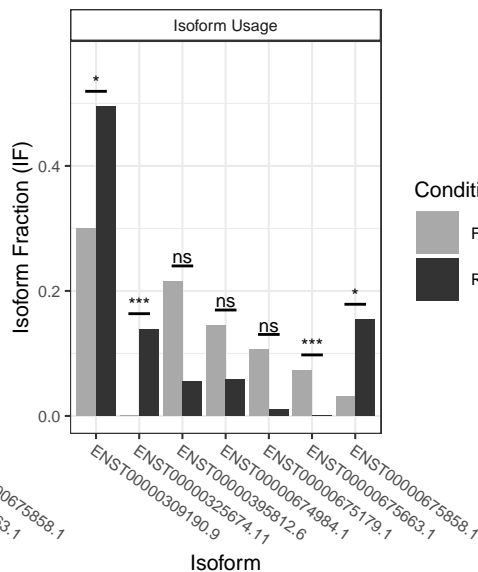
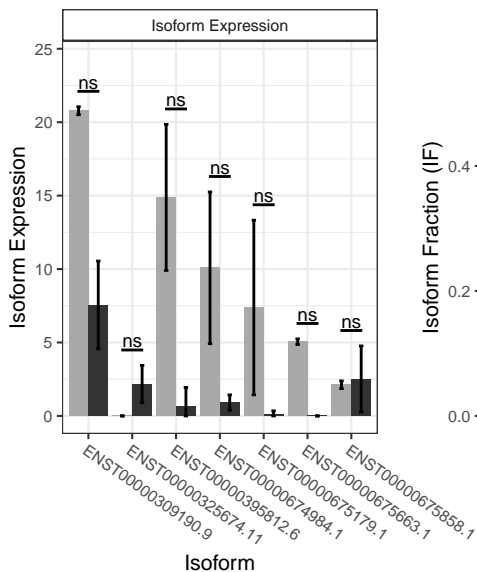
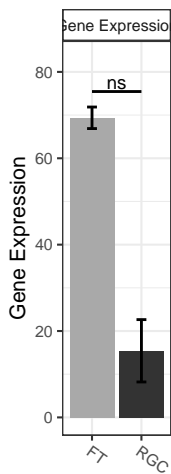
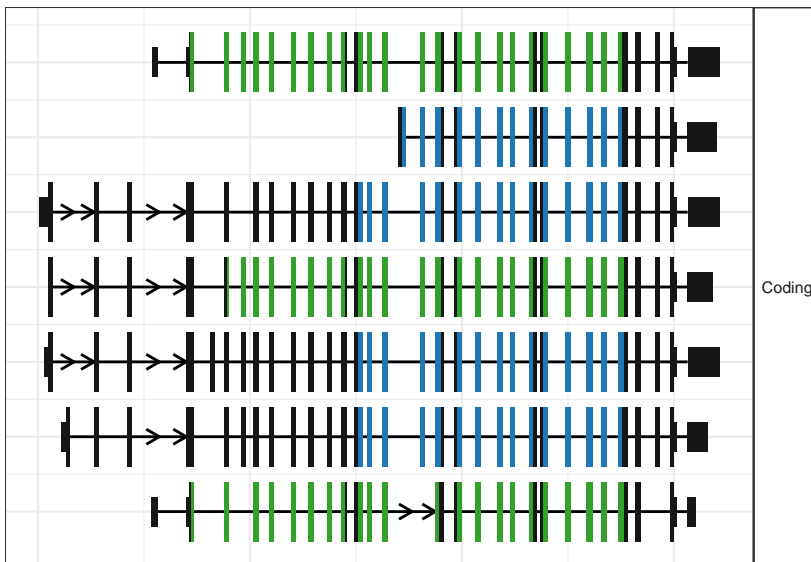
The isoform switch in AP5S1 (FT vs RGC)



Domain

 AP-5_subunit_s1

The isoform switch in CAST (FT vs RGC)



Domain

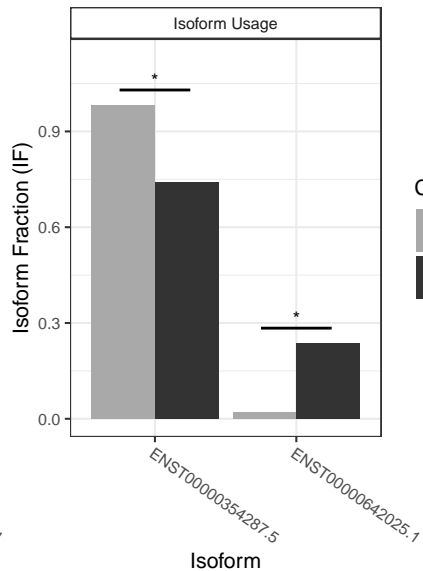
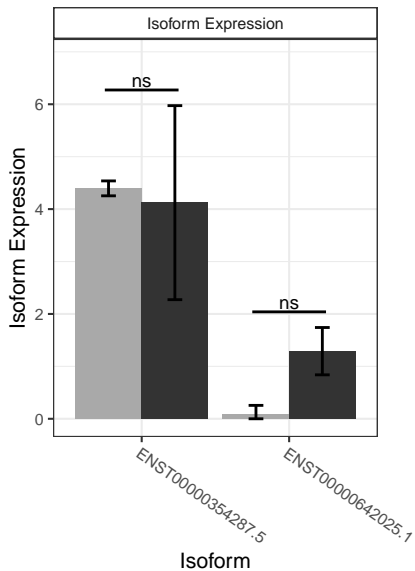
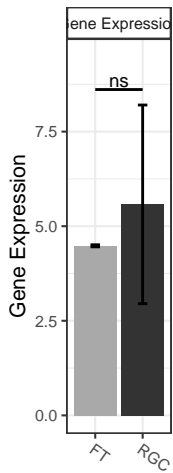
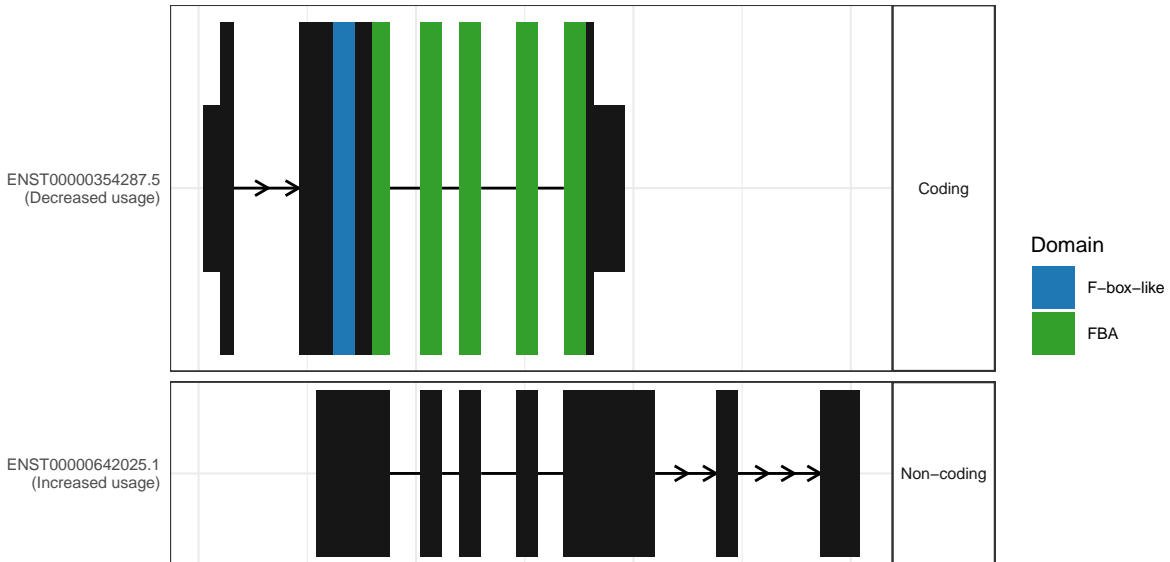


Calpain_inhib (x3)



Calpain_inhib (x4)

The isoform switch in FBXO2 (FT vs RGC)



Domain

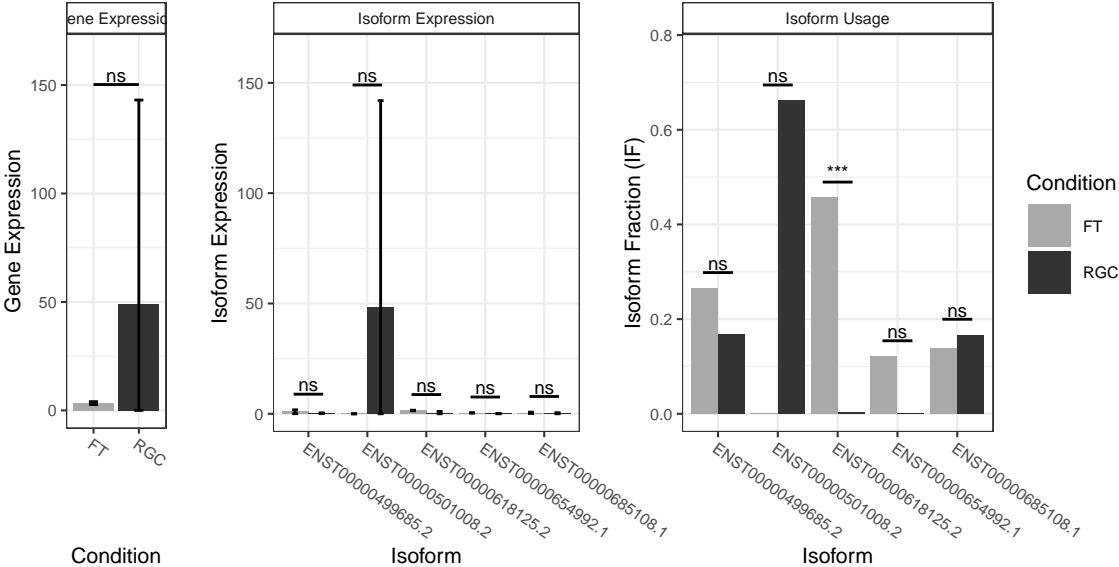
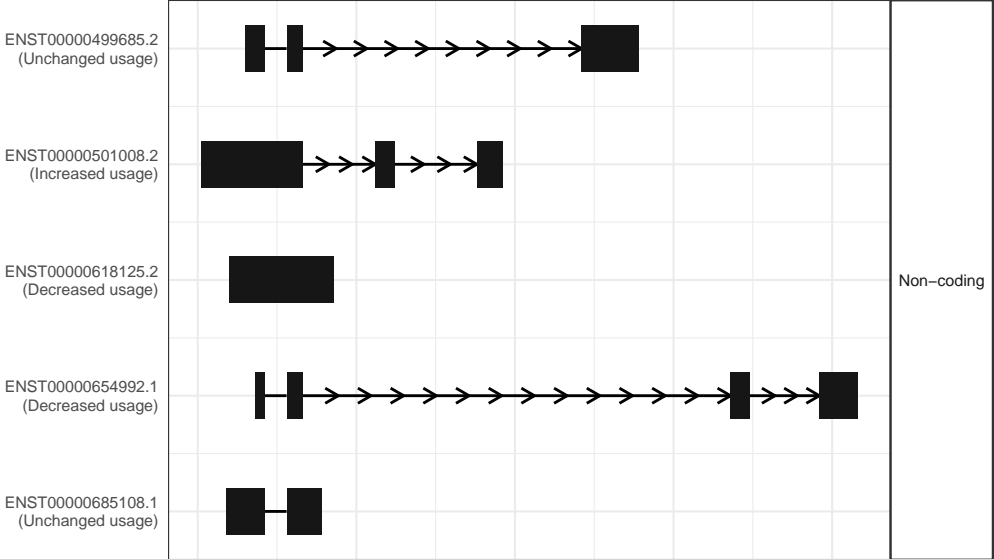


F-box-like

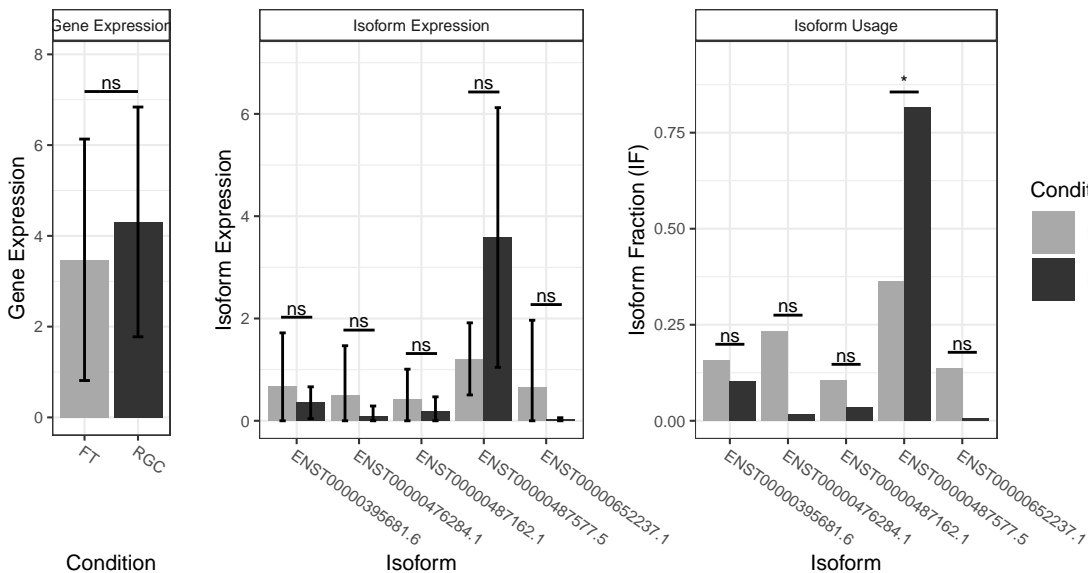
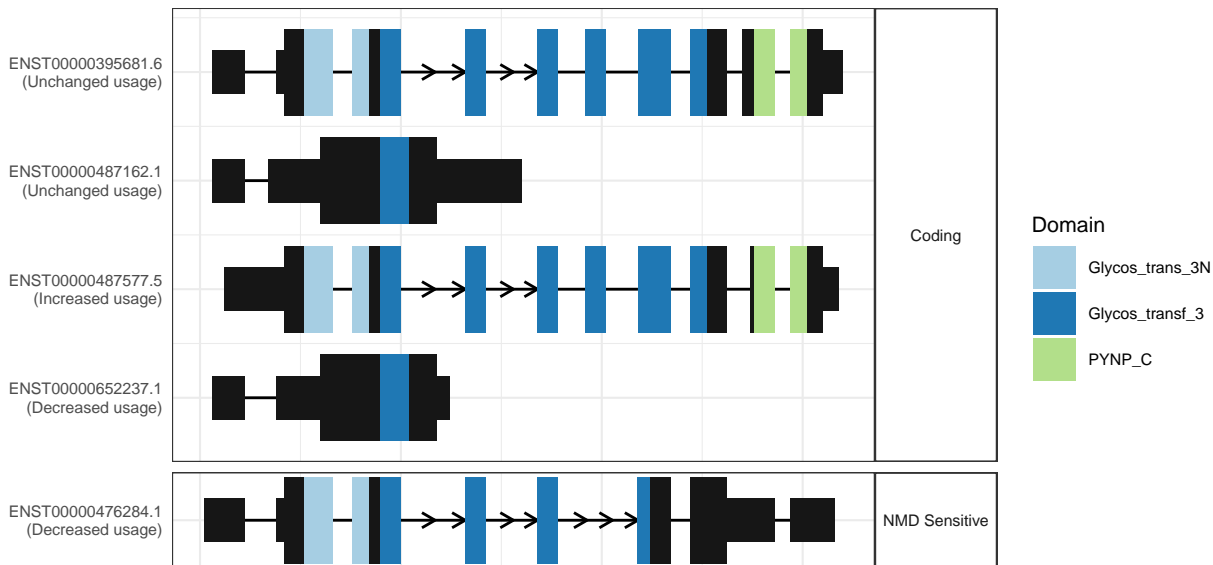


FBA




The isoform switch in BTG1-DT (FT vs RGC)



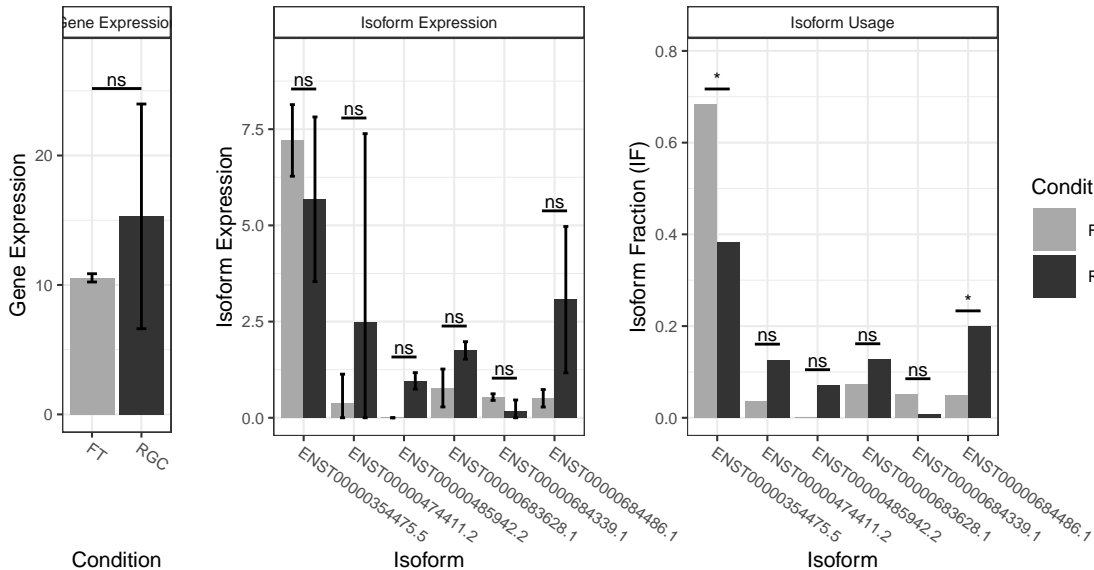
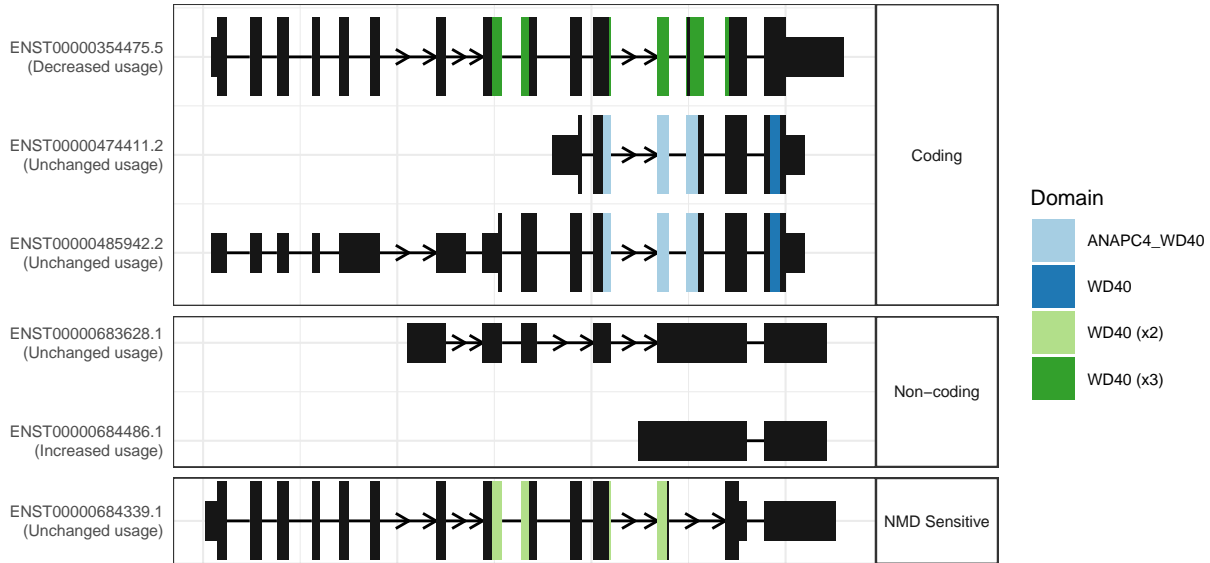
The isoform switch in TYMP (FT vs RGC)






Domain

	Glycos_trans_3N
	Glycos_transf_3
	PYNP_C

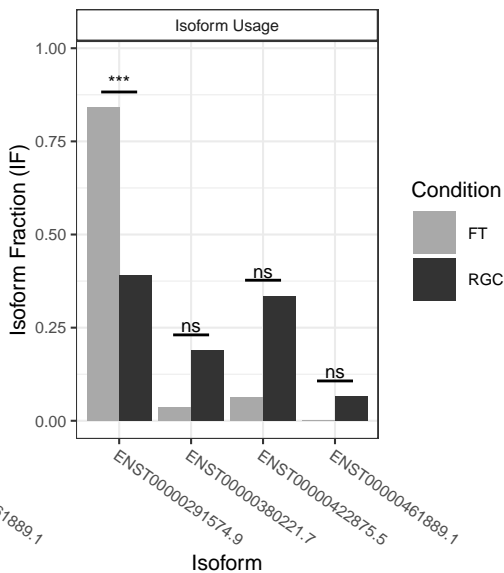
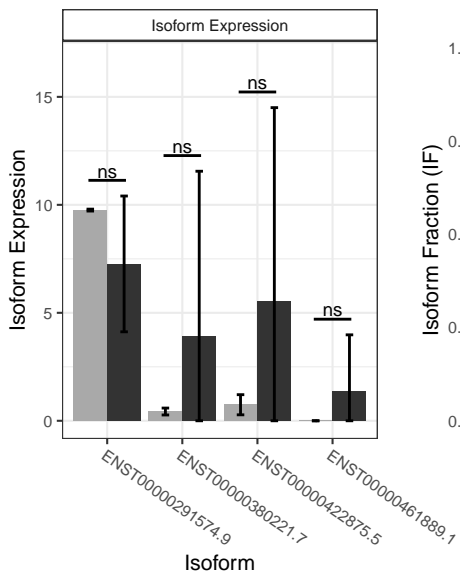
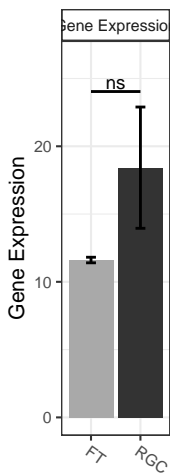
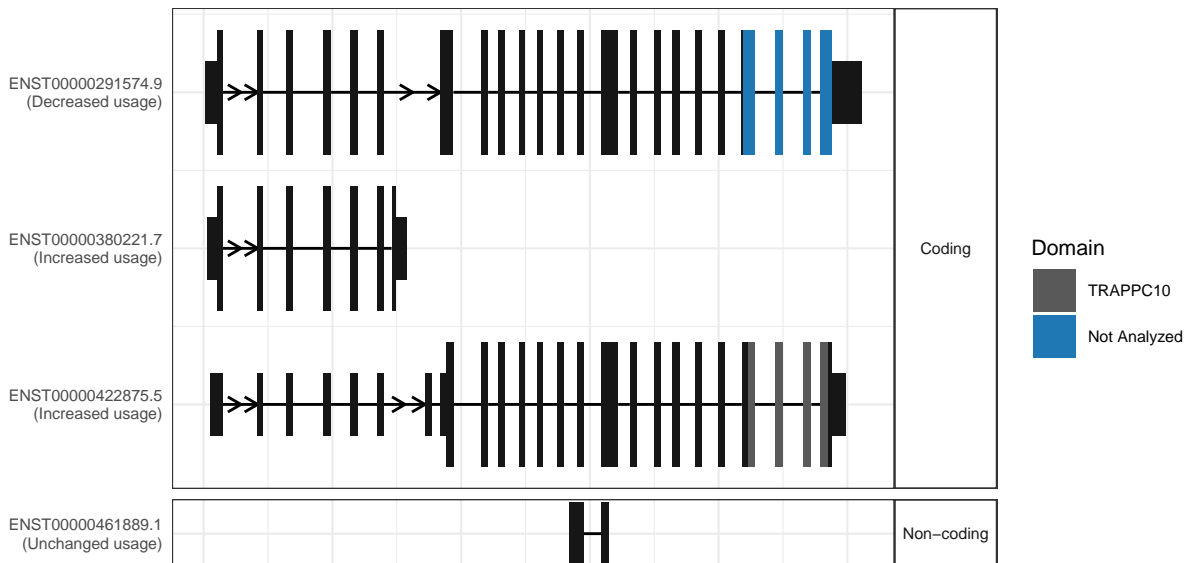
The isoform switch in WDR91 (FT vs RGC)



Domain

	ANAPC4_WD40
	WD40
	WD40 (x2)
	WD40 (x3)

The isoform switch in TRAPPC10 (FT vs RGC)



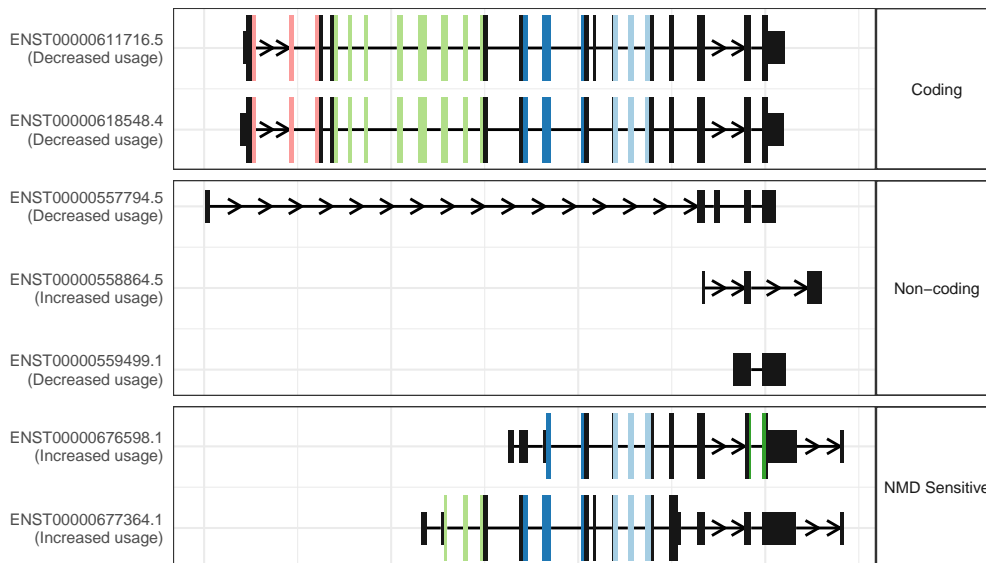
Domain



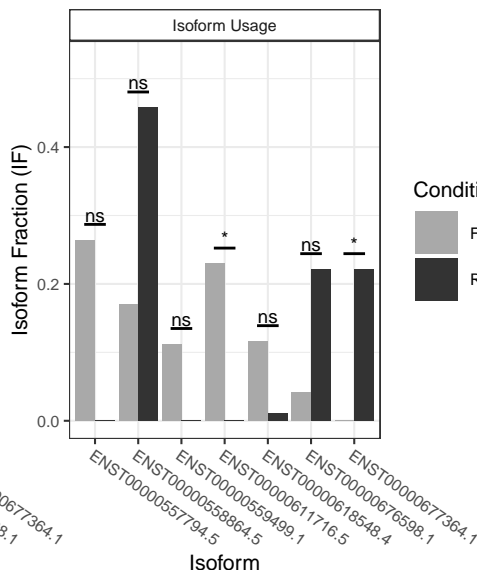
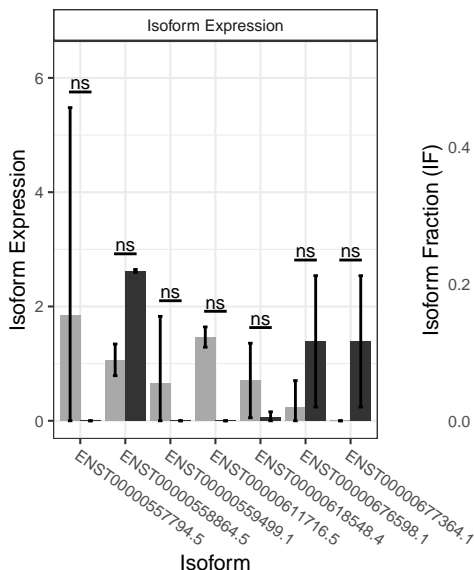
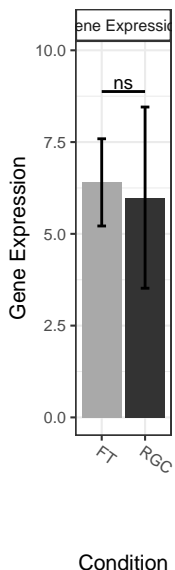
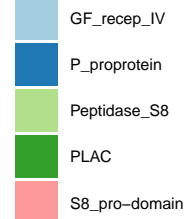
TRAPPC10

Not Analyzed

The isoform switch in PCSK6 (FT vs RGC)



Domain



Domain



GF_recep_IV



P_proprotein



Peptidase_S8

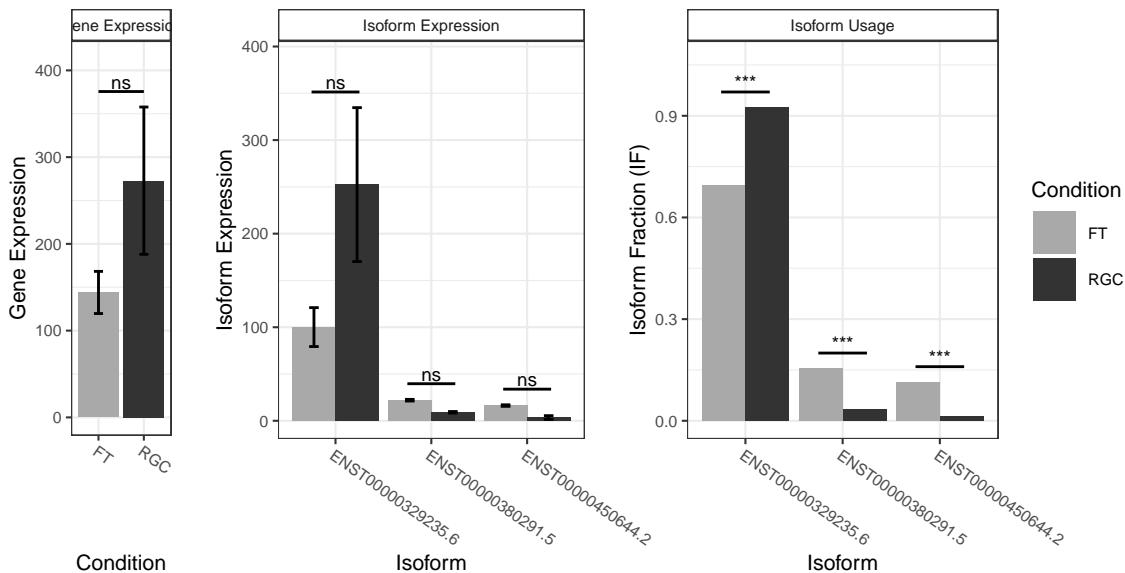
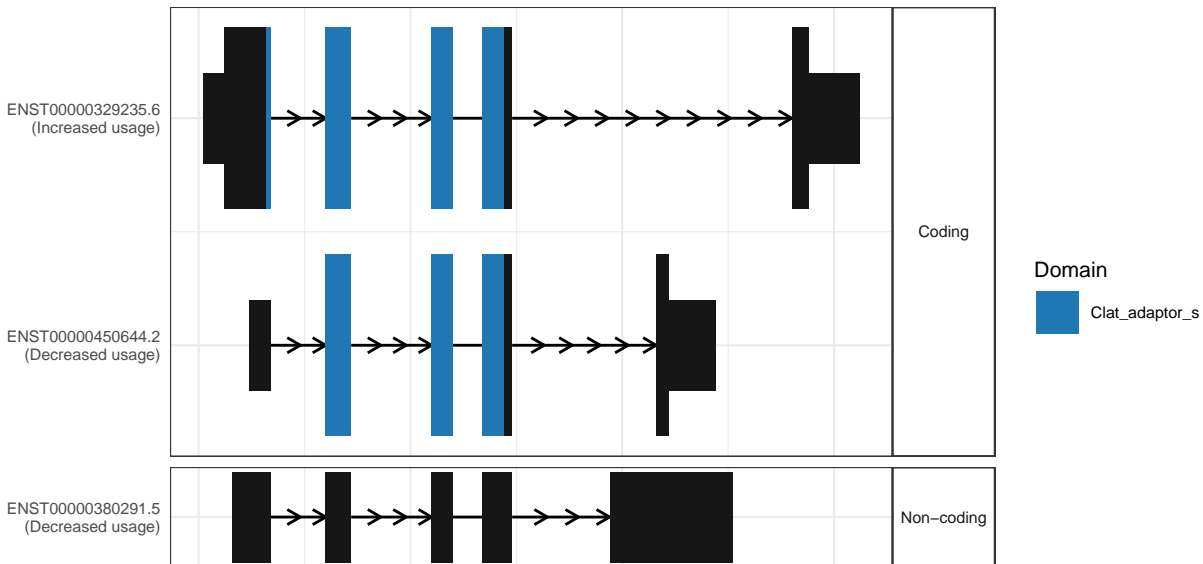


PLAC



S8_pro-domain

The isoform switch in AP1S2 (FT vs RGC)

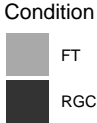
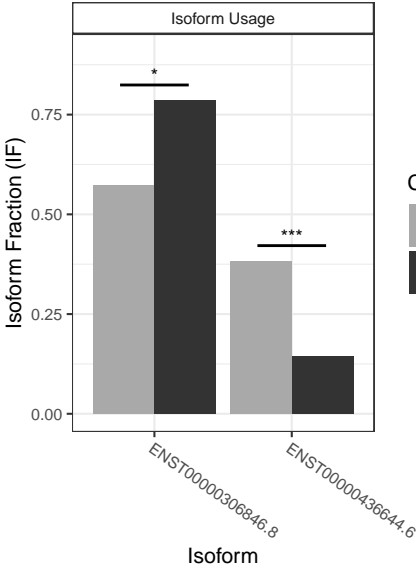
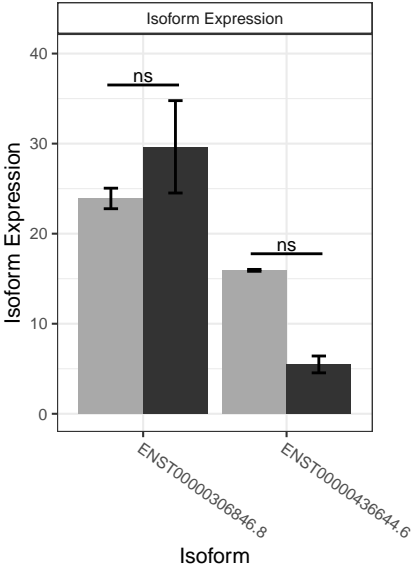
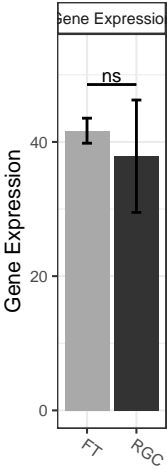
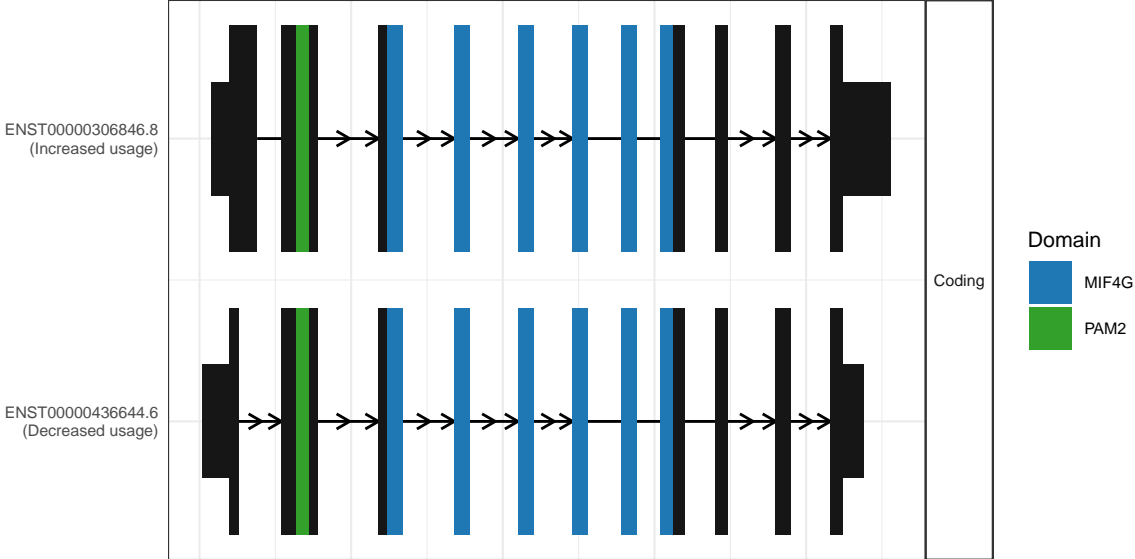


Domain



Clat_adaptor_s

The isoform switch in PAIP1 (FT vs RGC)



Domain

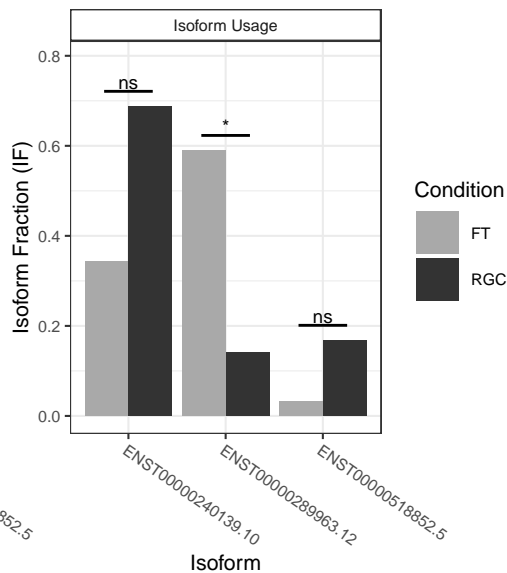
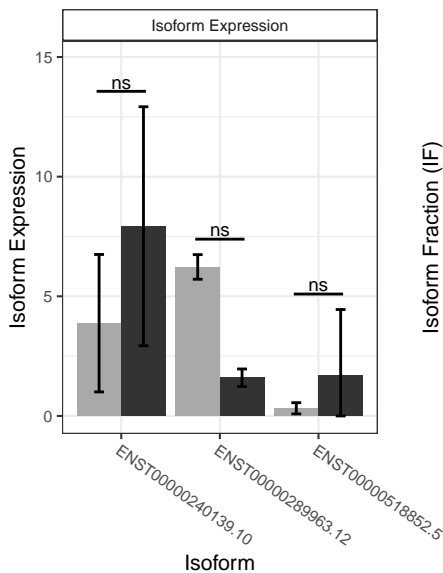
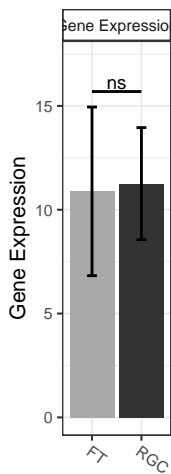
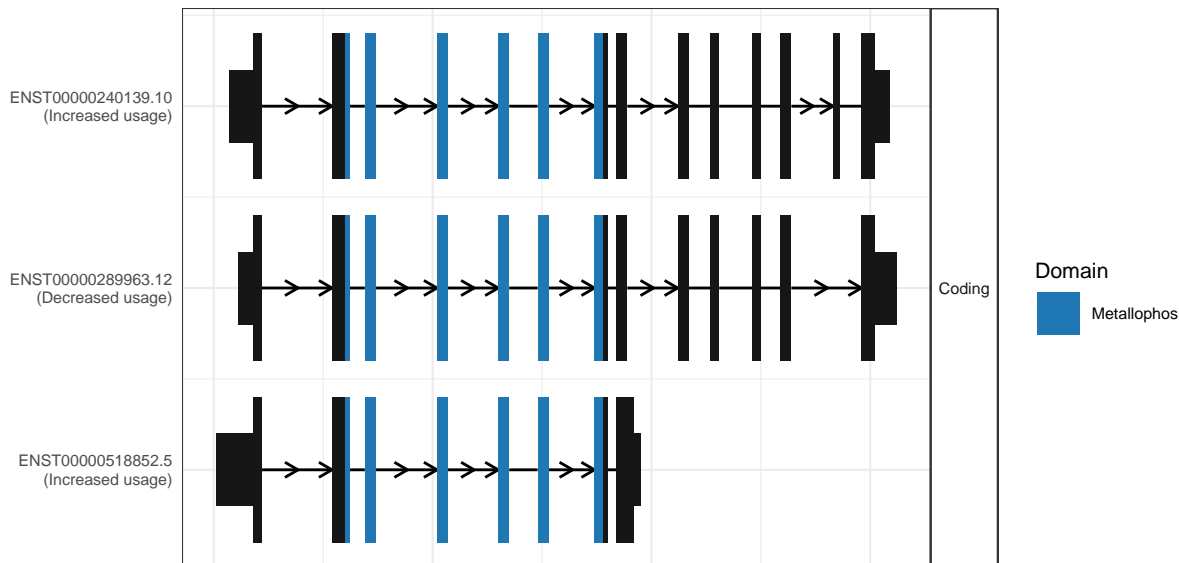


MIF4G



PAM2

The isoform switch in PPP3CC (FT vs RGC)

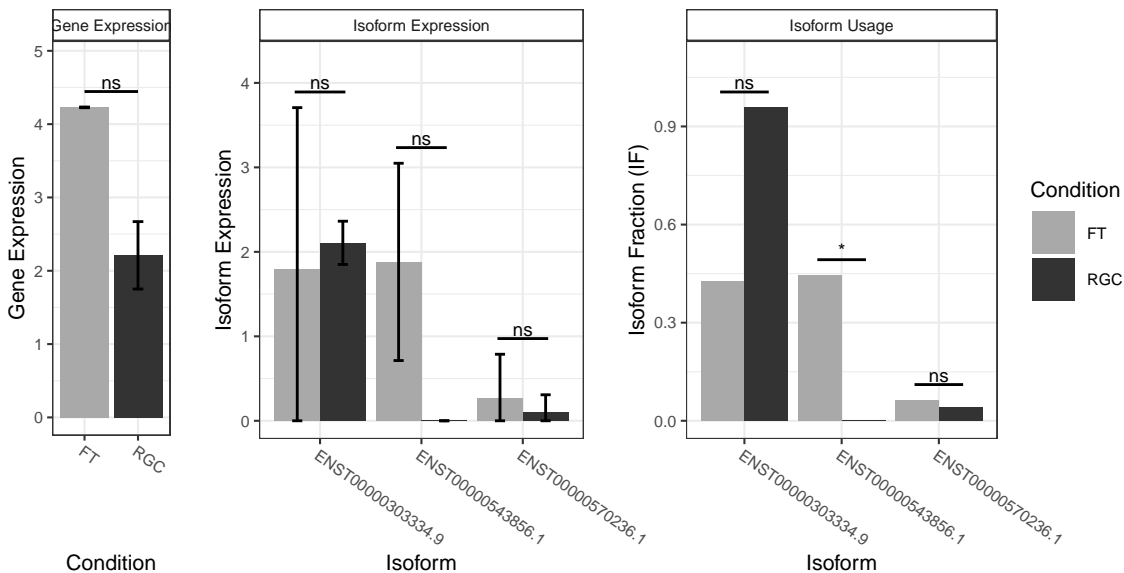
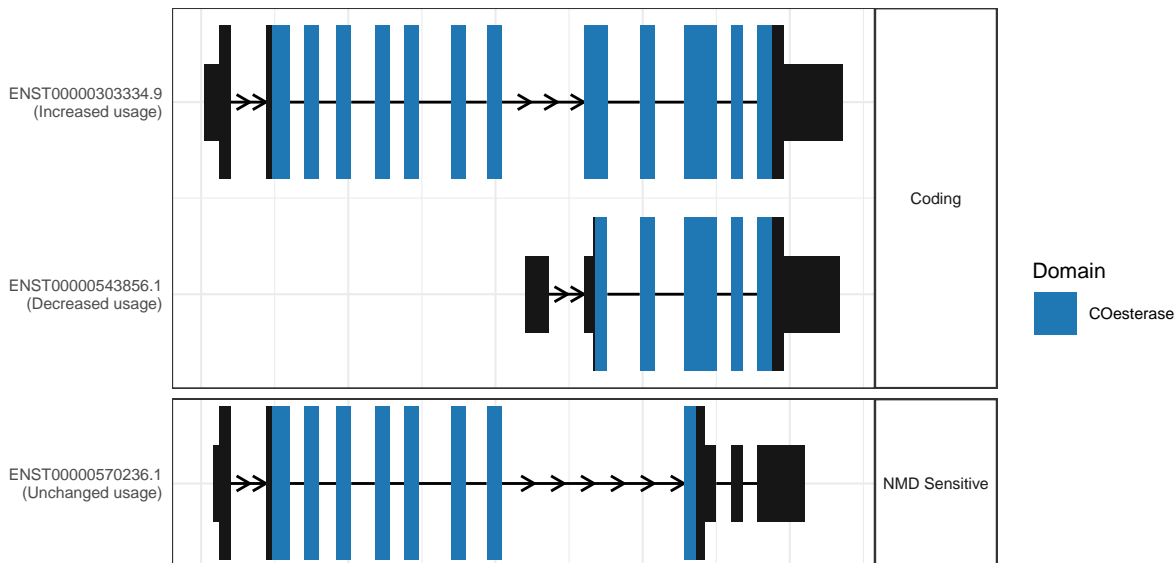


Domain



Metallophos

The isoform switch in CES3 (FT vs RGC)

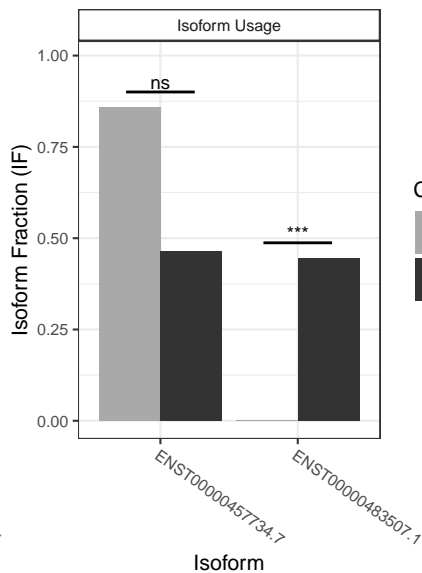
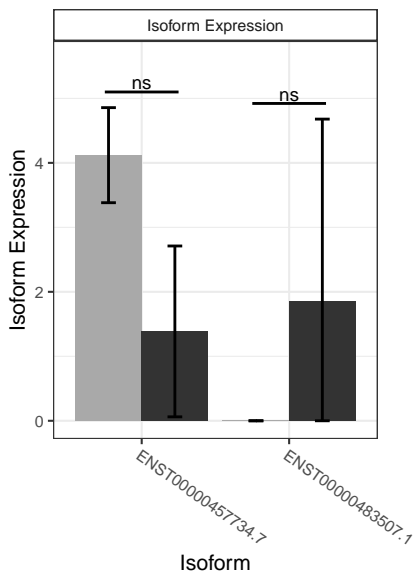
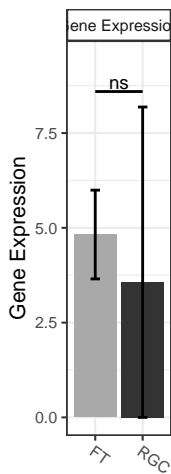
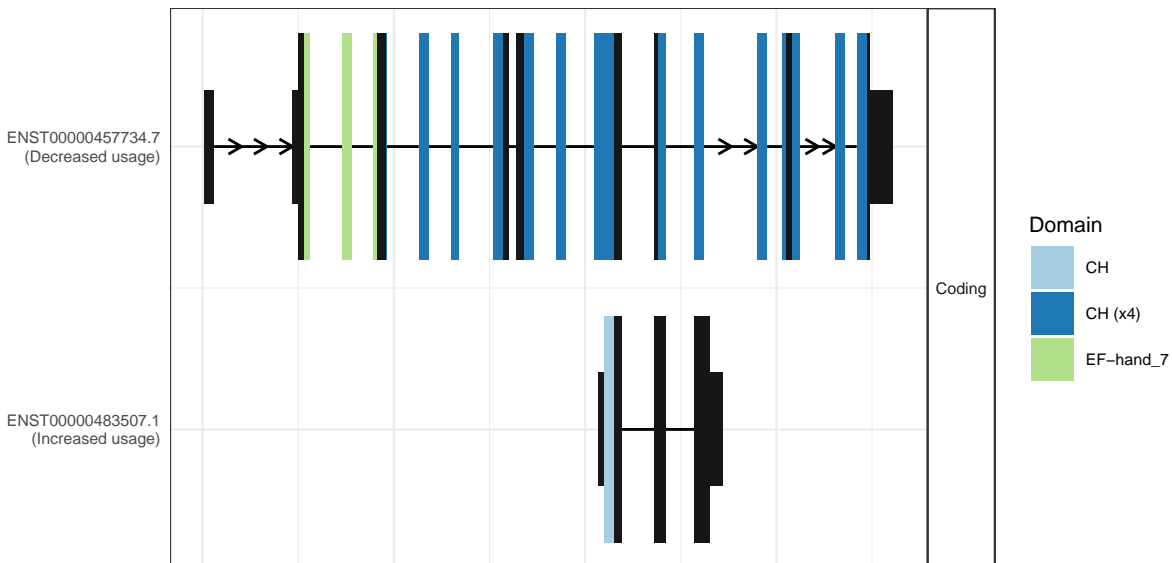


Domain



COesterase

The isoform switch in PLS1 (FT vs RGC)



Domain



CH

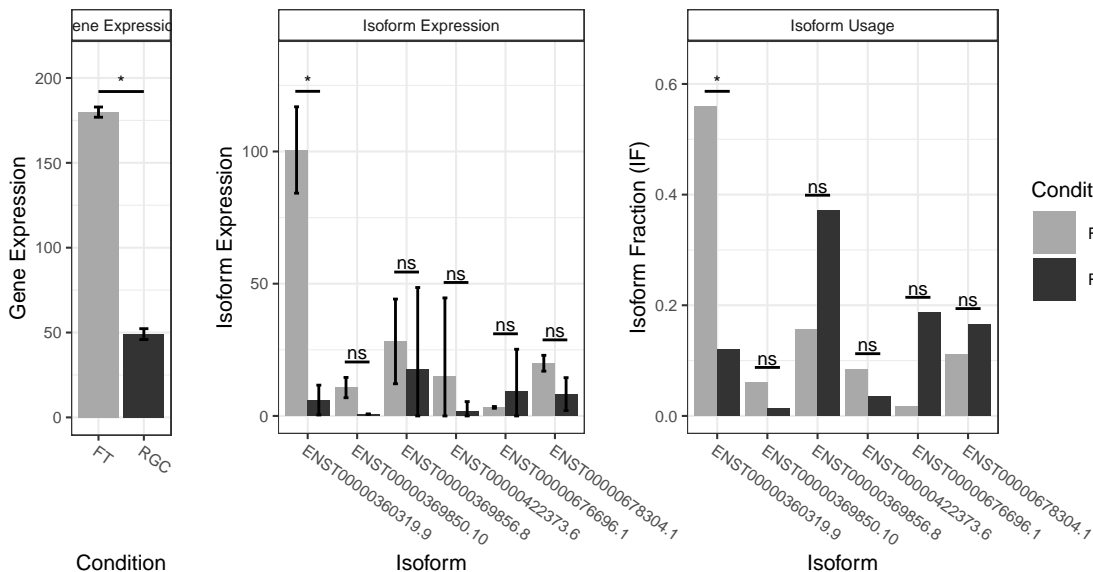
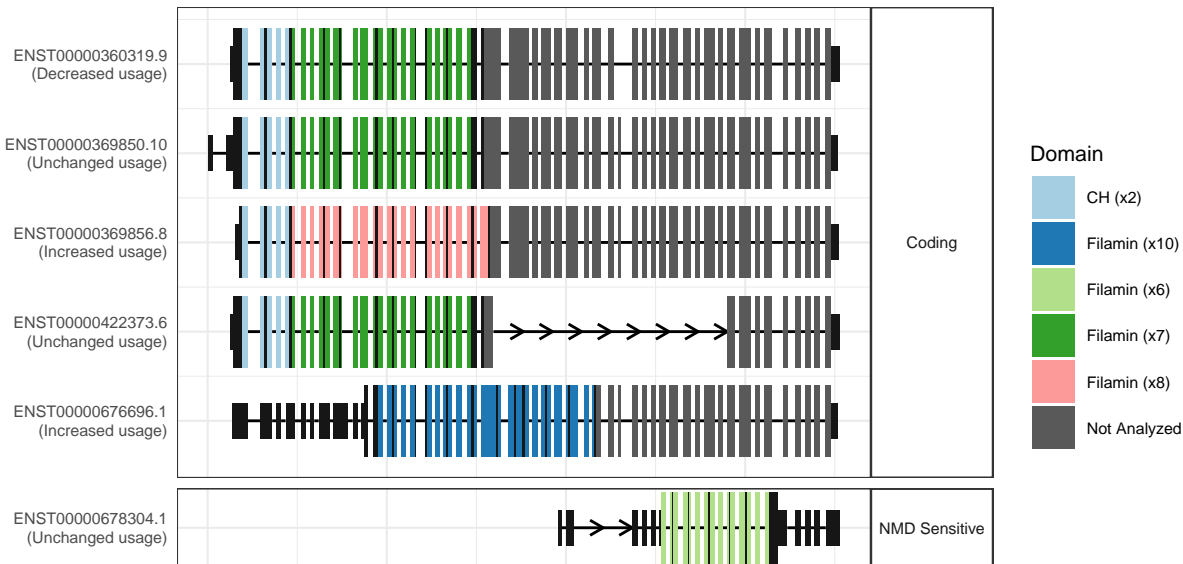


CH (x4)



EF-hand_7

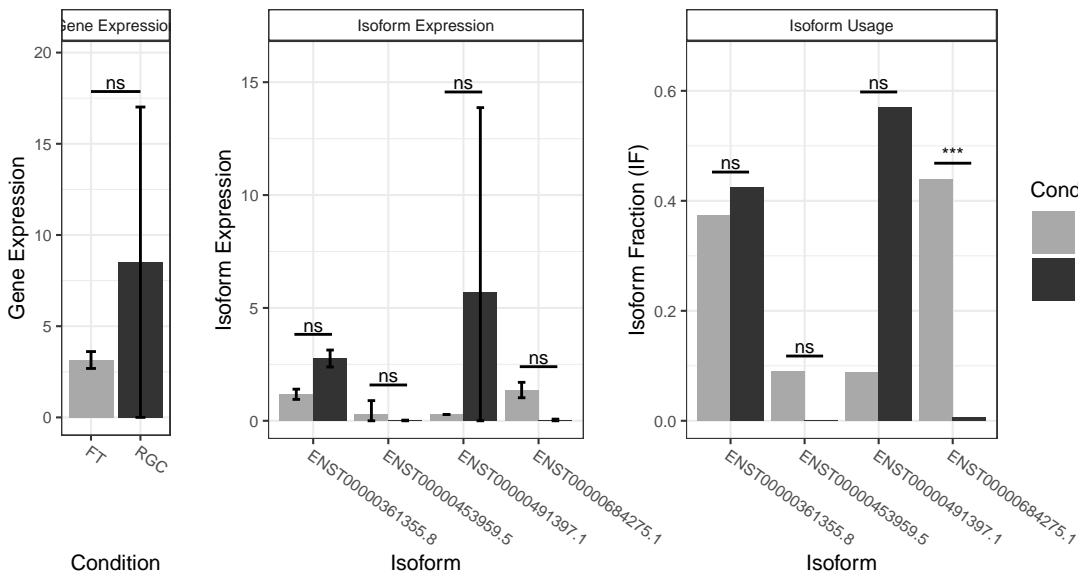
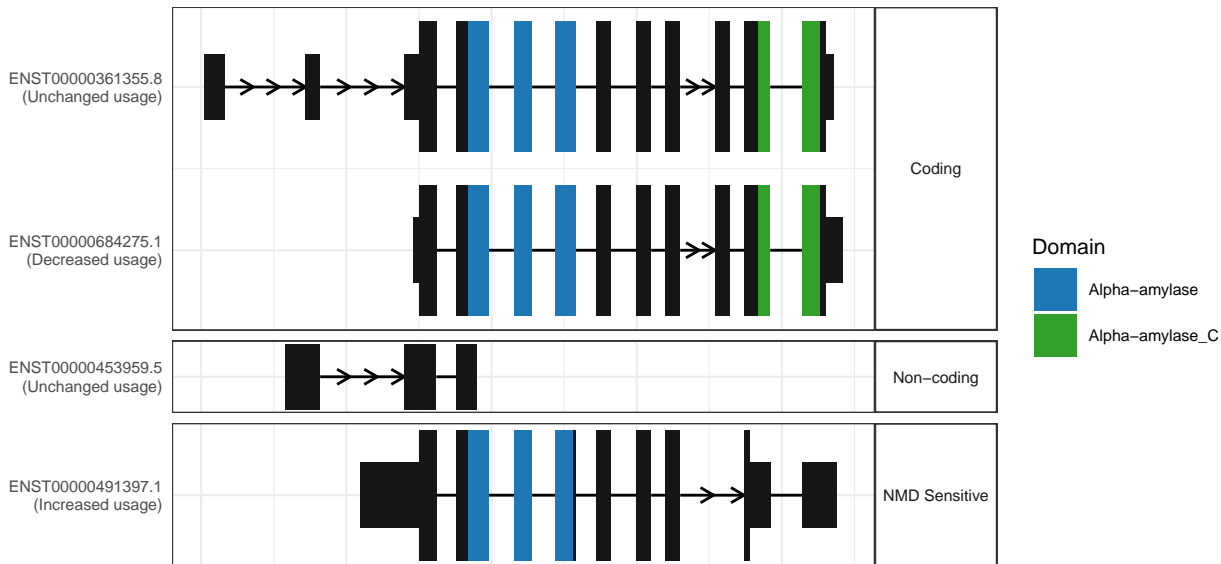
The isoform switch in FLNA (FT vs RGC)



Domain

	CH (x2)
	Filamin (x10)
	Filamin (x6)
	Filamin (x7)
	Filamin (x8)
	Not Analyzed

The isoform switch in AMY2B (FT vs RGC)



Domain

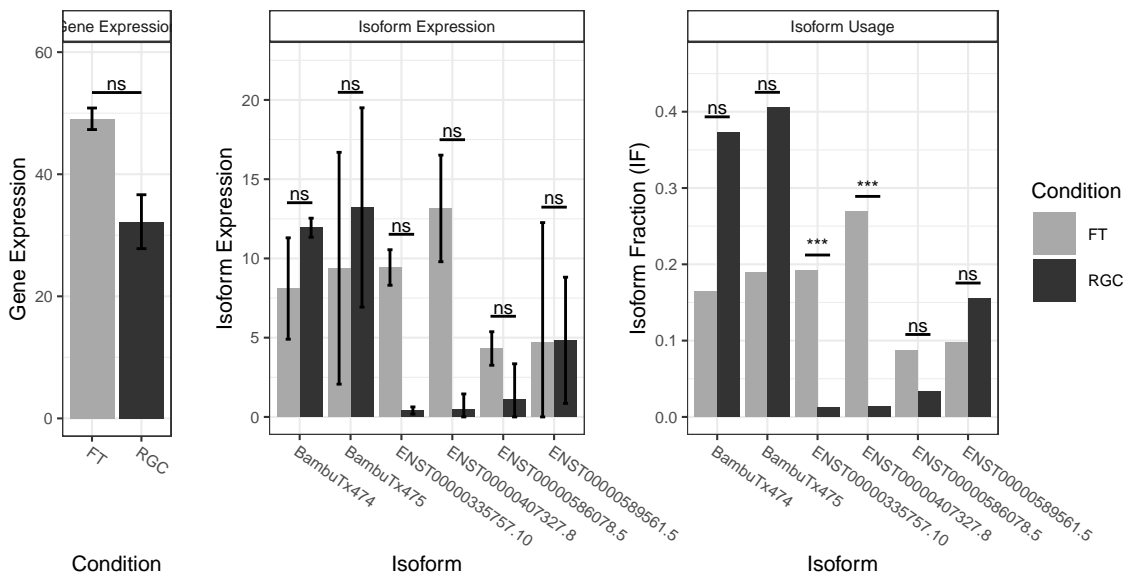
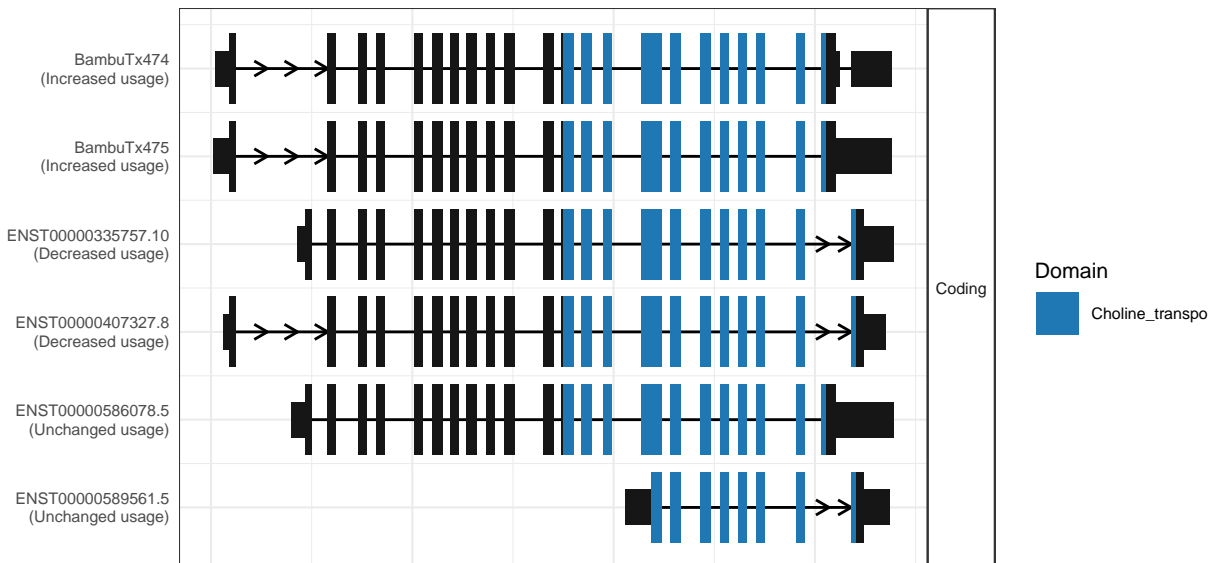


Alpha-amylase



Alpha-amylase_C

The isoform switch in SLC44A2 (FT vs RGC)

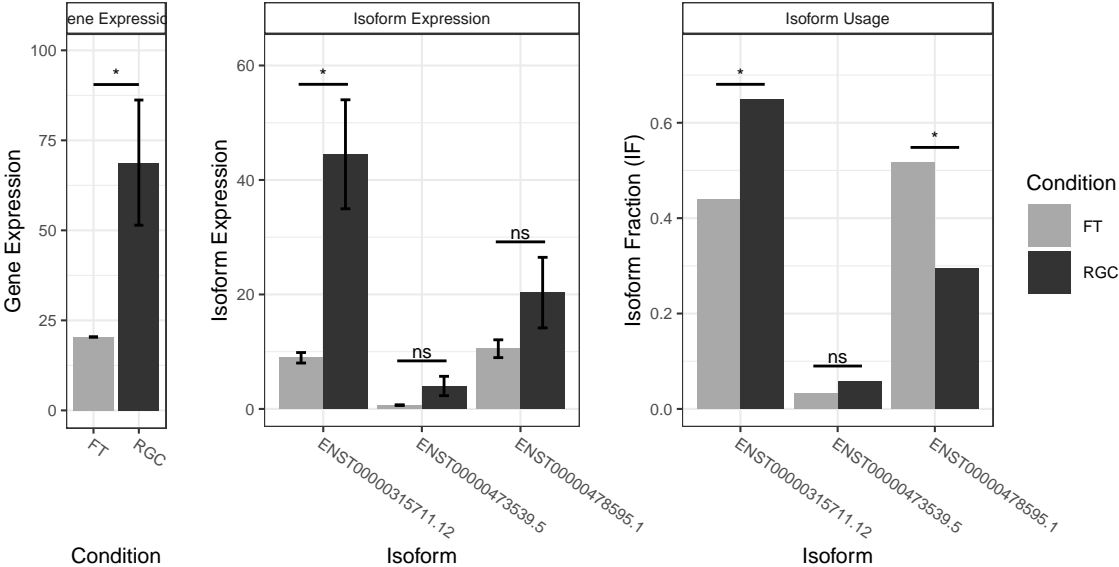
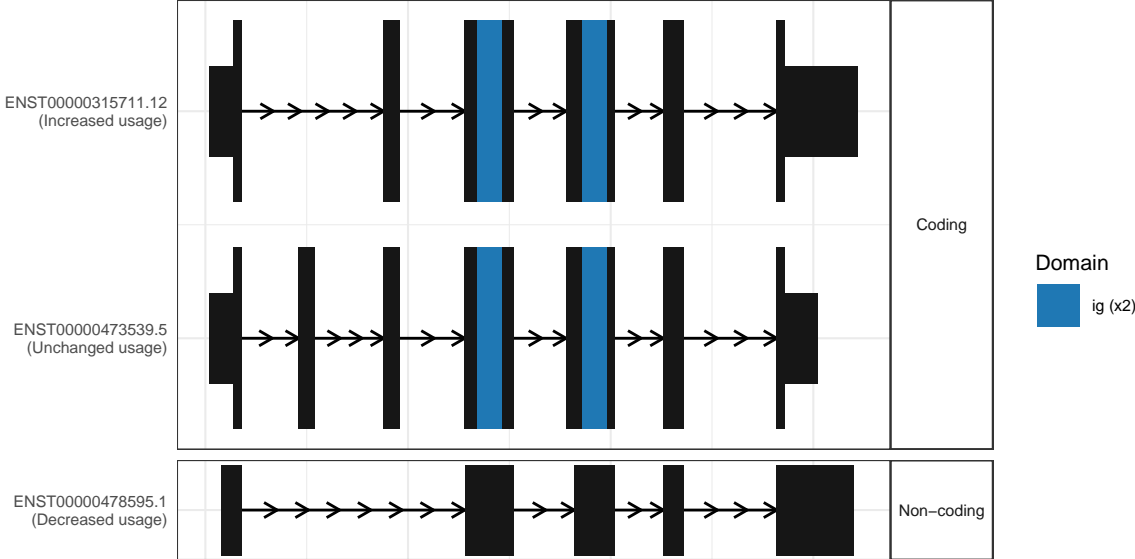


Domain



Choline_transpo

The isoform switch in CD200 (FT vs RGC)

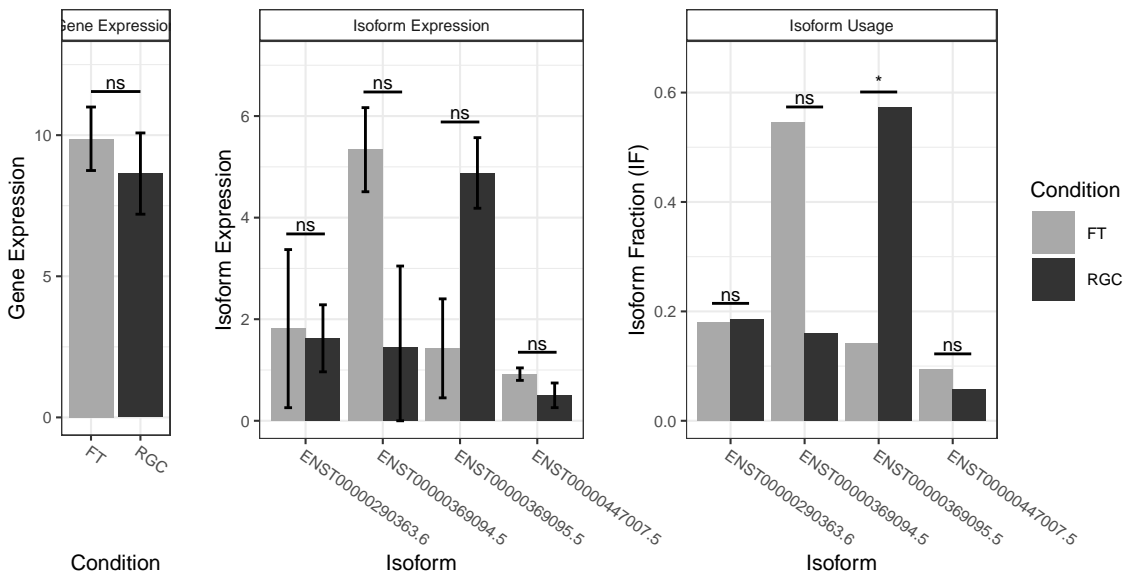
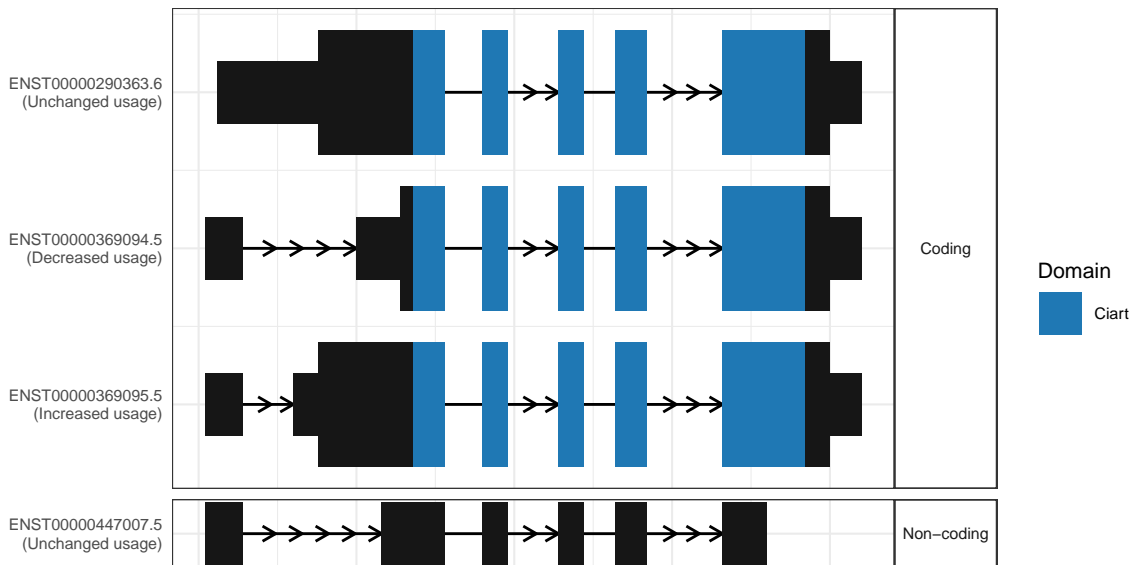


Domain



ig (x2)

The isoform switch in CIART (FT vs RGC)

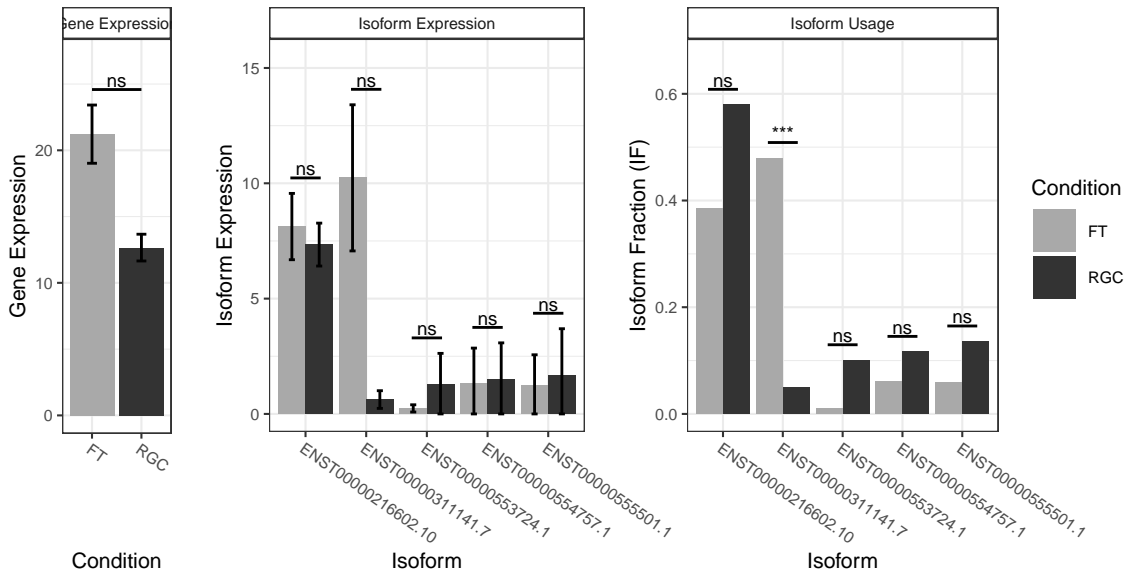
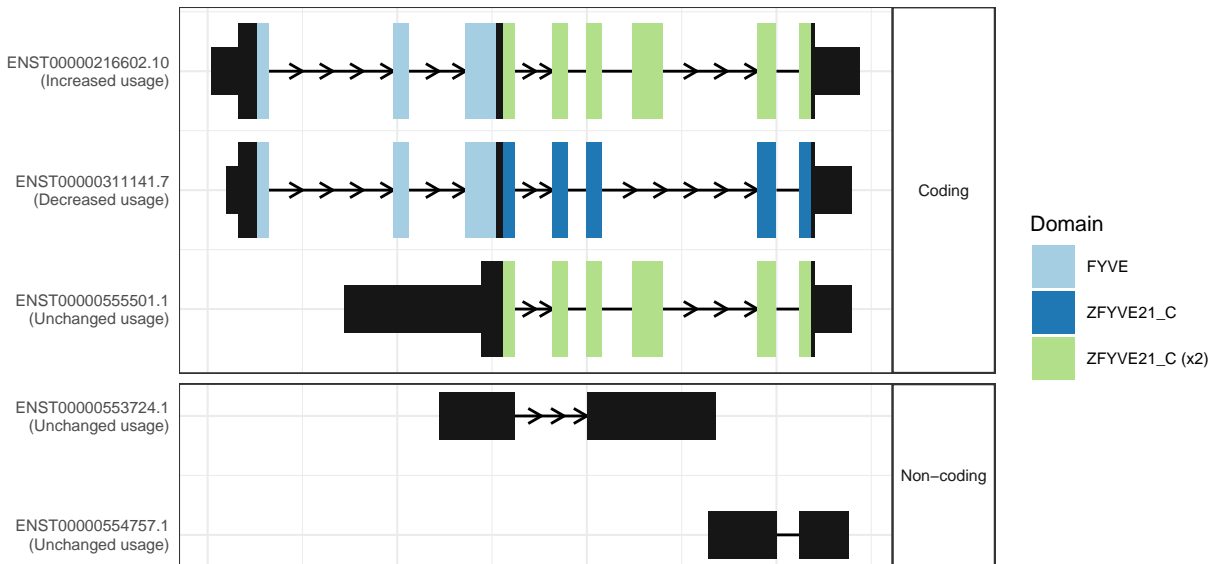


Domain



Ciart

The isoform switch in ZFYVE21 (FT vs RGC)



Domain



FYVE

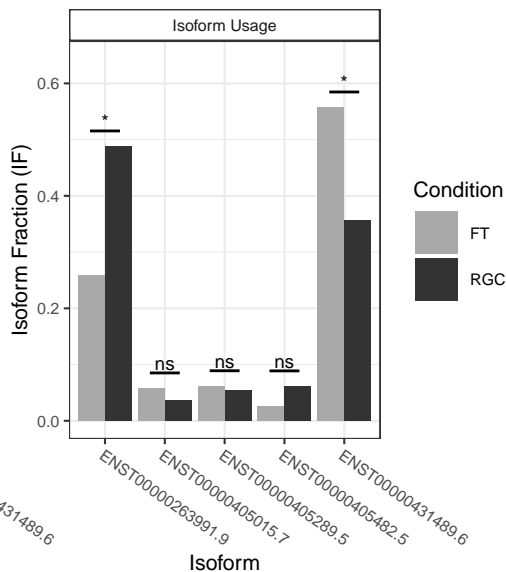
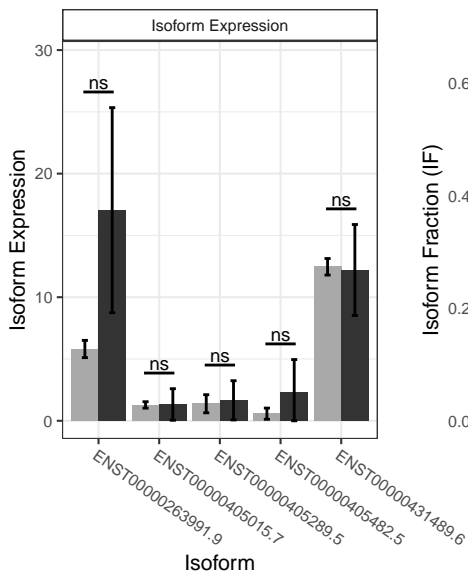
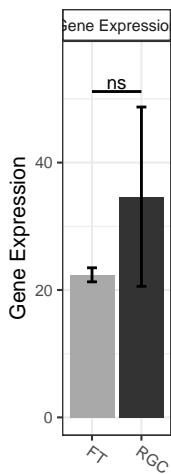
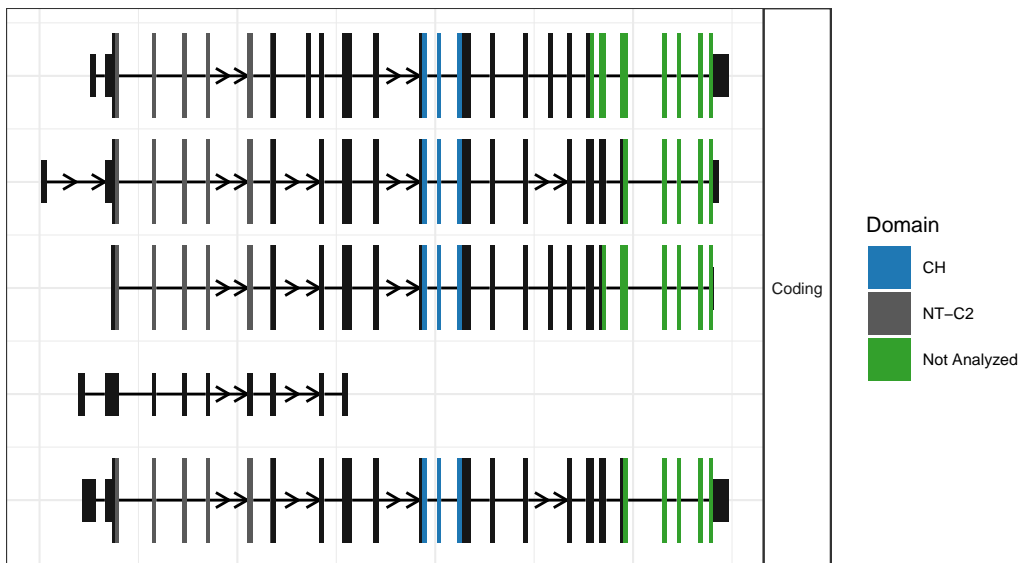


ZFYVE21_C



ZFYVE21_C (x2)

The isoform switch in EHBP1 (FT vs RGC)



Domain



CH

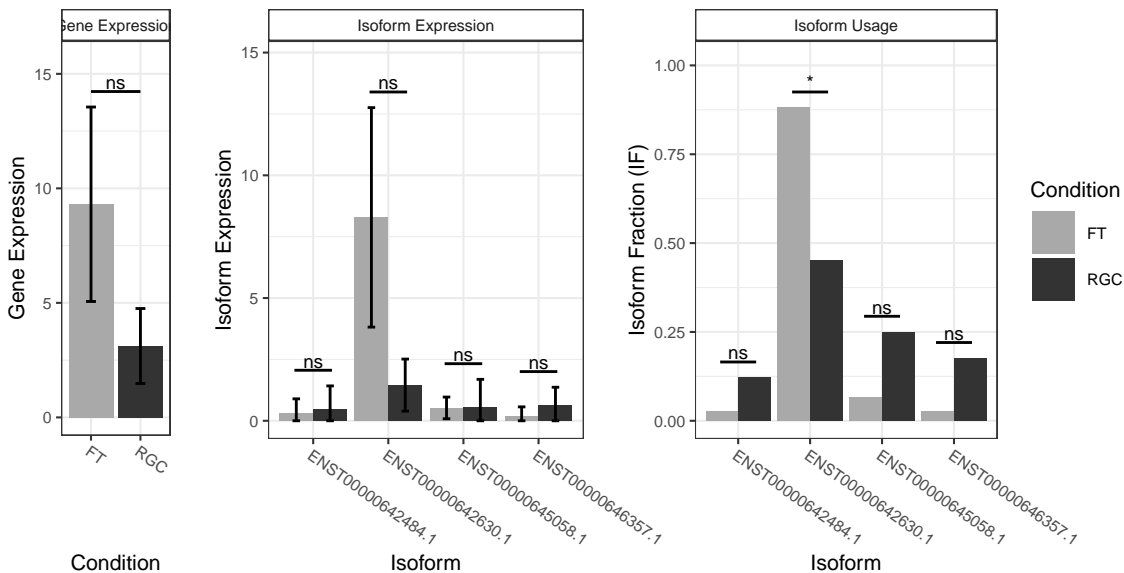
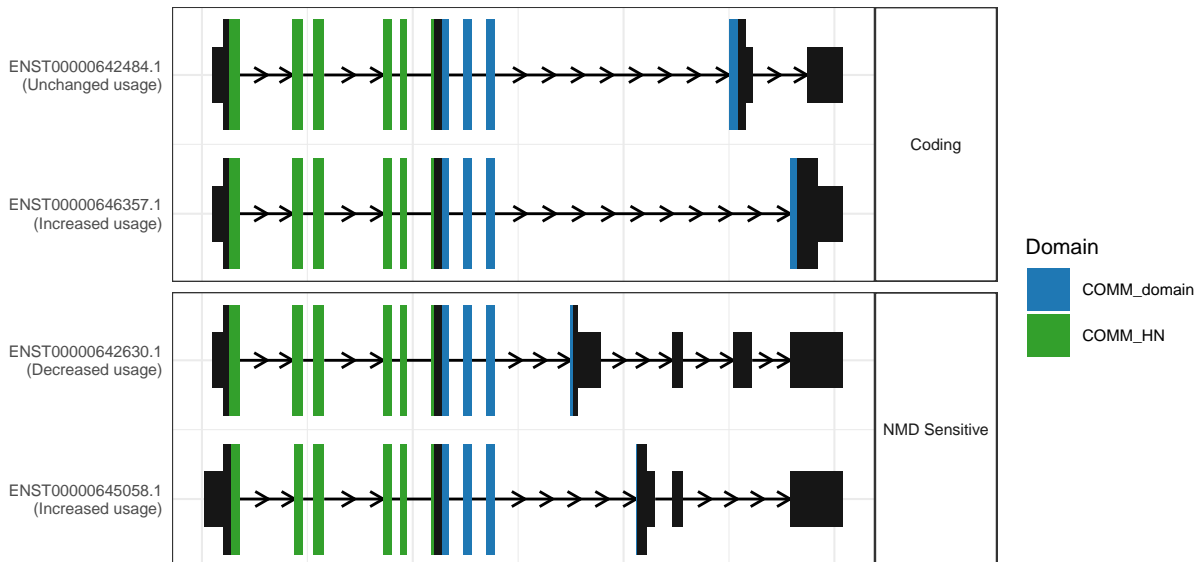


NT-C2



Not Analyzed

The isoform switch in (FT vs RGC)



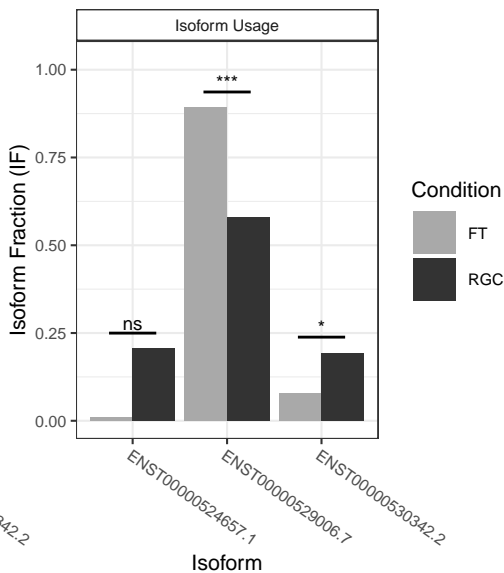
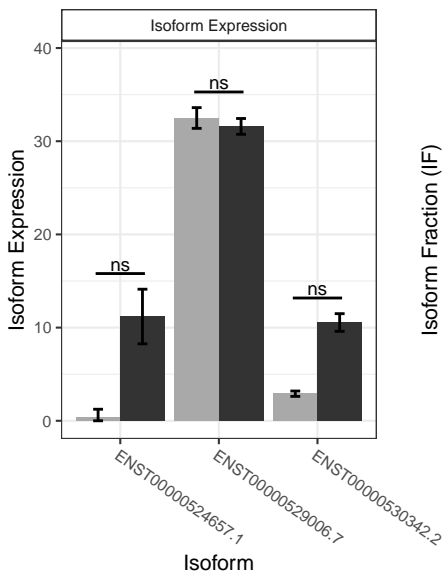
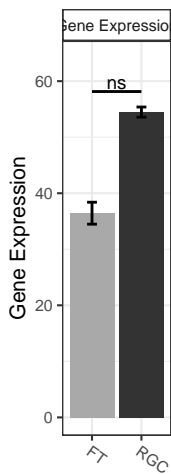
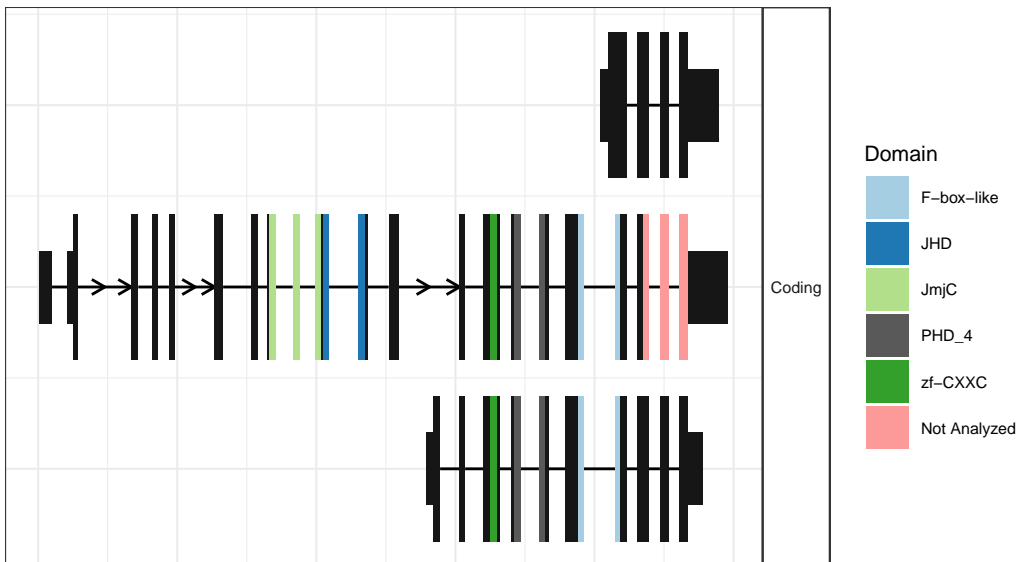
Domain



COMM_domain

COMM_HN

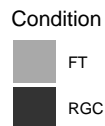
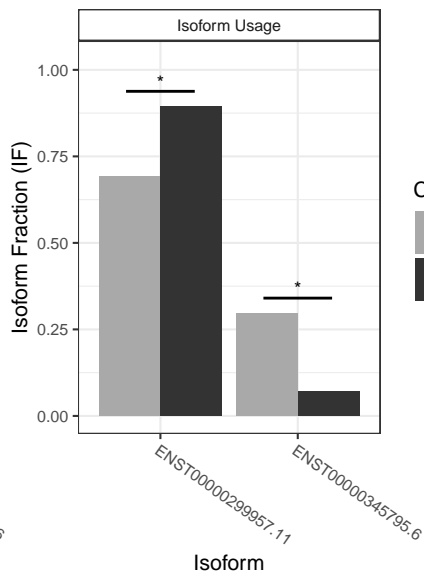
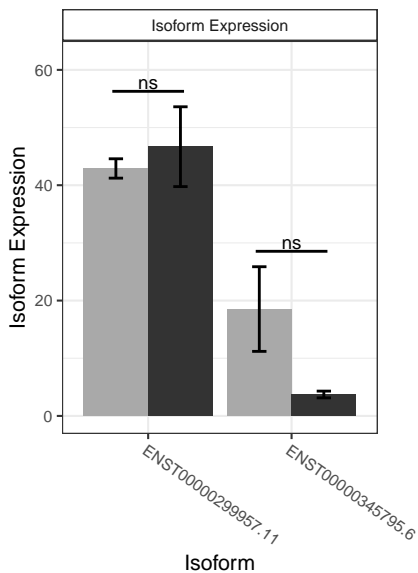
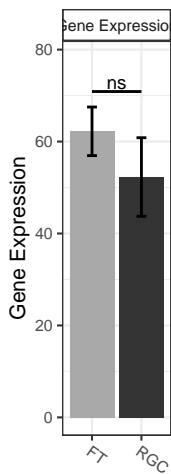
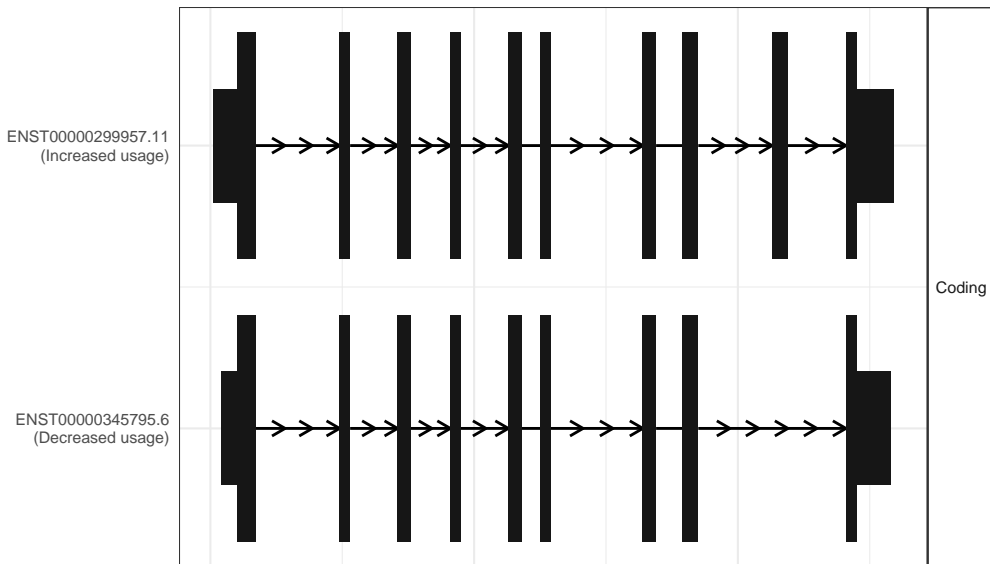
The isoform switch in KDM2A (FT vs RGC)



Domain

	F-box-like
	JHD
	JmjC
	PHD_4
	zf-CXXC
	Not Analyzed

The isoform switch in GOLM2 (FT vs RGC)

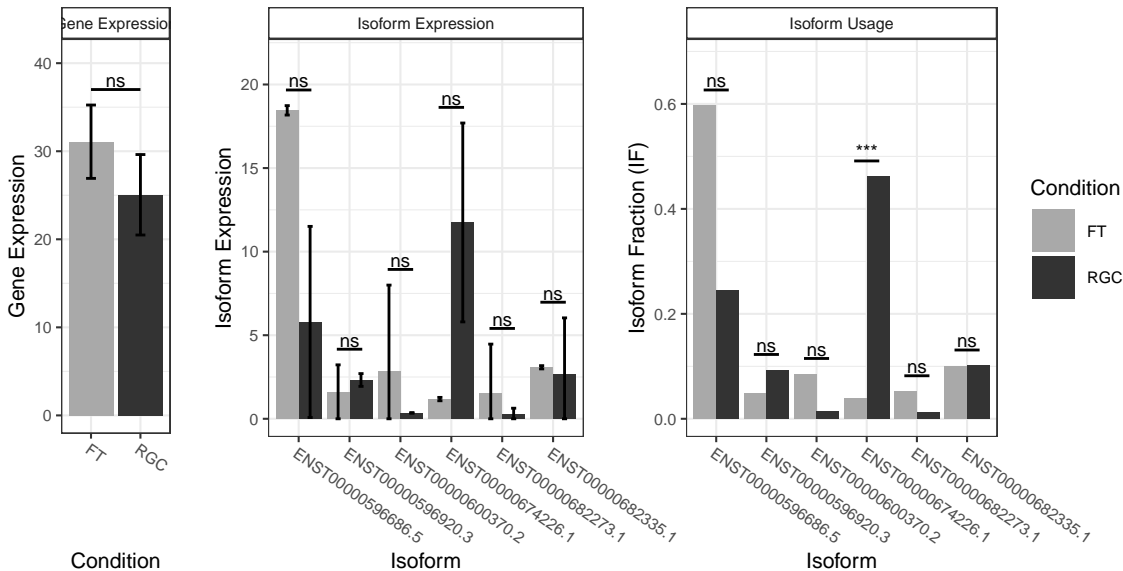
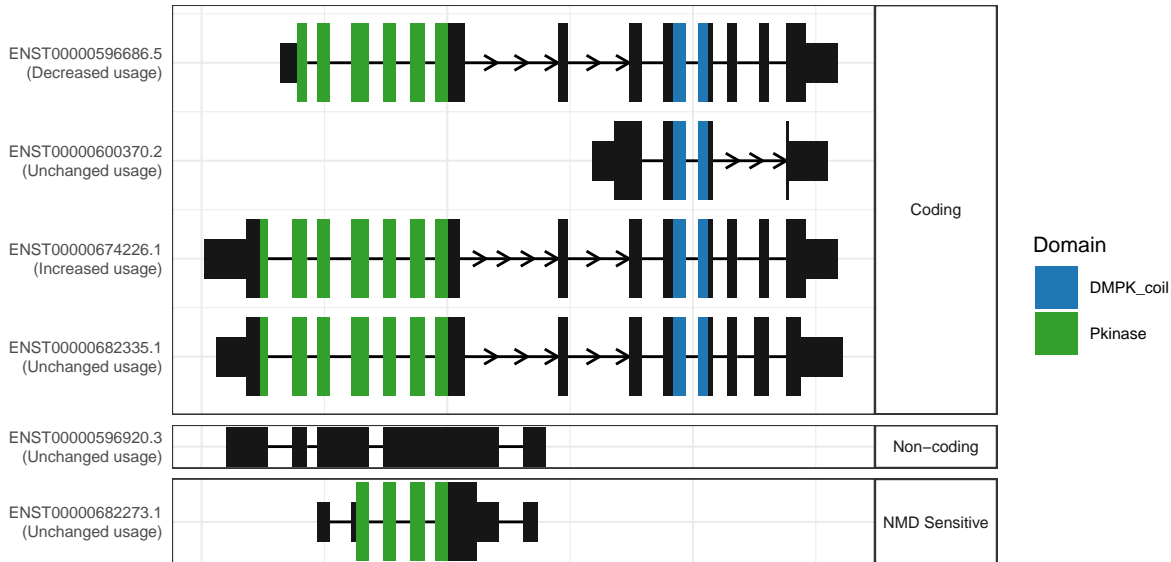


Domain



RAMP

The isoform switch in DMPK (FT vs RGC)



Domain

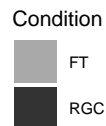
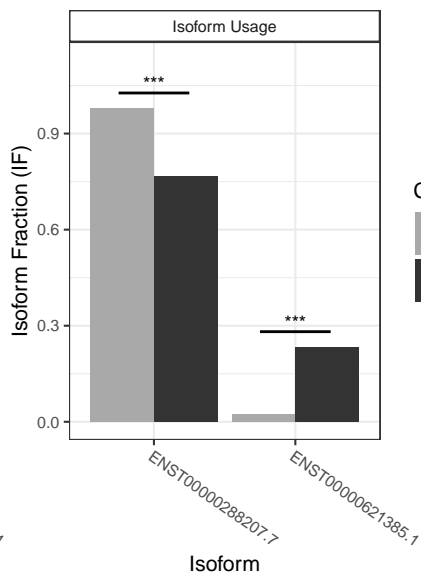
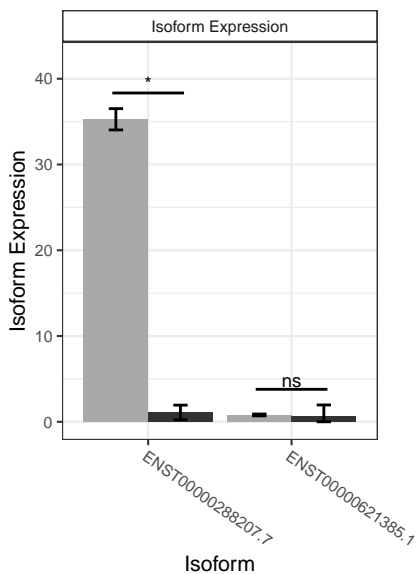
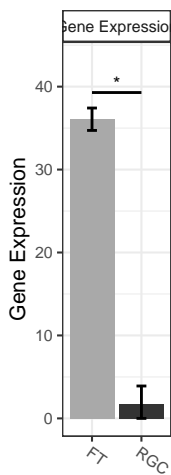
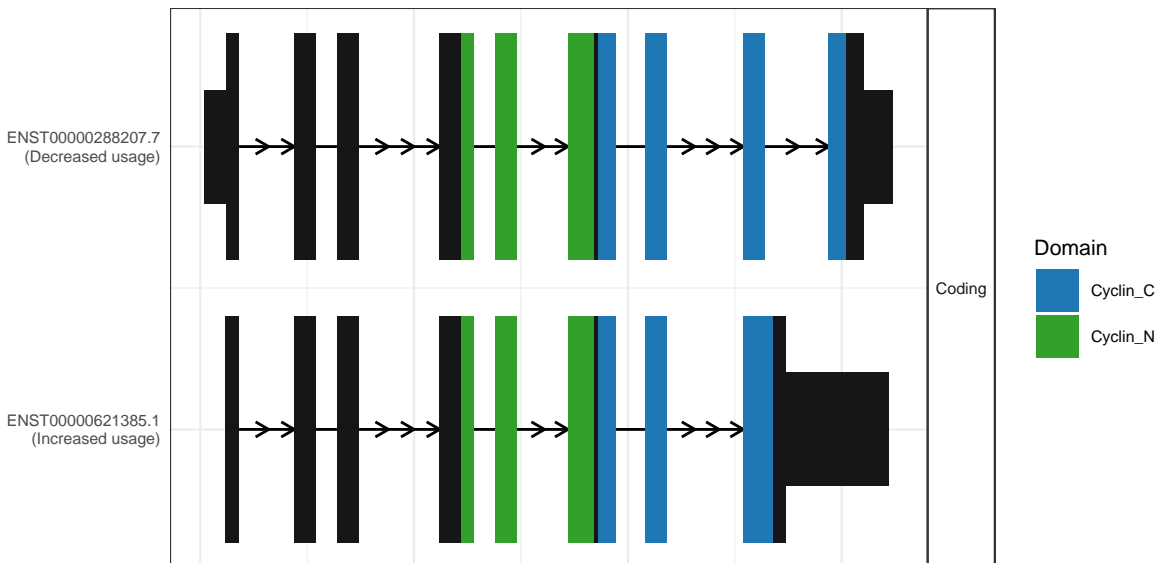


DMPK_coil



Pkinase

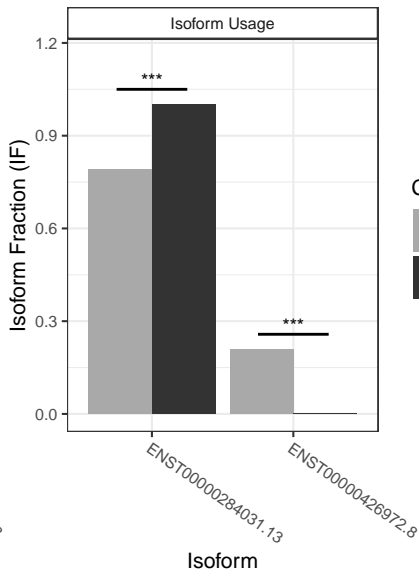
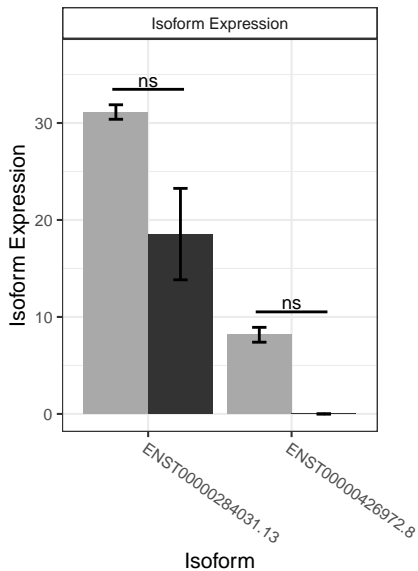
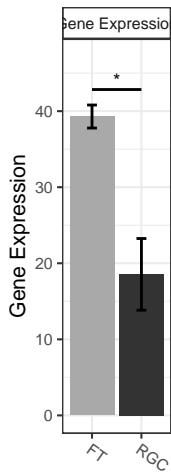
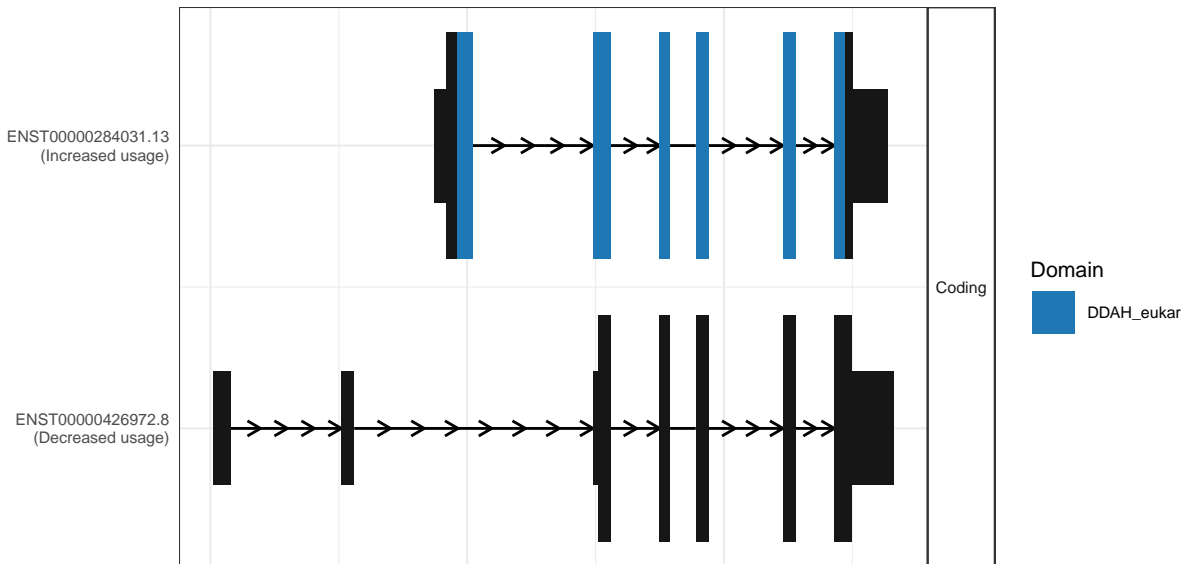
The isoform switch in CCNB2 (FT vs RGC)



Domain



The isoform switch in DDAH1 (FT vs RGC)

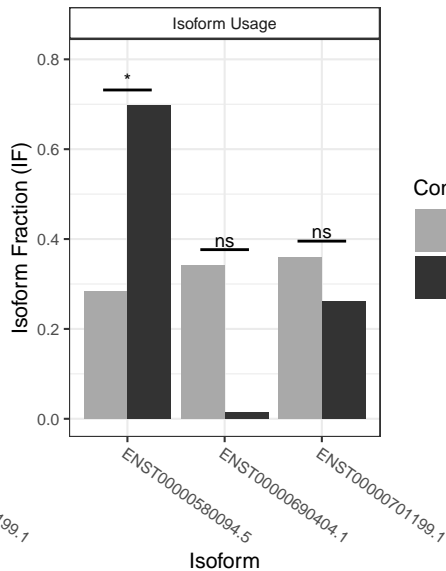
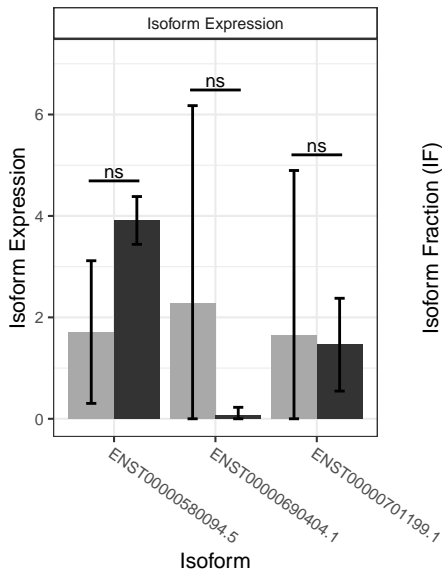
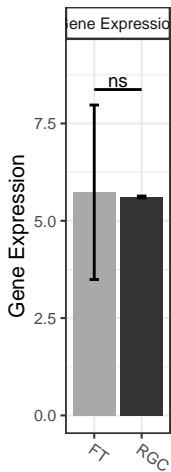
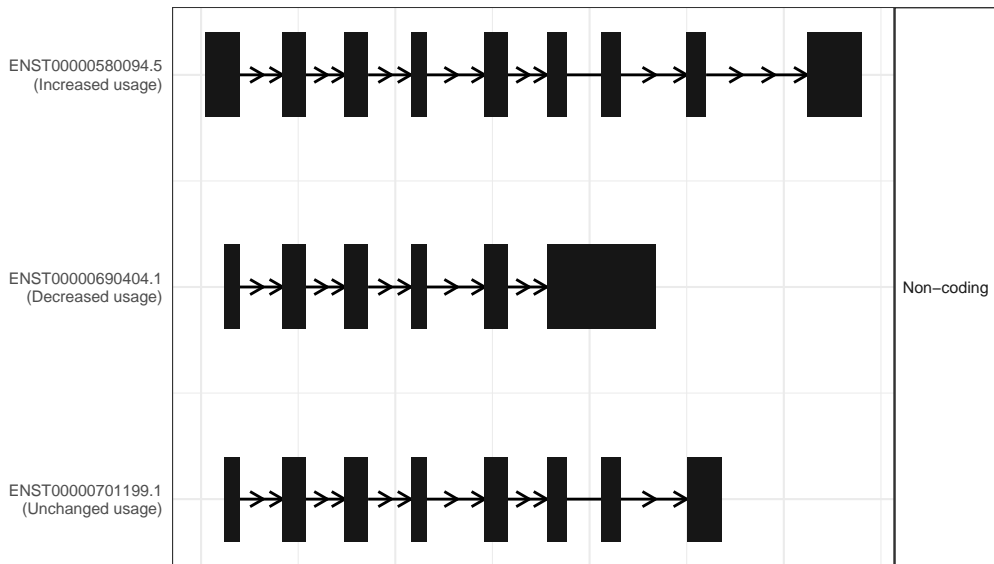


Domain



DDAH_eukar

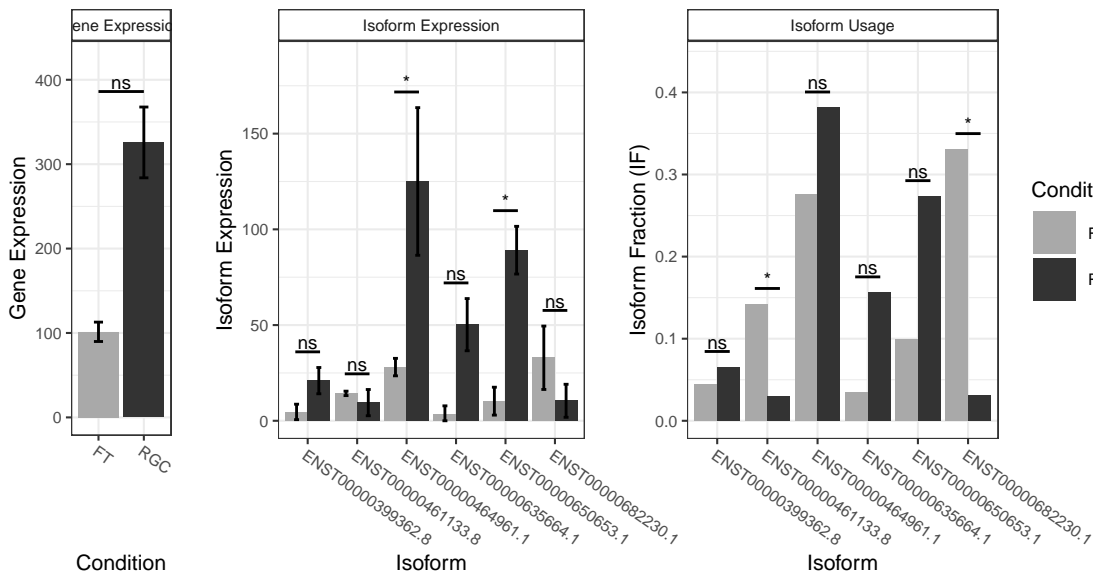
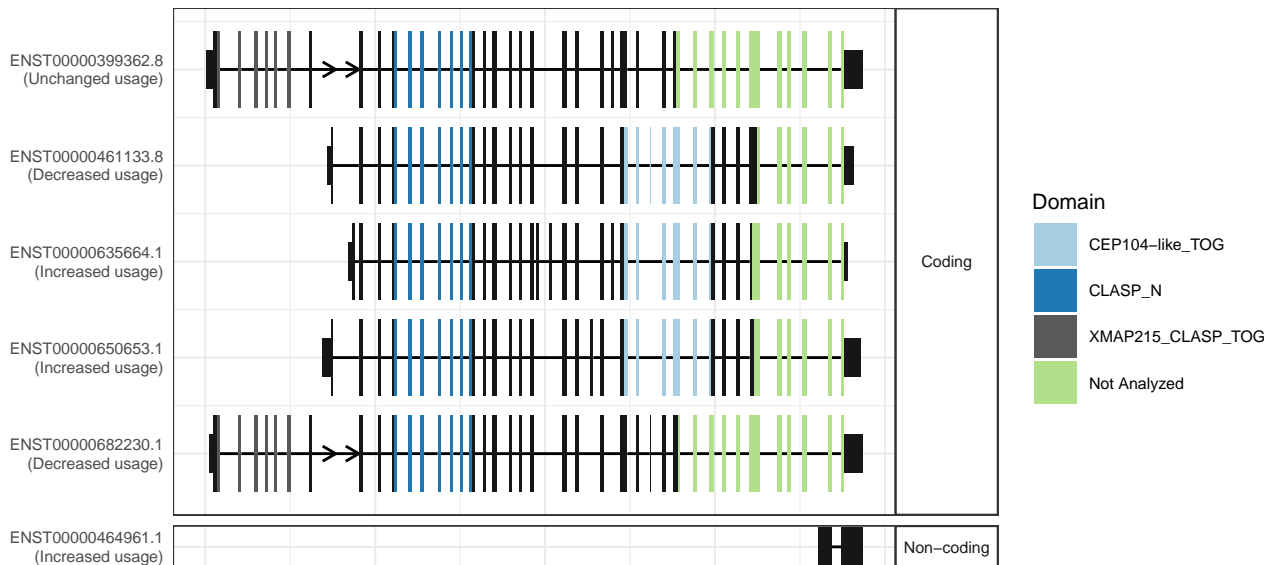
The isoform switch in (FT vs RGC)



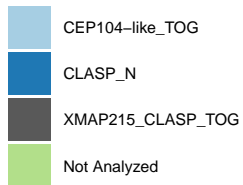
Condition



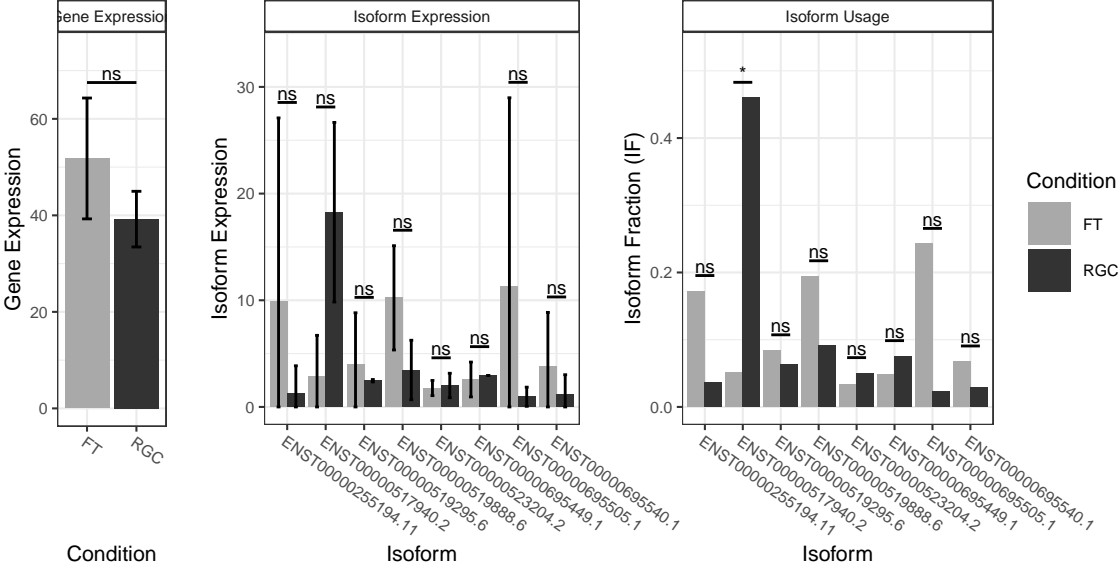
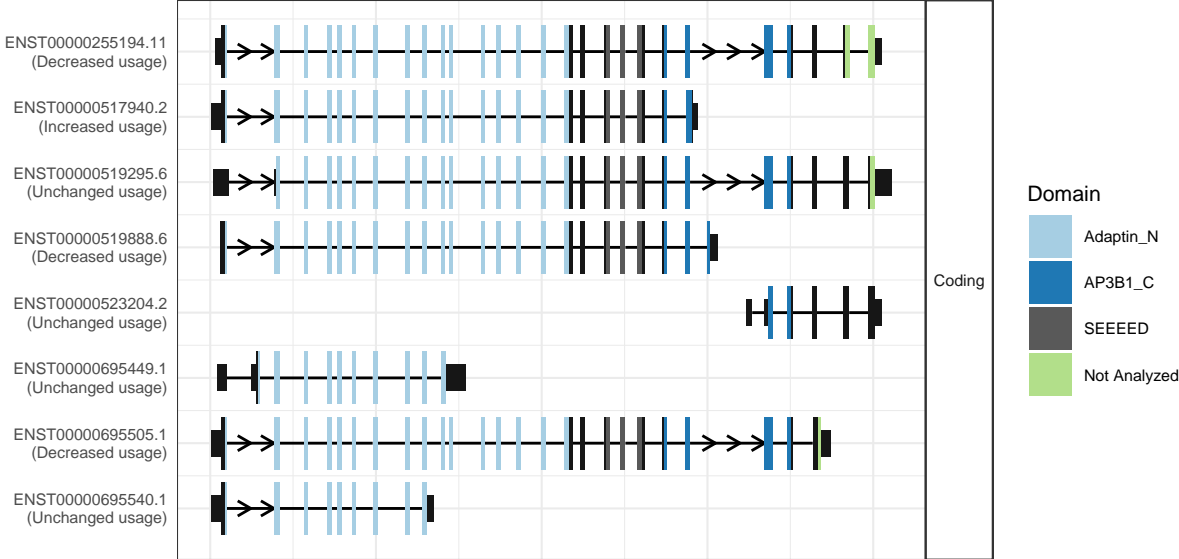
The isoform switch in CLASP2 (FT vs RGC)







Domain



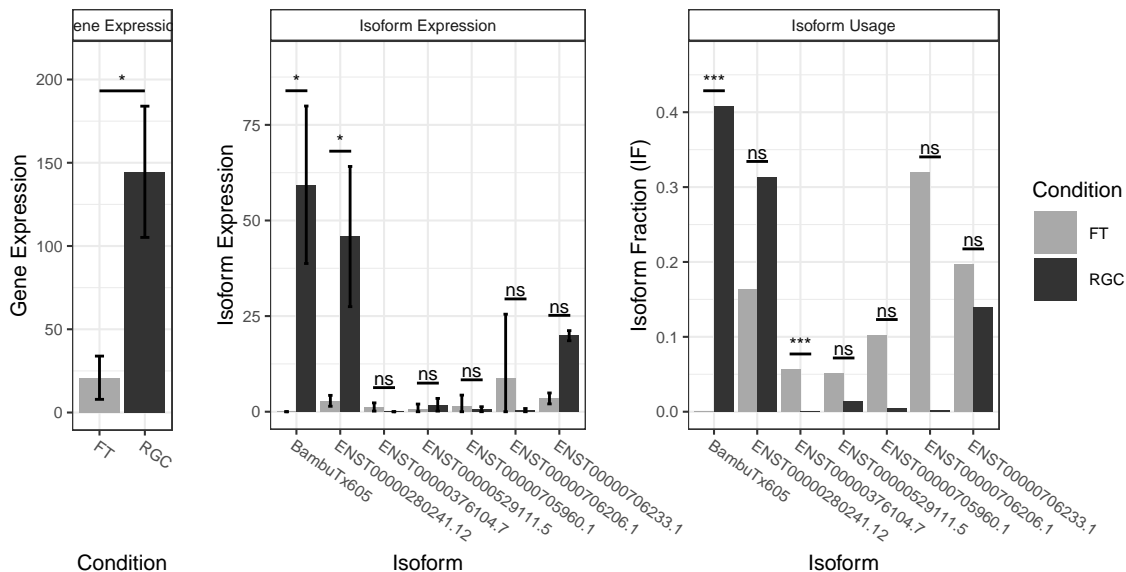
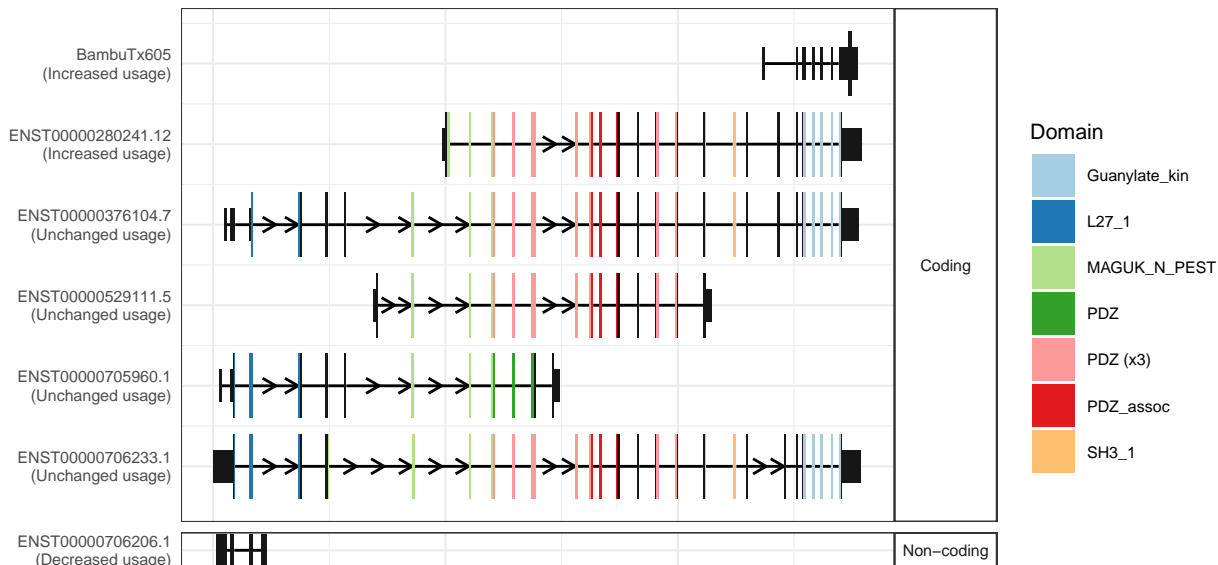
The isoform switch in AP3B1 (FT vs RGC)










Domain

	Adaptin_N
	AP3B1_C
	SEEEED
	Not Analyzed

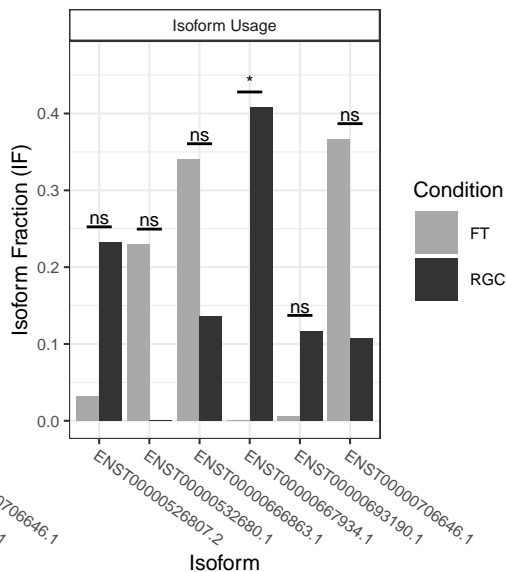
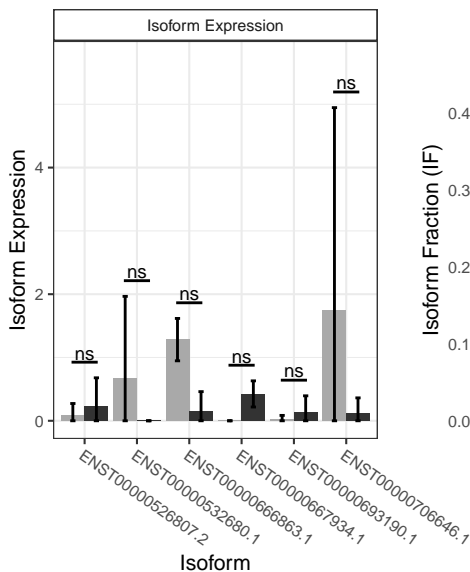
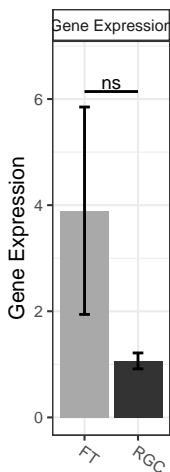
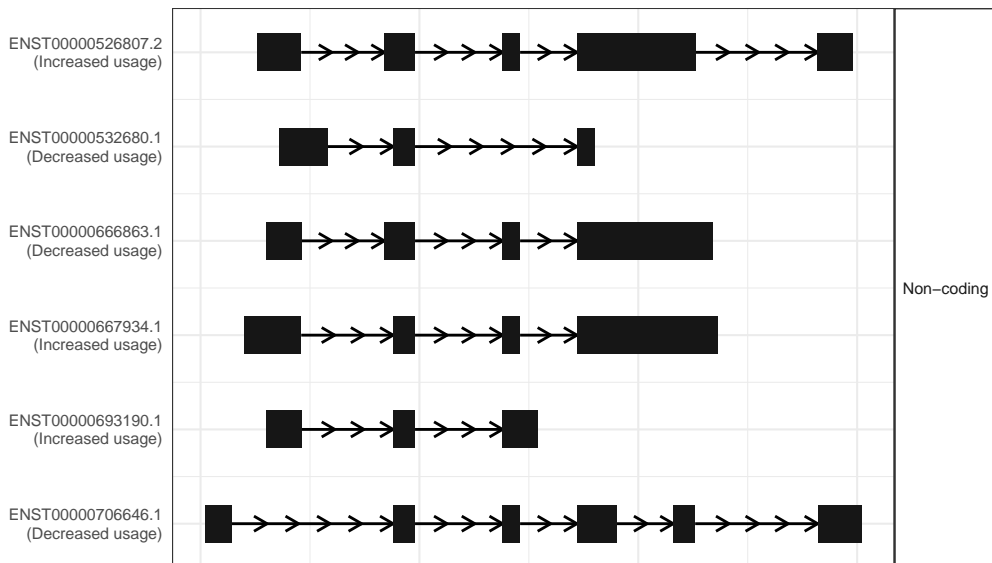
The isoform switch in DLG2 (FT vs RGC)



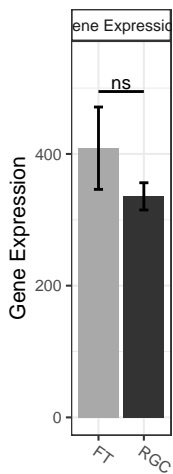
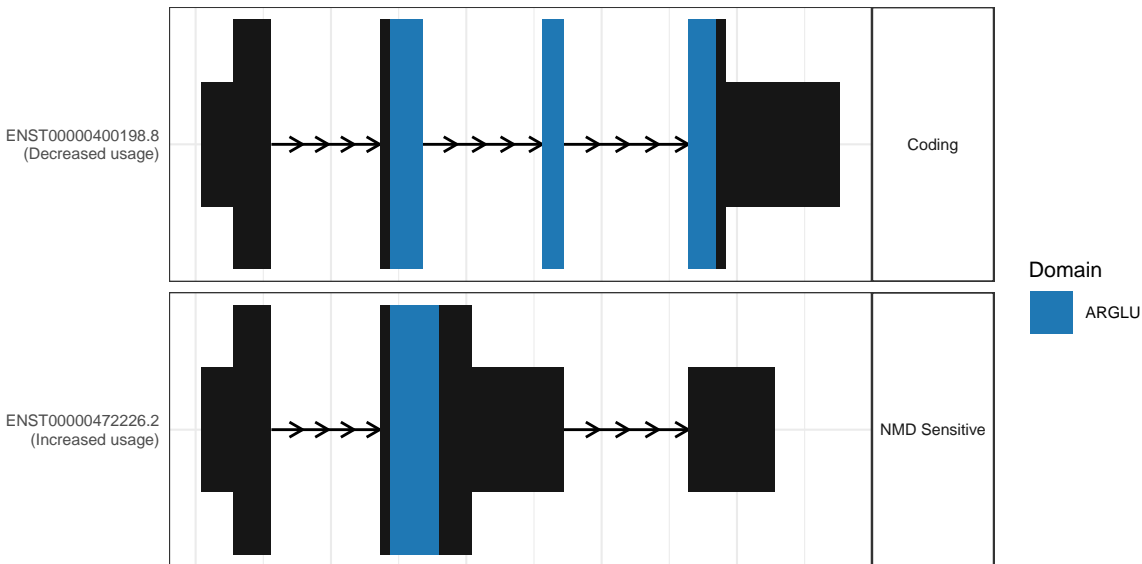
Domain

	Guanylate_kin
	L27_1
	MAGUK_N_PEST
	PDZ
	PDZ (x3)
	PDZ_assoc
	SH3_1

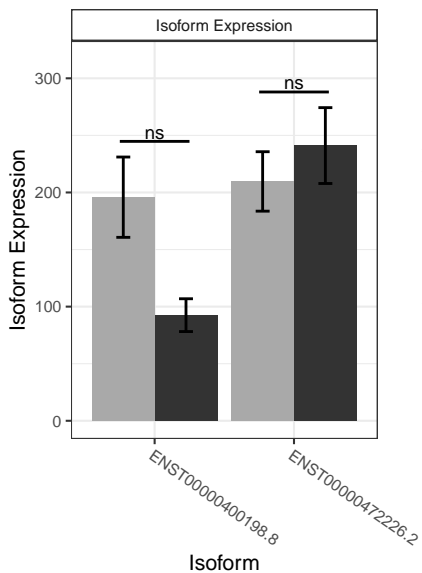
The isoform switch in SMIM14-DT (FT vs RGC)



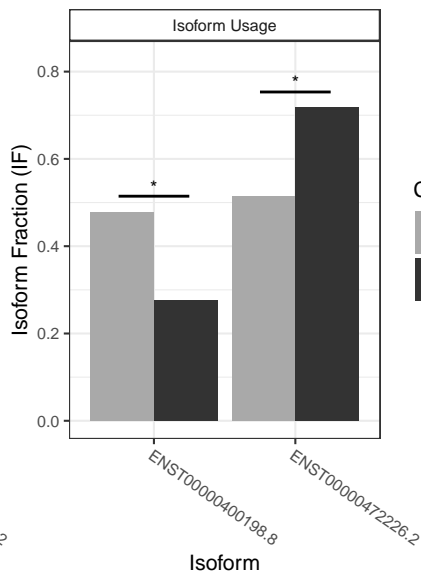
The isoform switch in ARGLU1 (FT vs RGC)



Condition



Isoform



Isoform

Condition

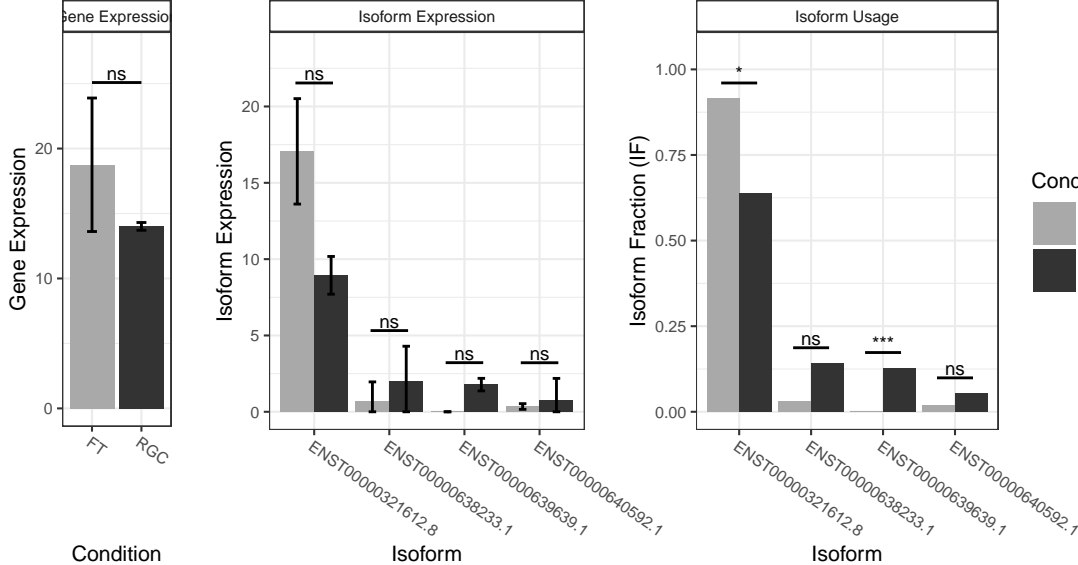
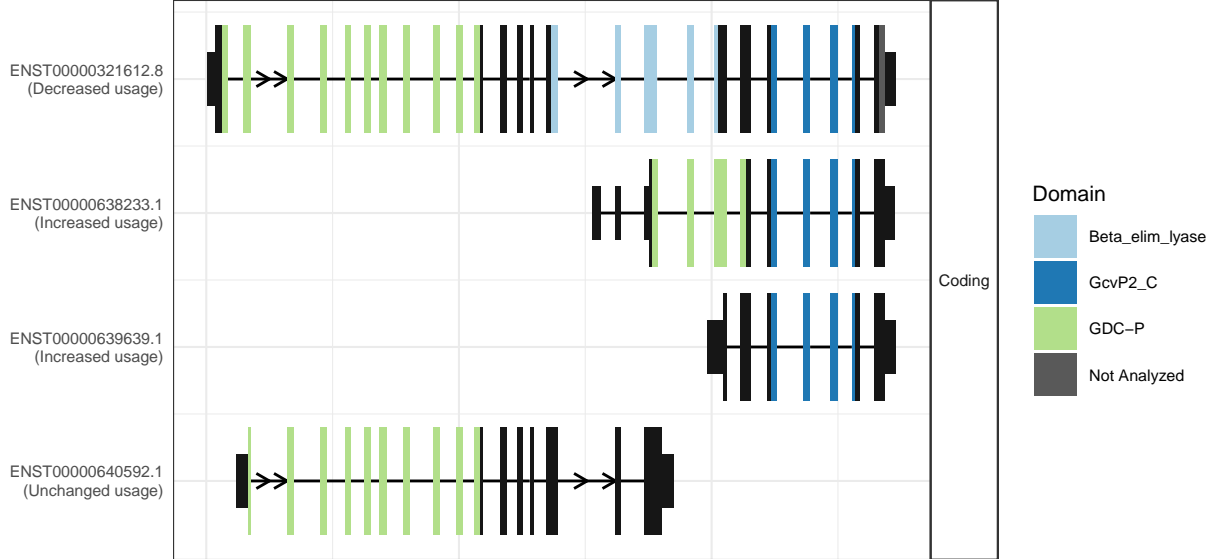


Domain

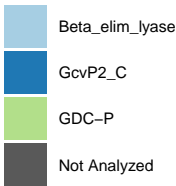


ARGLU

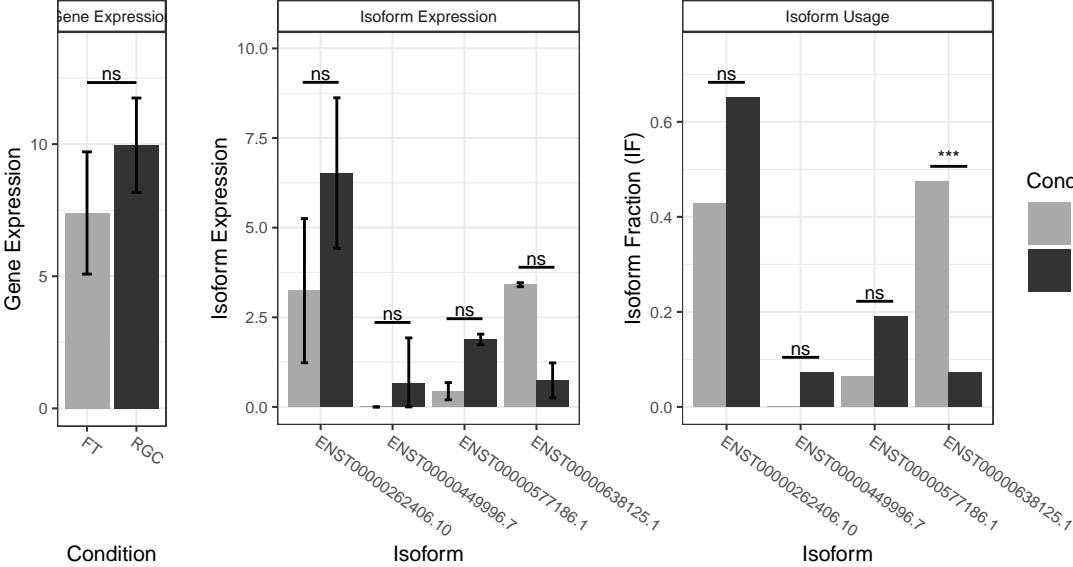
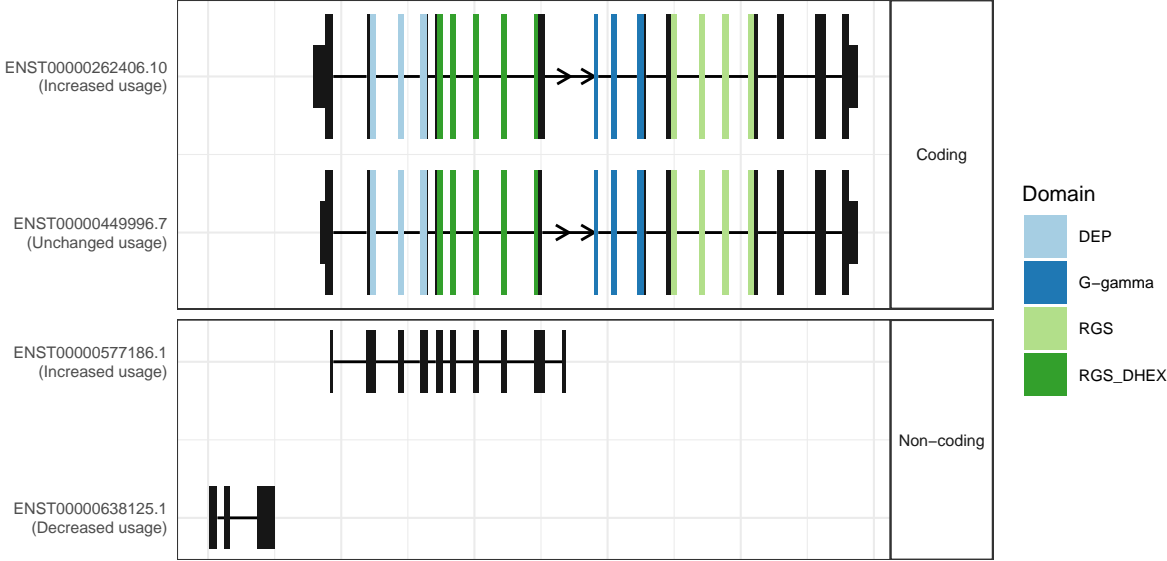
The isoform switch in GLDC (FT vs RGC)



Domain



The isoform switch in RGS9 (FT vs RGC)



Domain



DEP



G-gamma

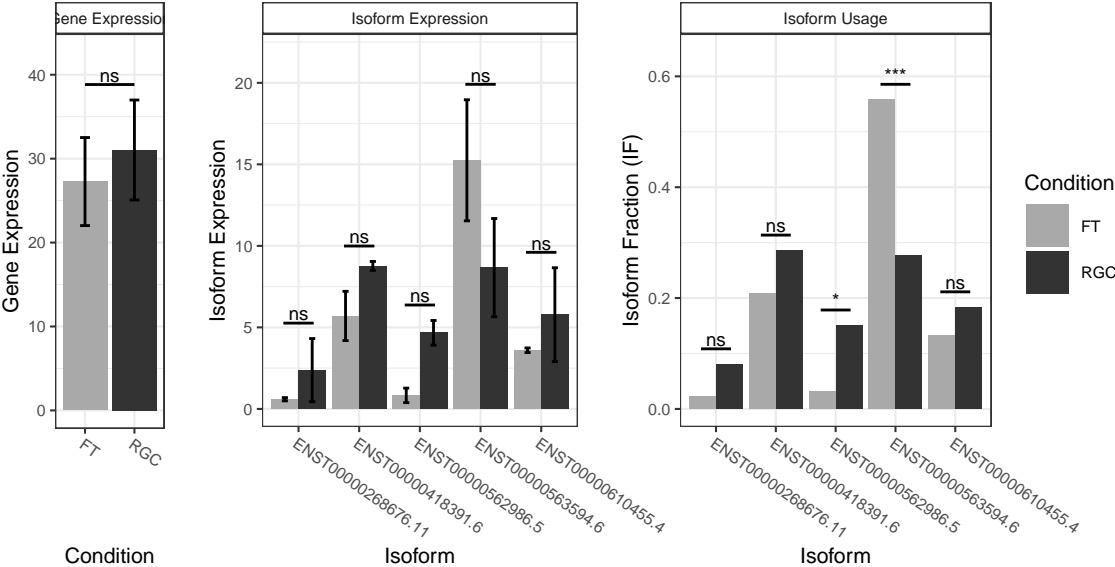
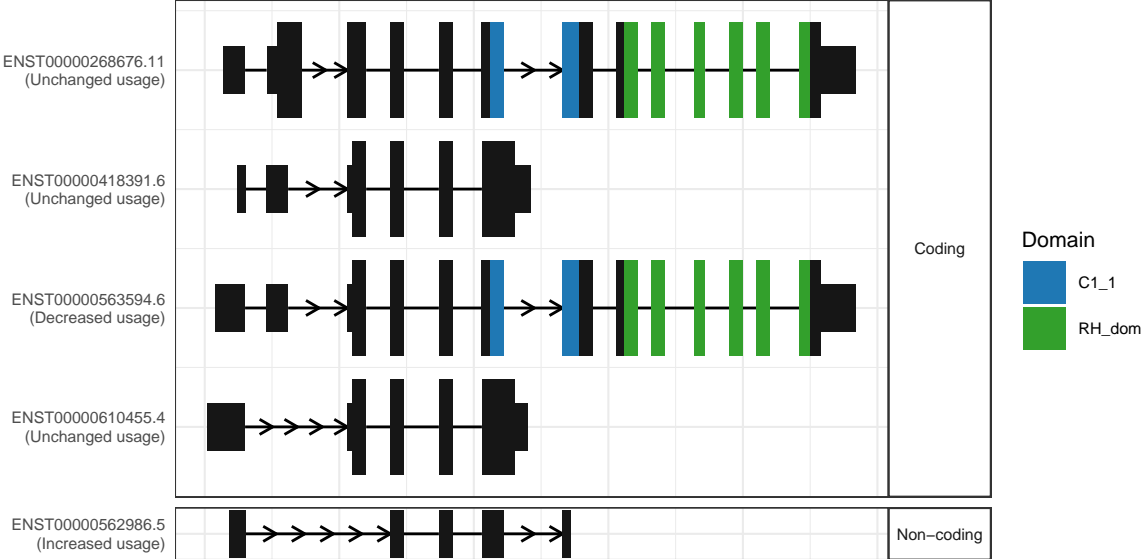


RGS



RGS_DHEX

The isoform switch in DEF8 (FT vs RGC)



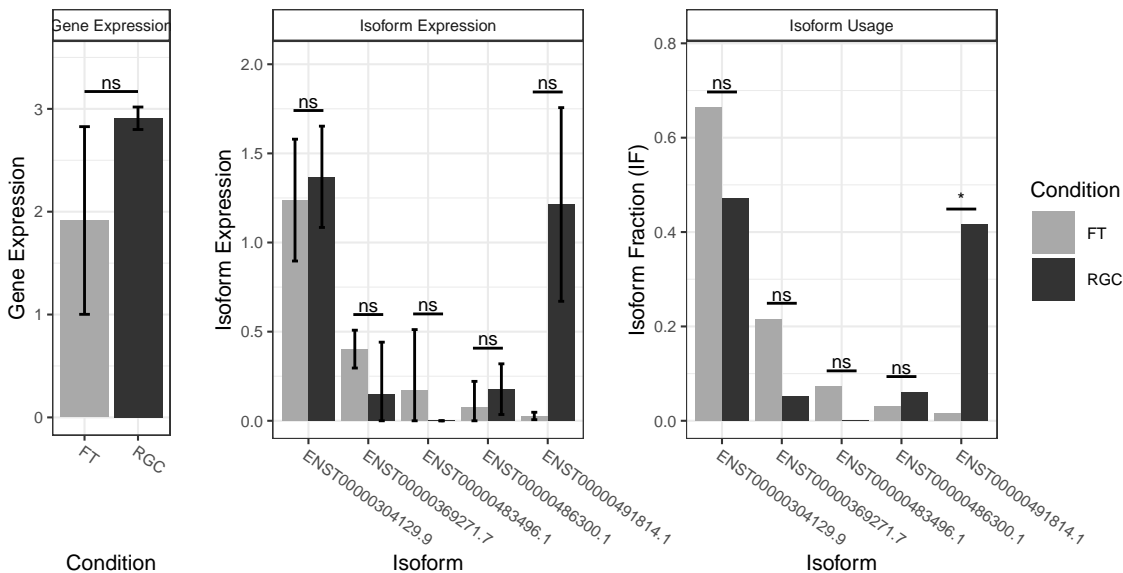
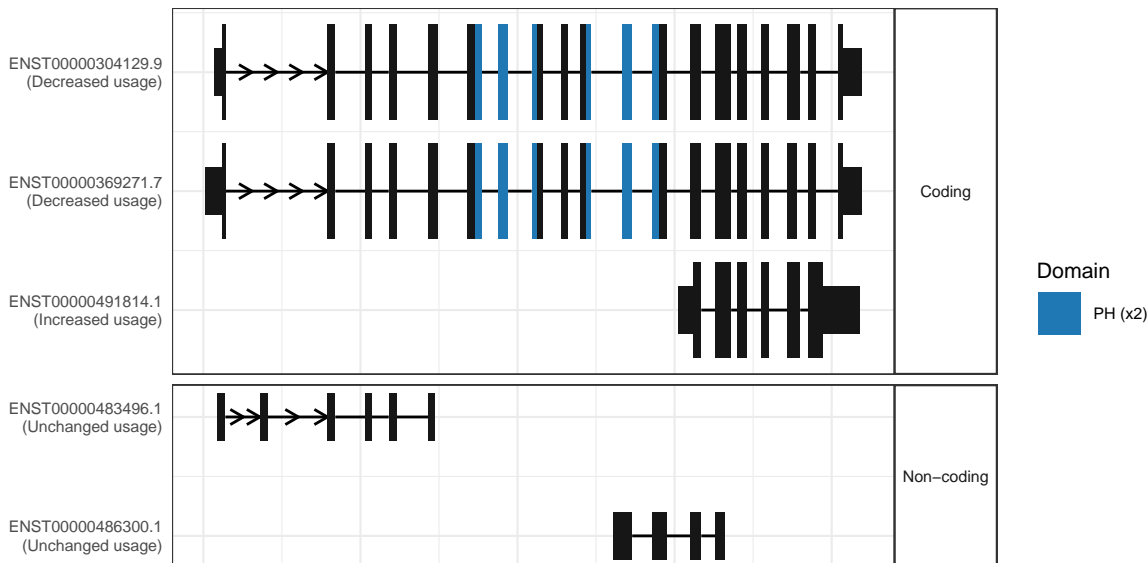
Domain



C1_1

RH_dom

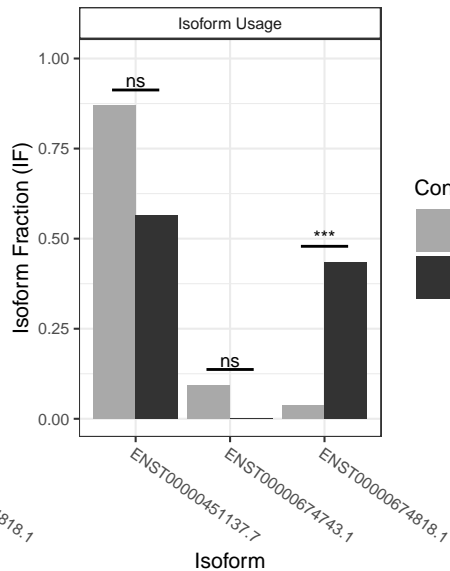
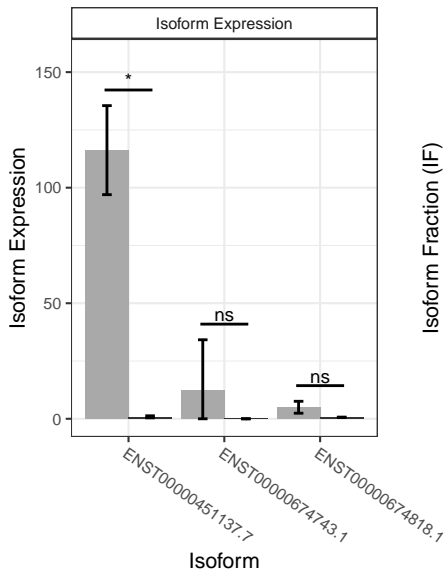
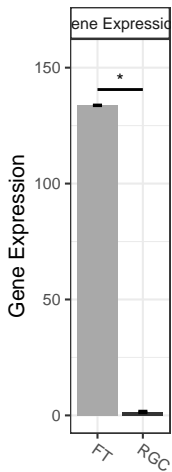
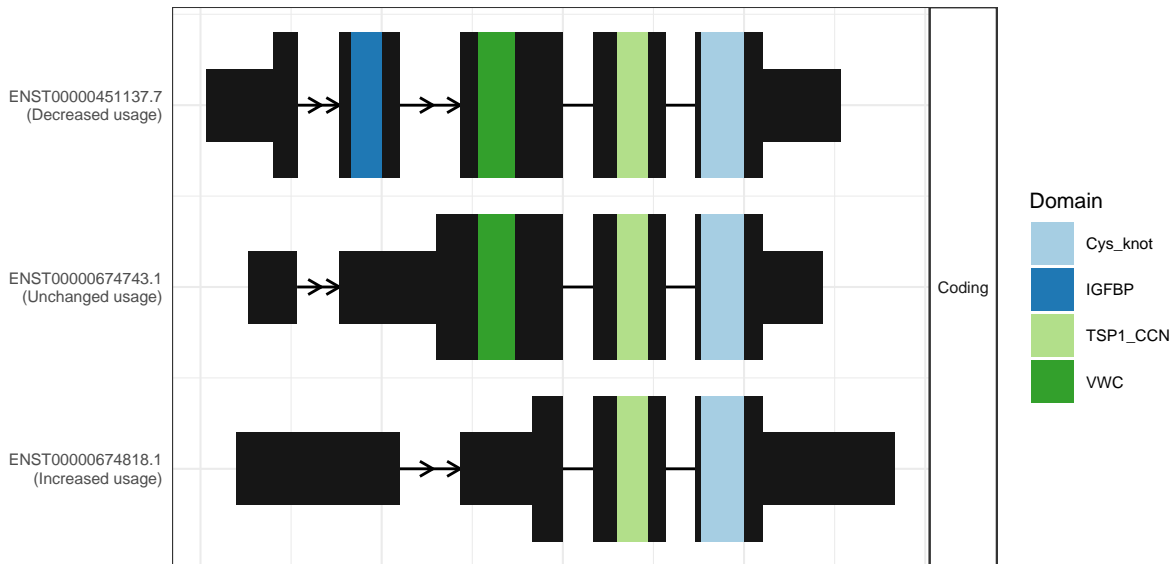
The isoform switch in AFAP1L2 (FT vs RGC)



Domain

 PH (x2)

The isoform switch in CCN1 (FT vs RGC)



Domain



Cys_knot



IGFBP

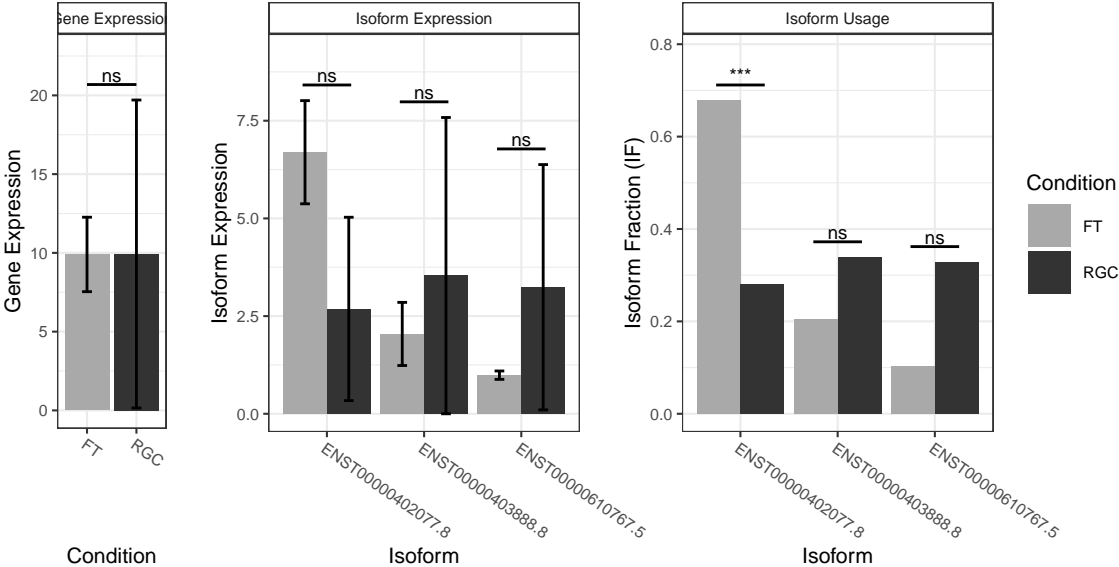
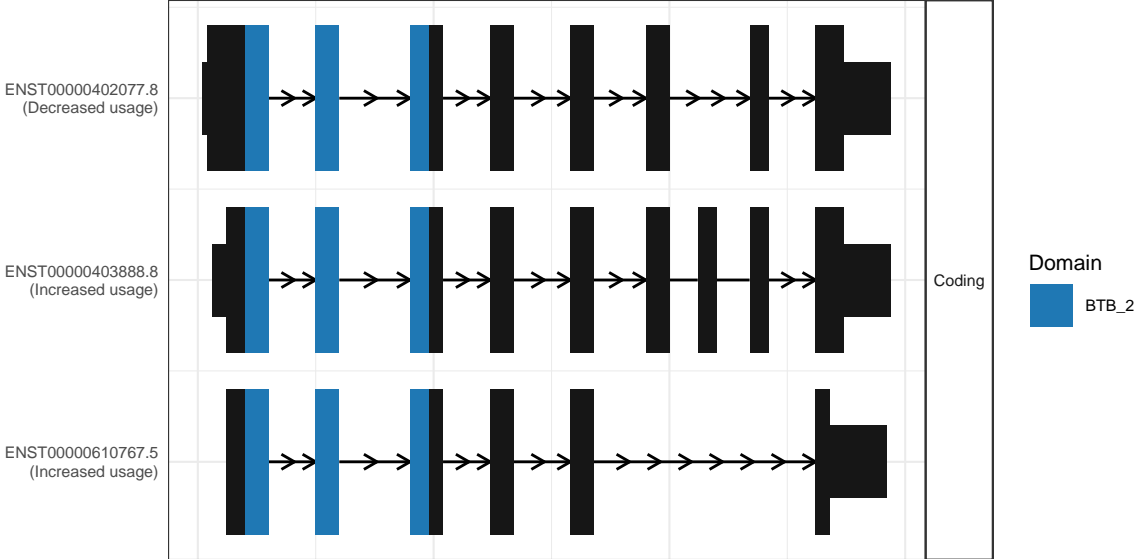


TSP1_CCN



VWC

The isoform switch in KCTD17 (FT vs RGC)

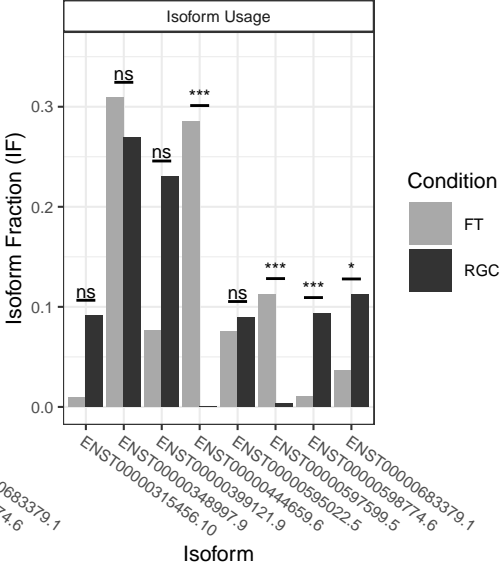
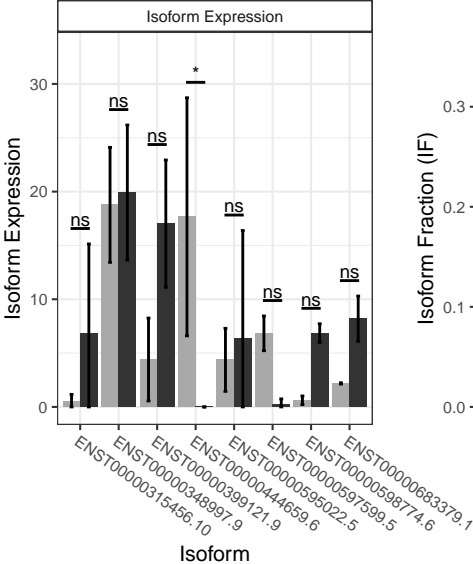
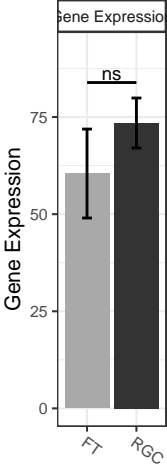
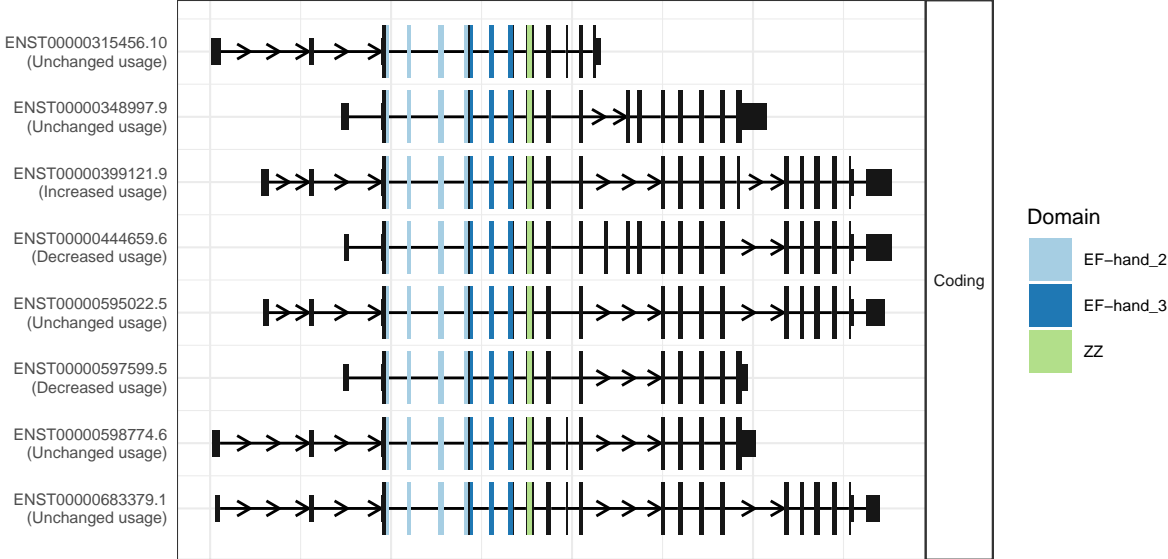


Domain



BTB_2

The isoform switch in DTNA (FT vs RGC)



Domain



EF-hand_2

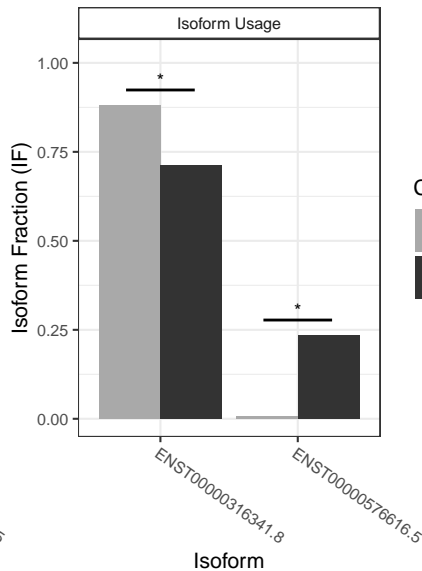
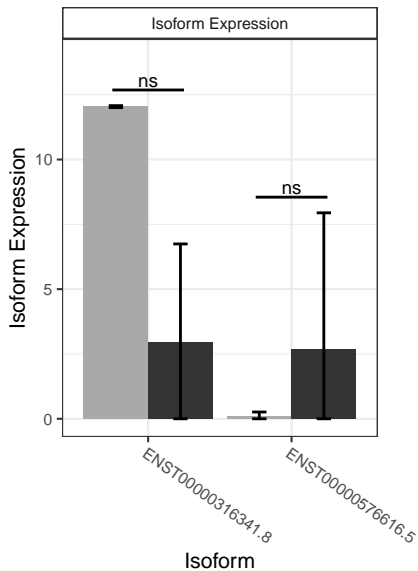
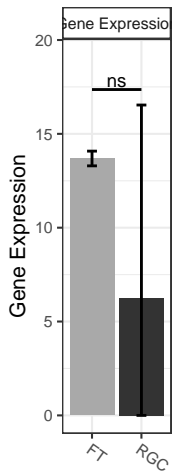
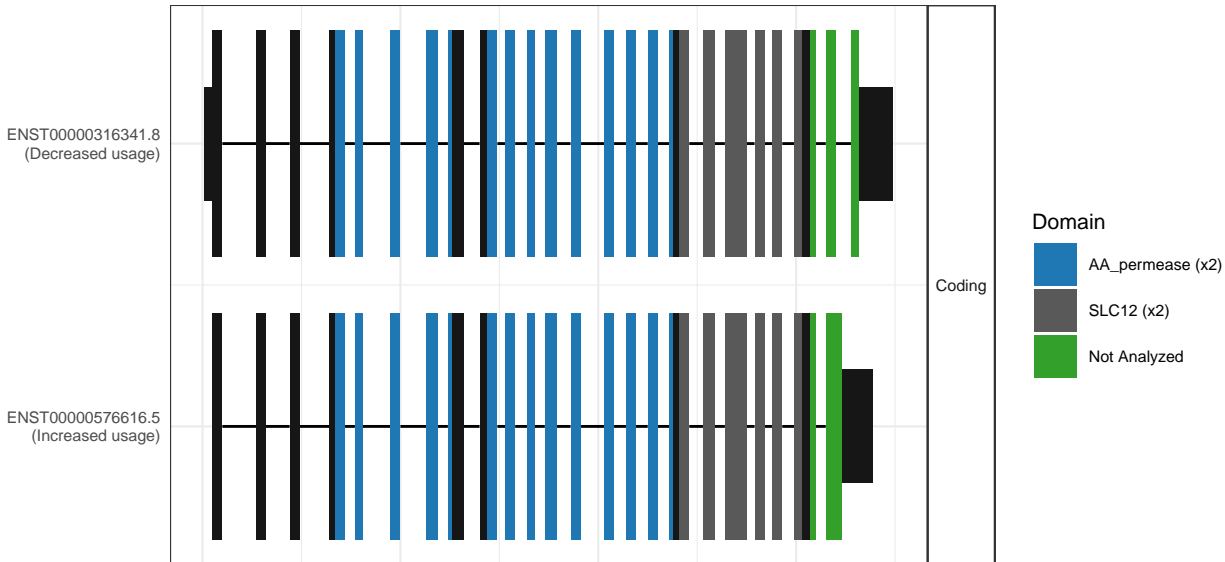


EF-hand_3






ZZ

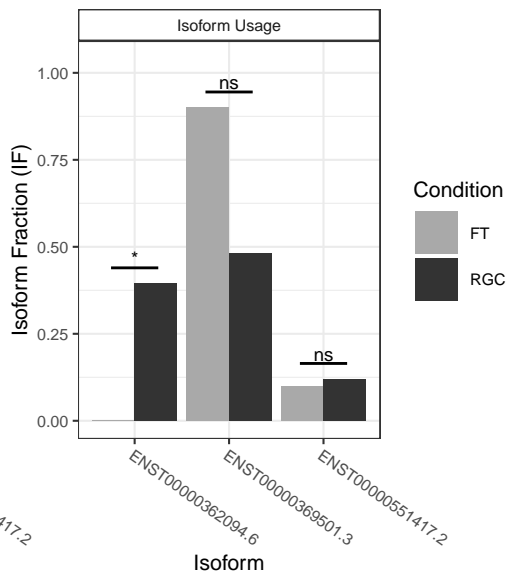
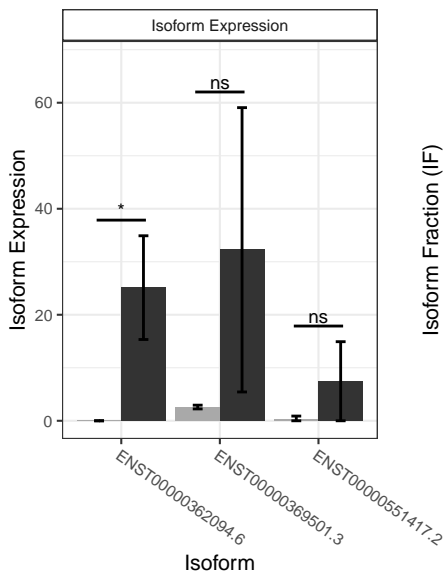
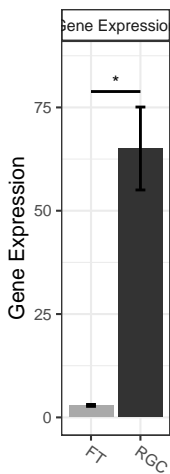
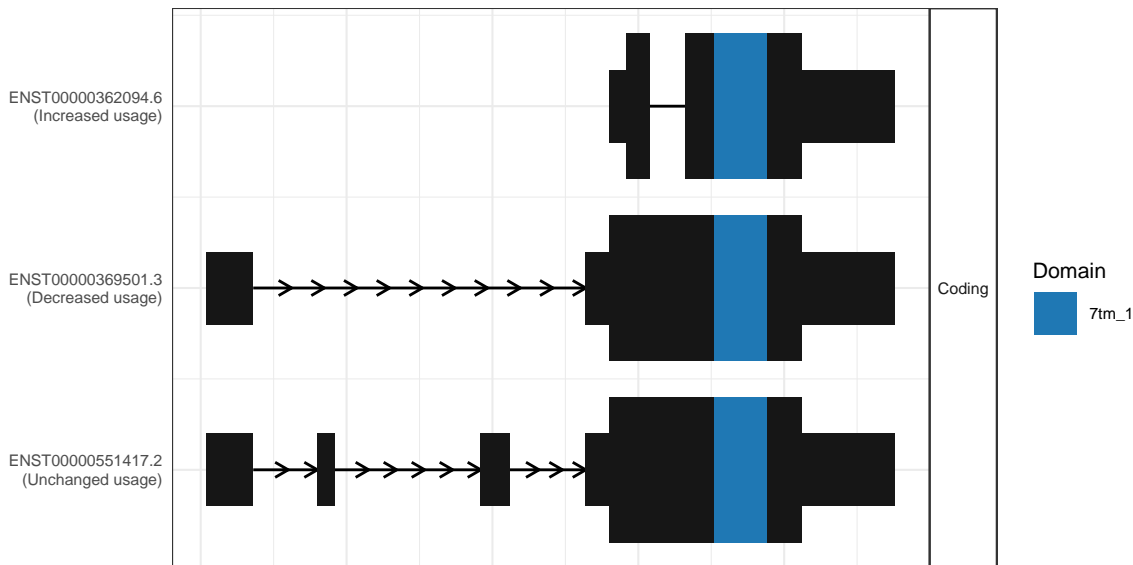
The isoform switch in SLC12A4 (FT vs RGC)



Domain

	AA_permease (x2)
	SLC12 (x2)
	Not Analyzed

The isoform switch in CNR1 (FT vs RGC)

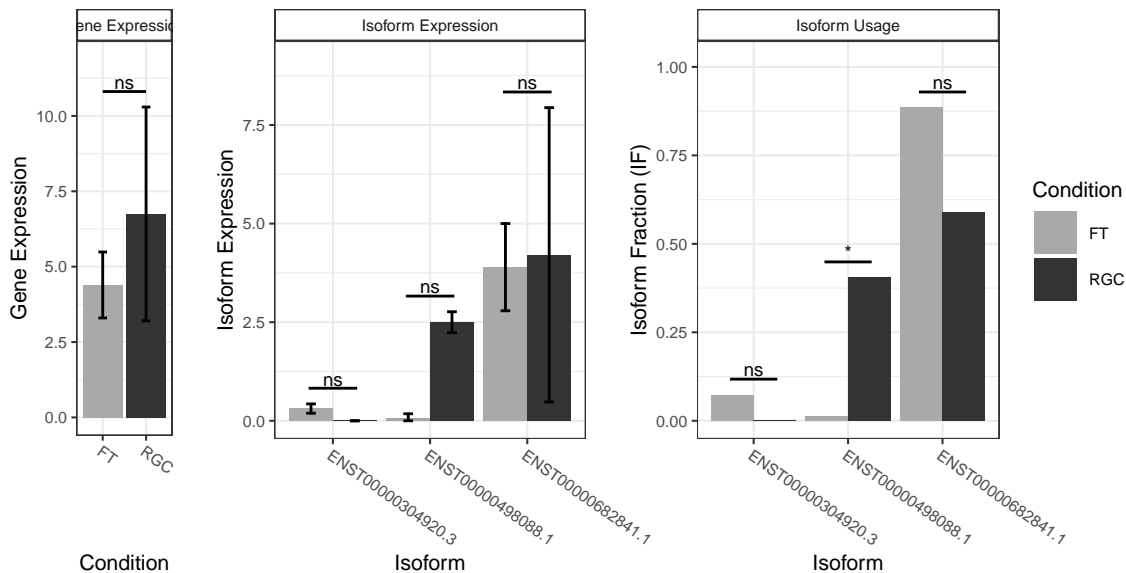
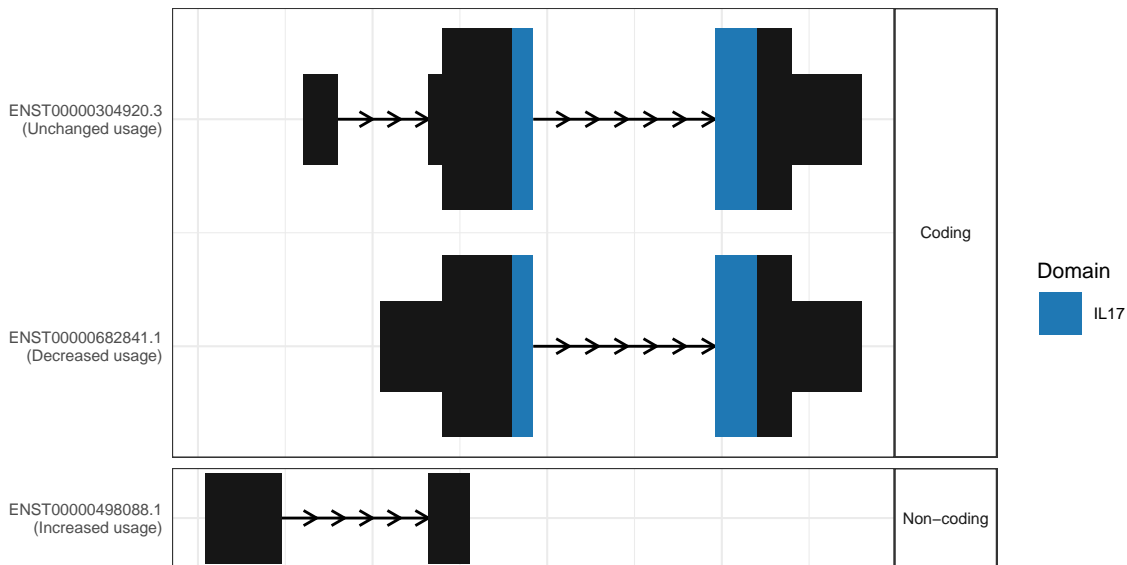


Domain



7tm_1

The isoform switch in IL17D (FT vs RGC)

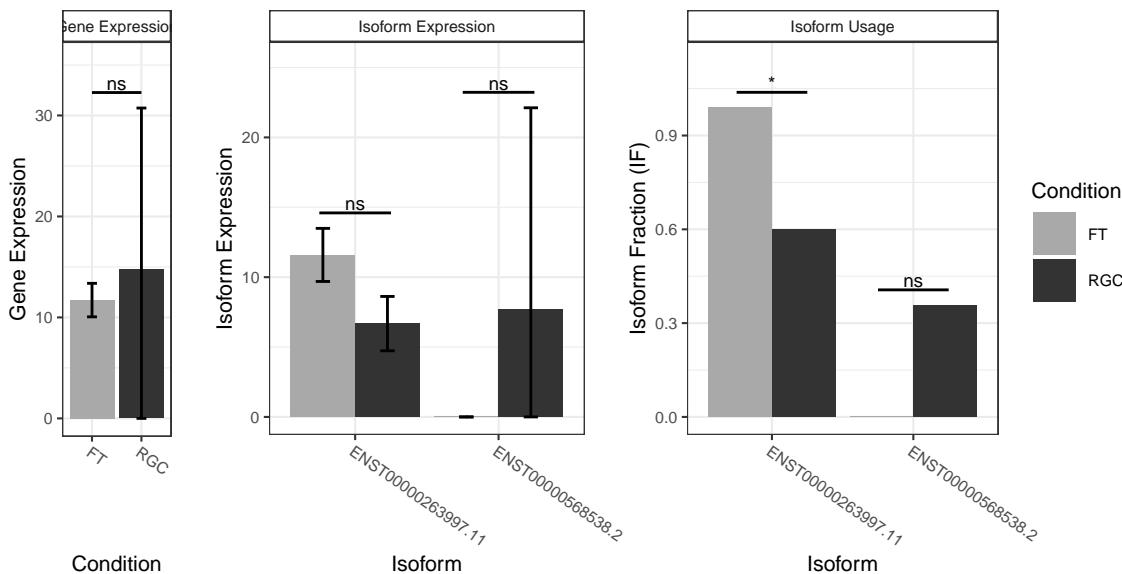
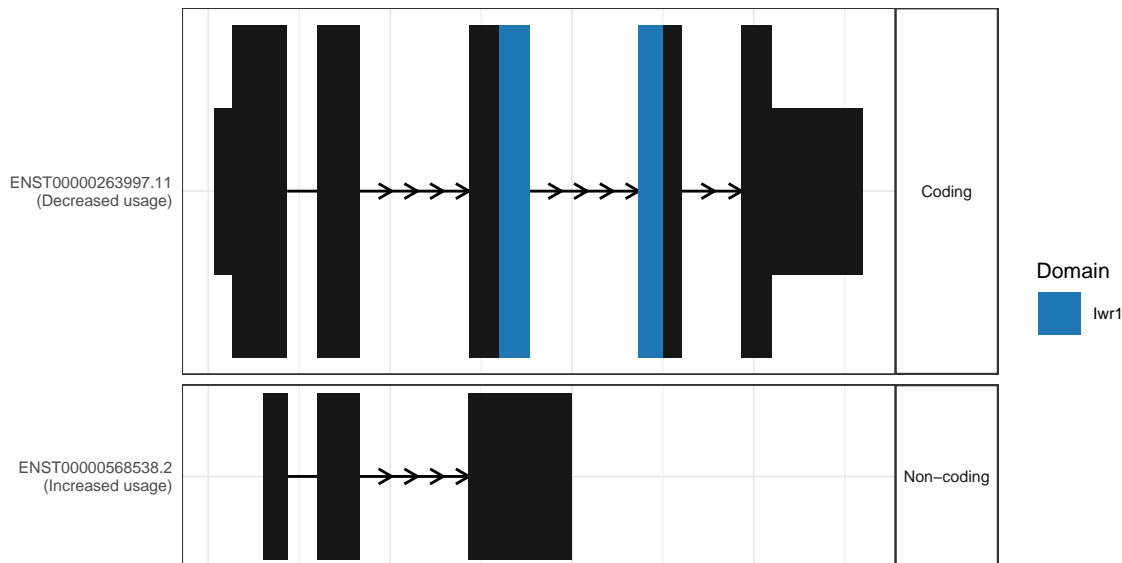


Domain



IL17

The isoform switch in SLC7A6OS (FT vs RGC)

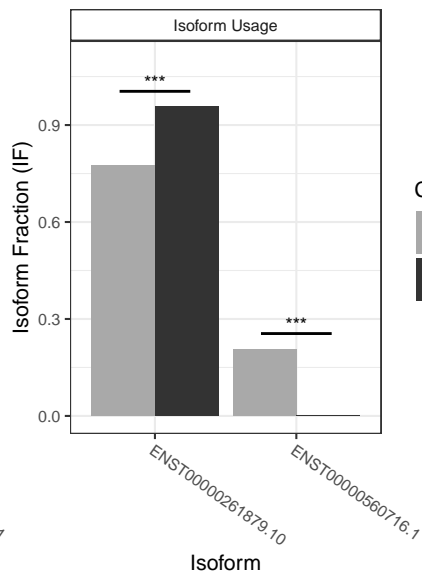
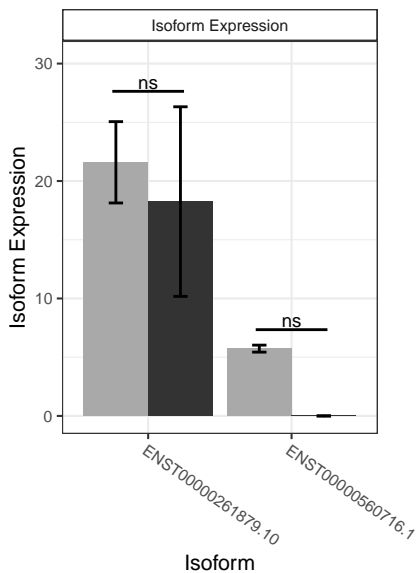
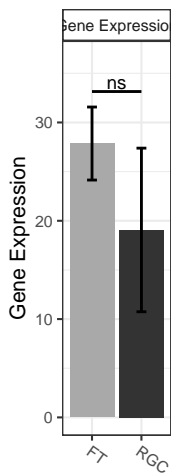
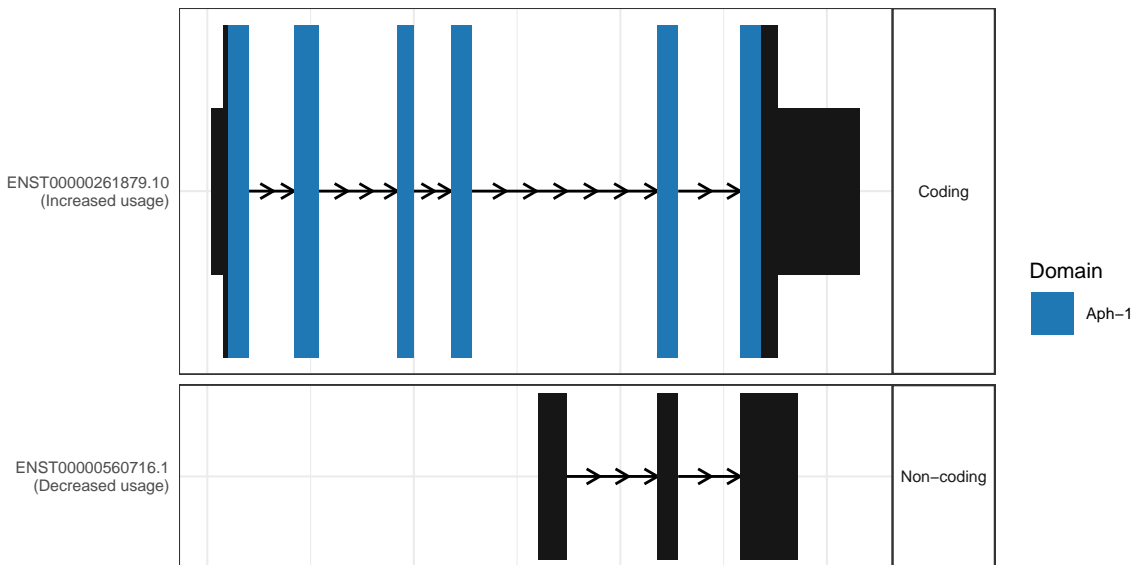


Domain



lwr1

The isoform switch in APh1B (FT vs RGC)

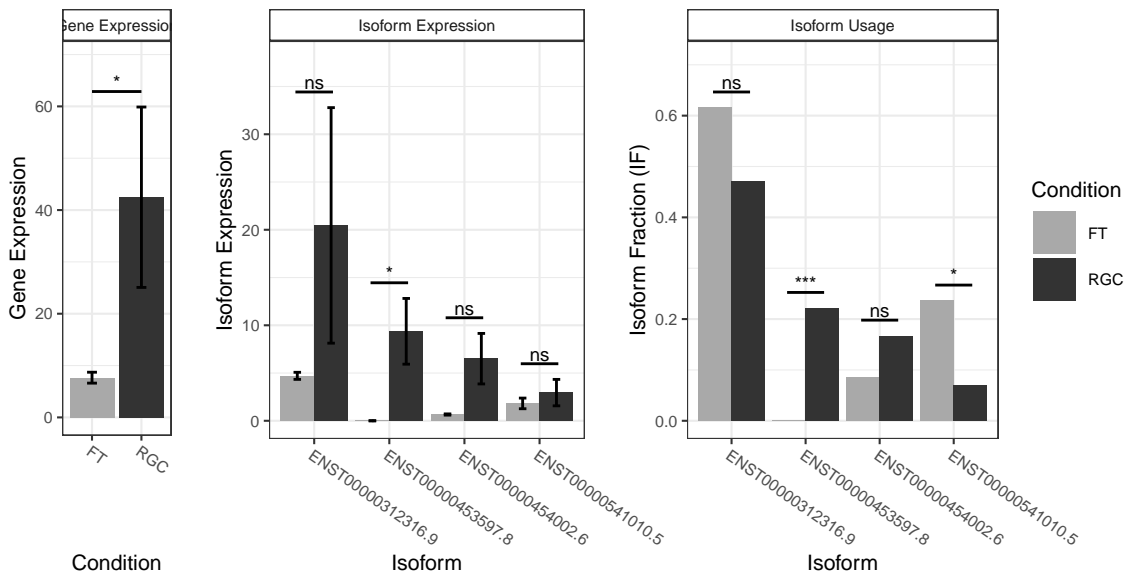
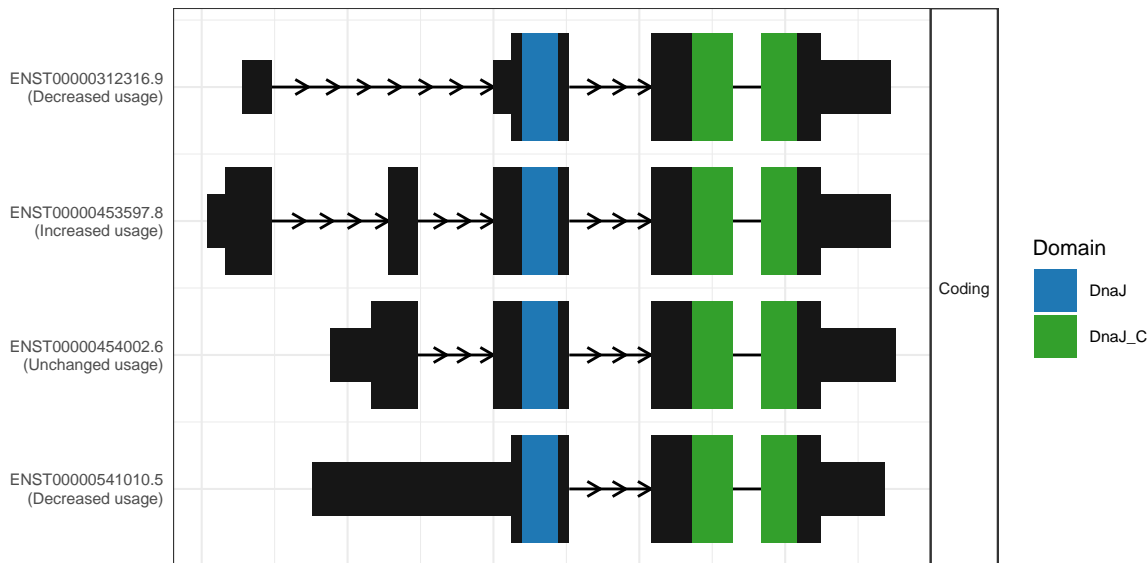


Domain



Aph-1

The isoform switch in DNAJB5 (FT vs RGC)



Domain

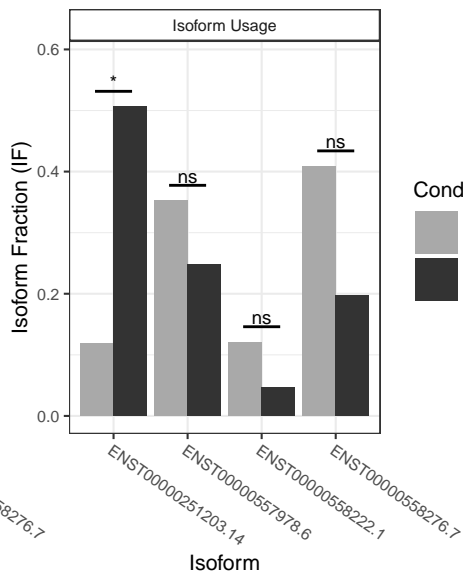
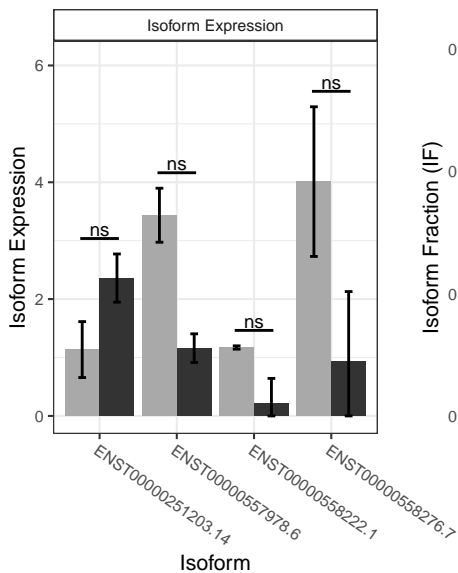
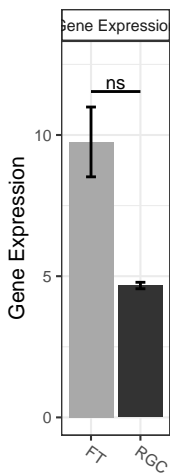
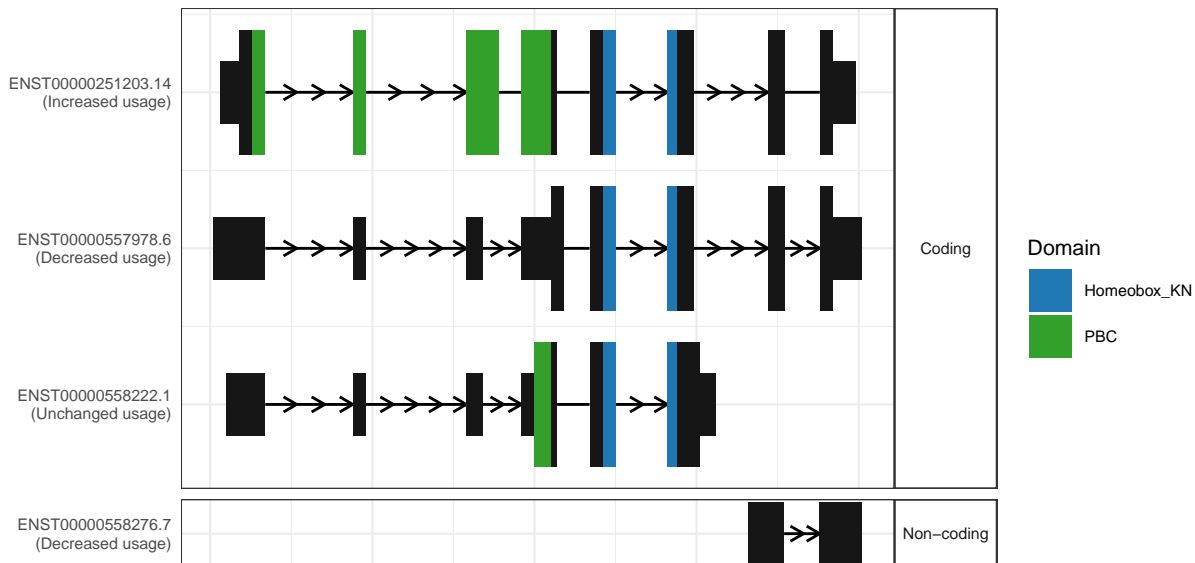


DnaJ



DnaJ_C

The isoform switch in PBX4 (FT vs RGC)



Condition

Isoform

Isoform

Condition



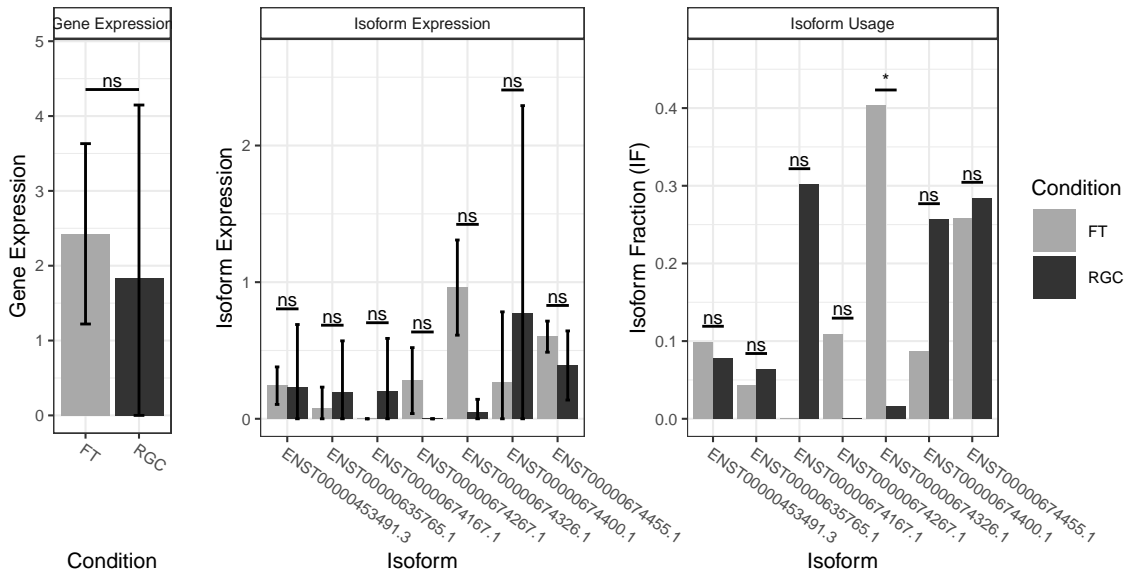
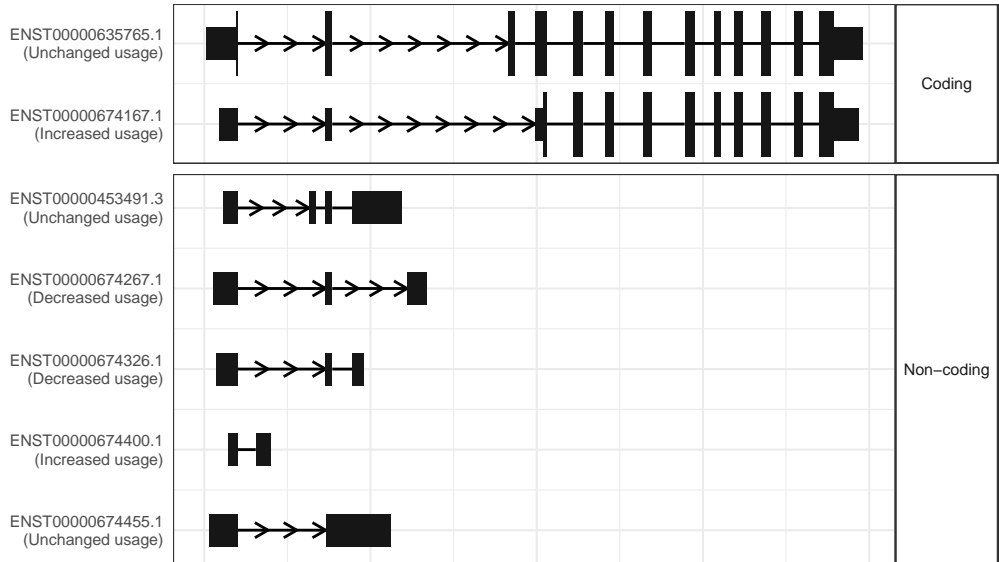
Domain



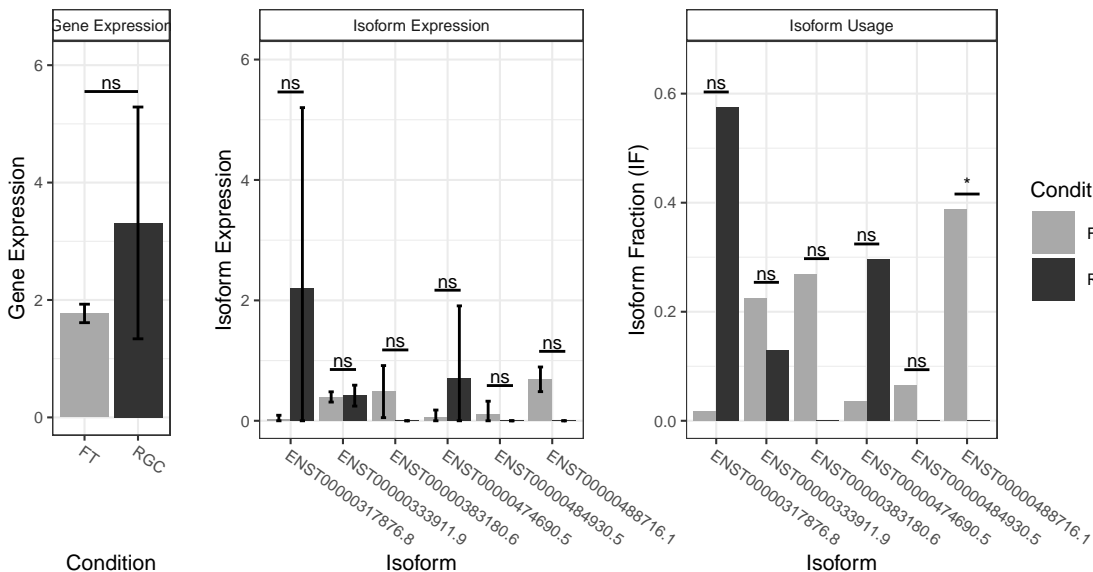
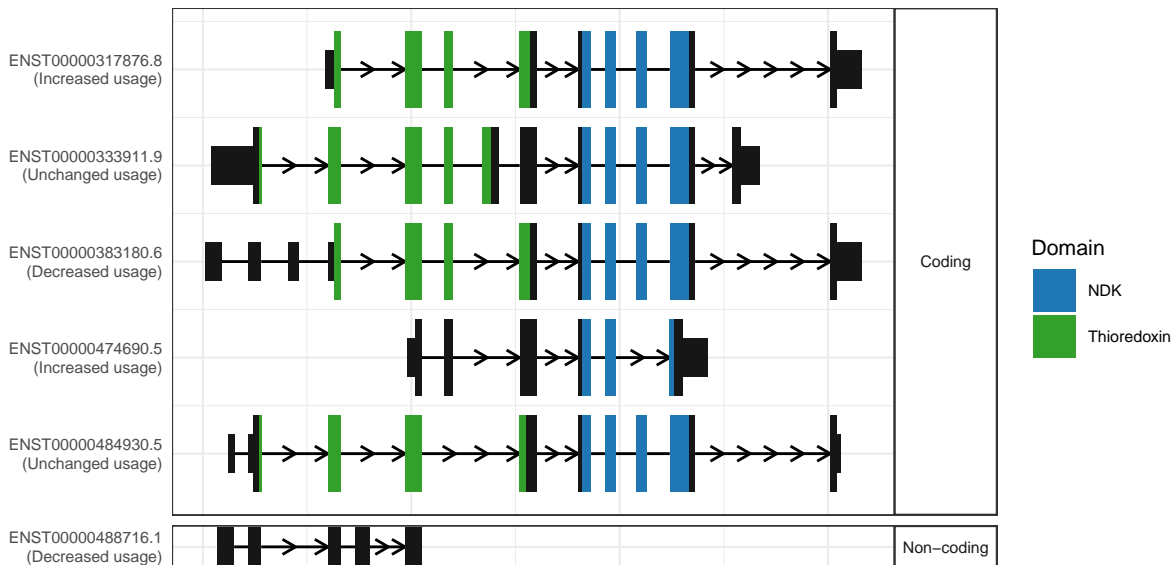
Homeobox_KN

PBC

The isoform switch in HSPA12A (FT vs RGC)



The isoform switch in NME9 (FT vs RGC)



Domain

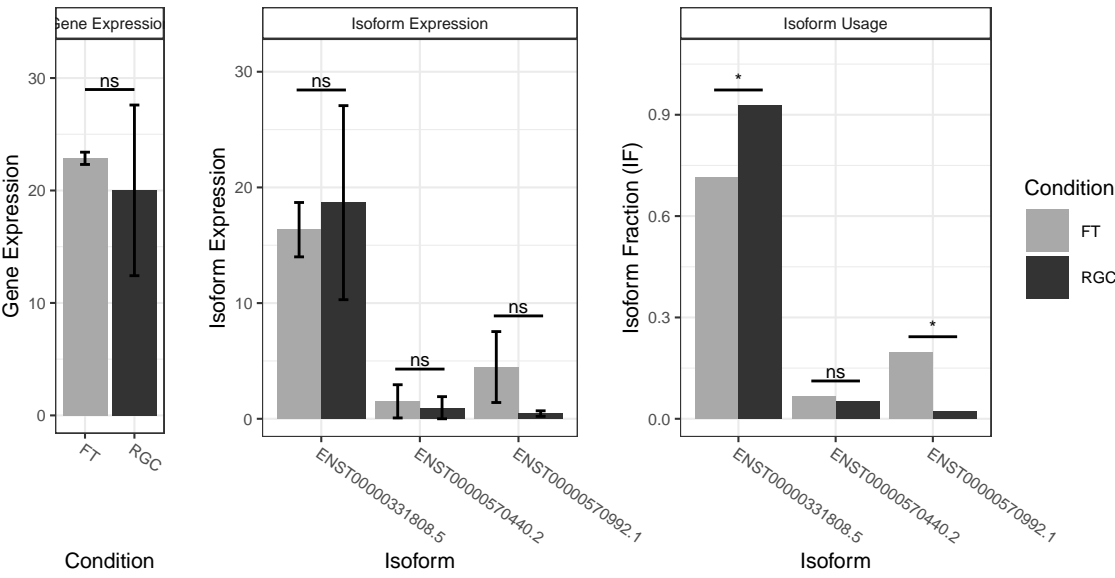
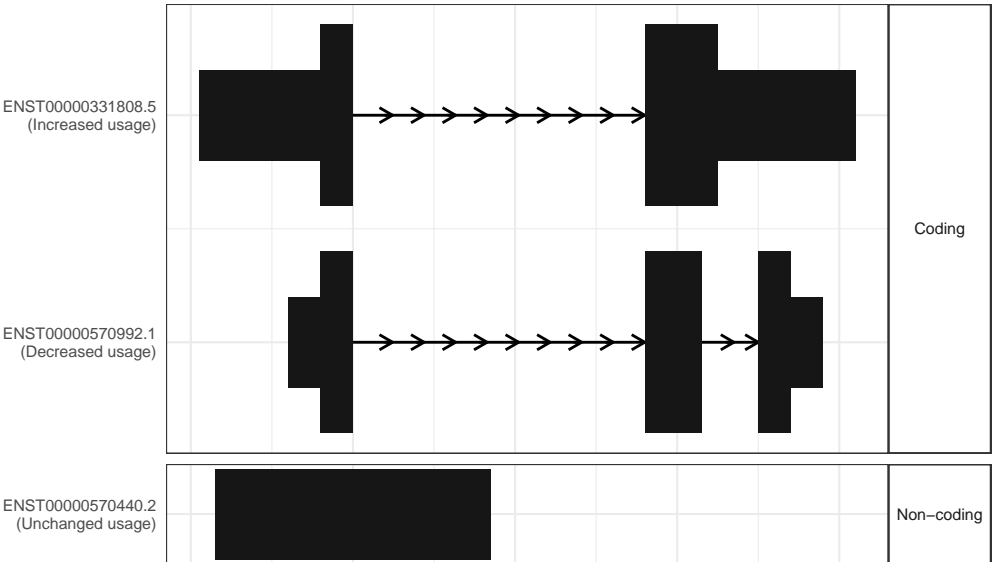


NDK

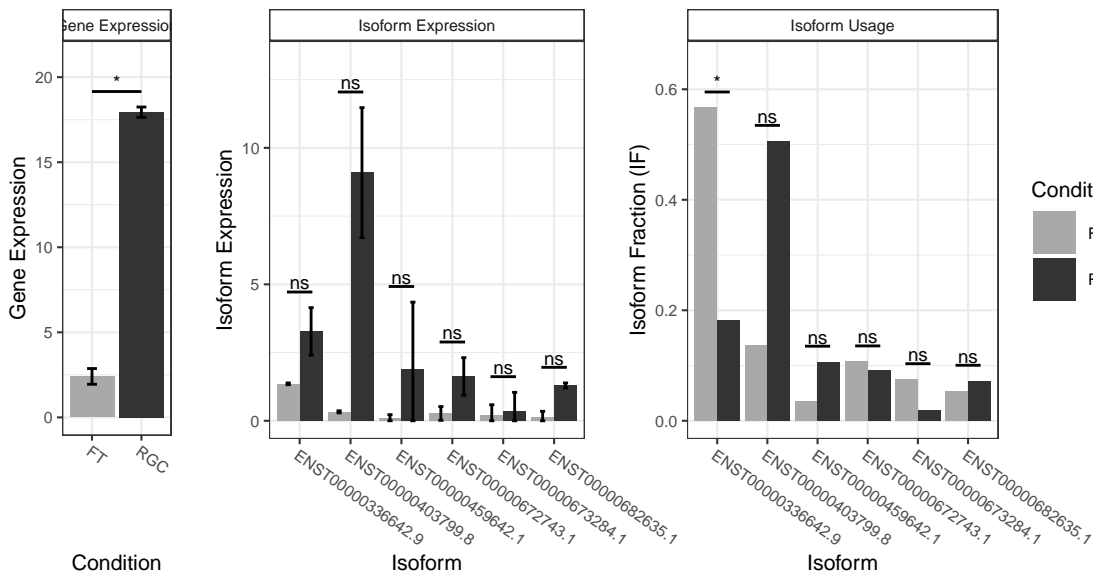
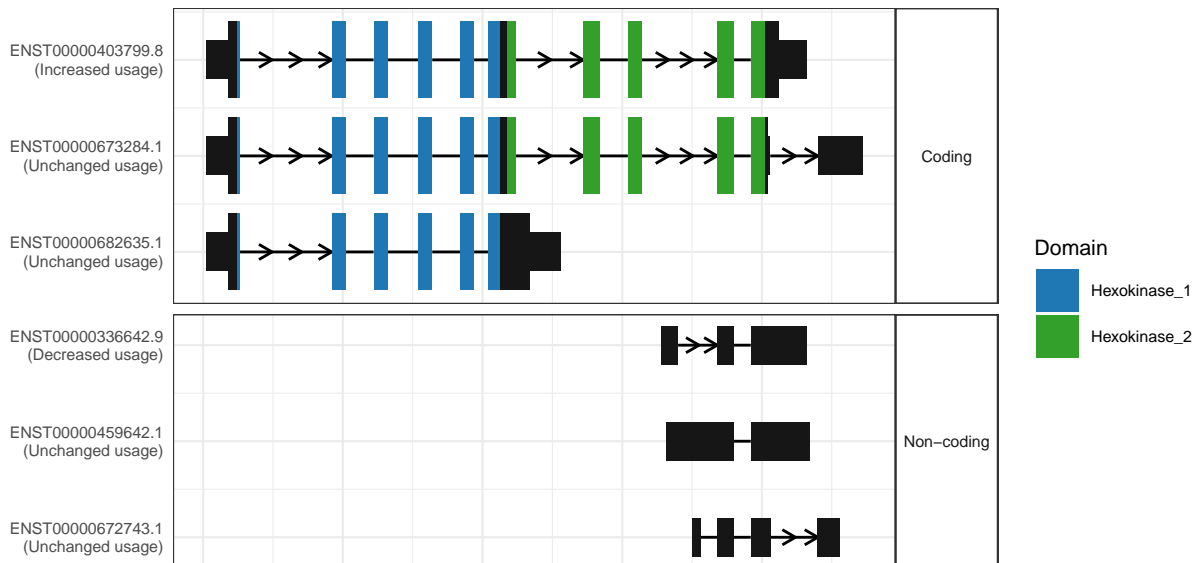


Thioredoxin

The isoform switch in DEXI (FT vs RGC)



The isoform switch in GCK (FT vs RGC)



Domain

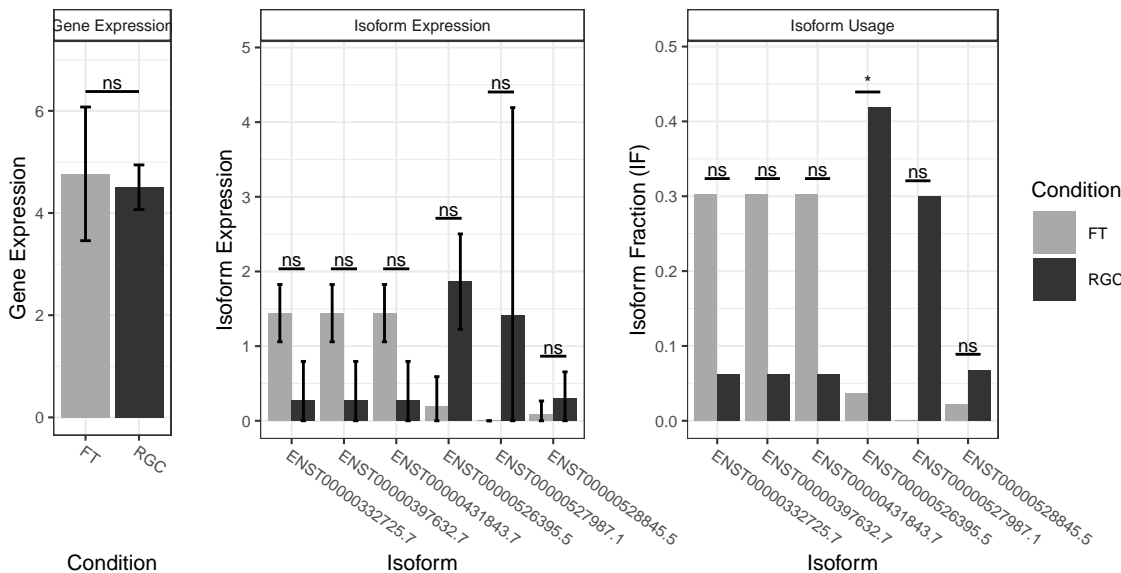
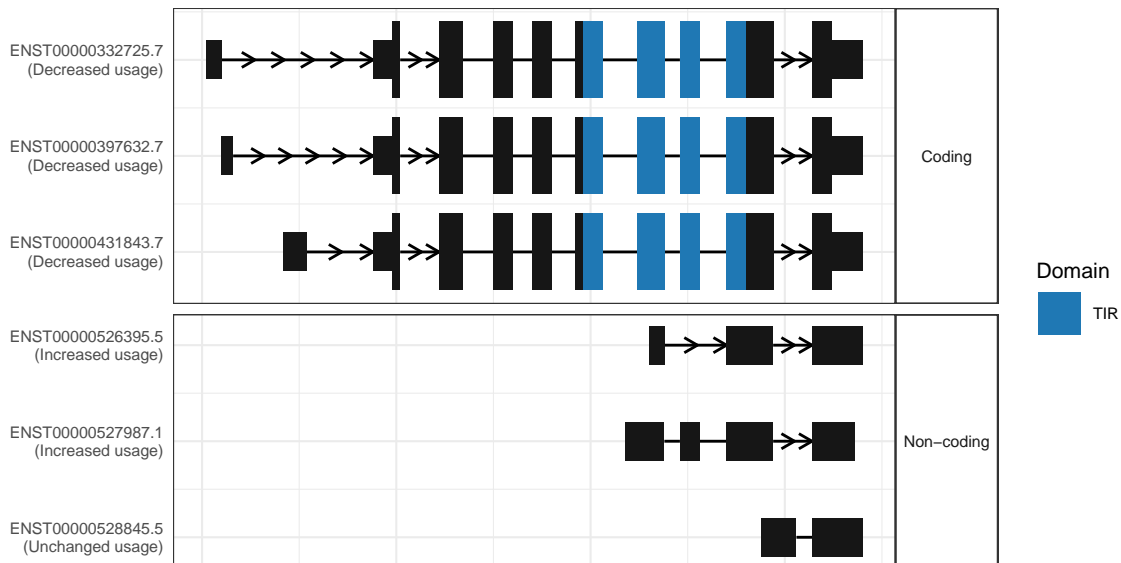


Hexokinase_1



Hexokinase_2

The isoform switch in SIGIRR (FT vs RGC)

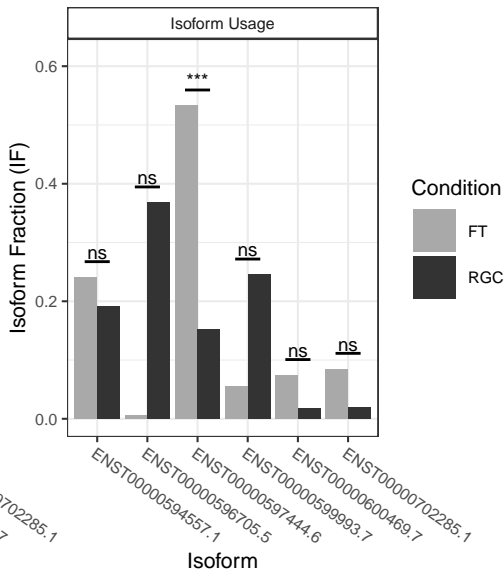
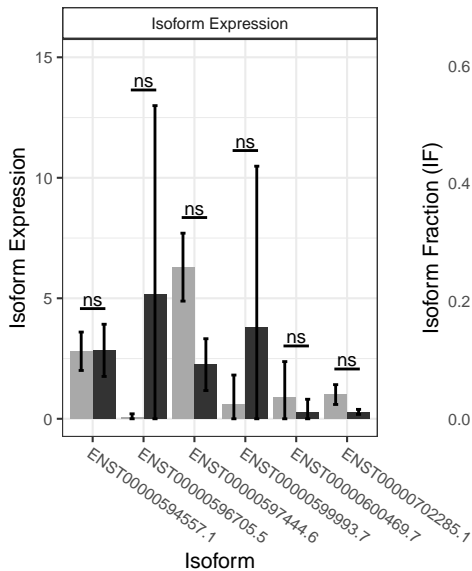
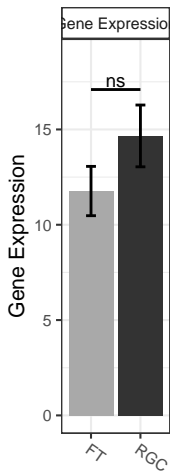
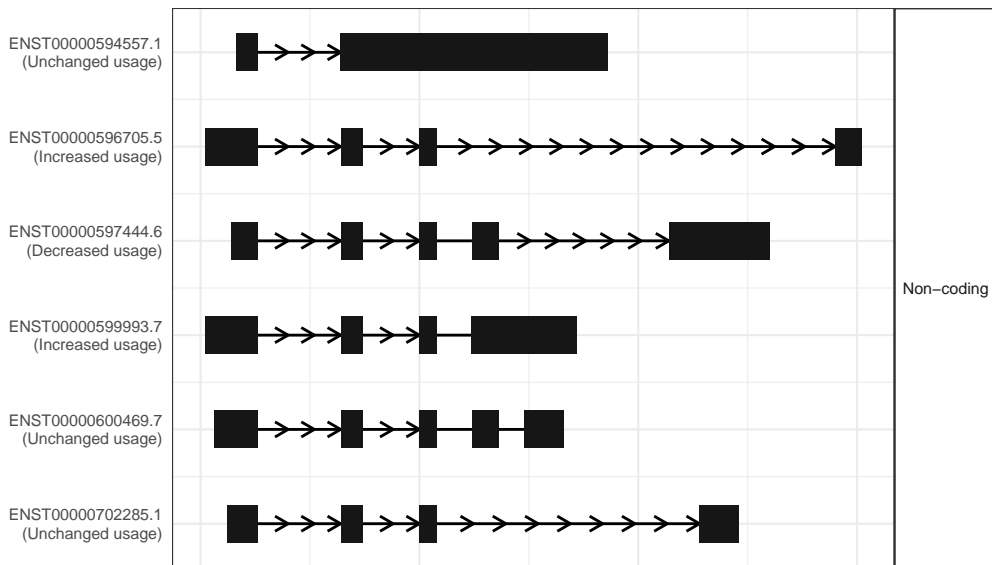


Domain

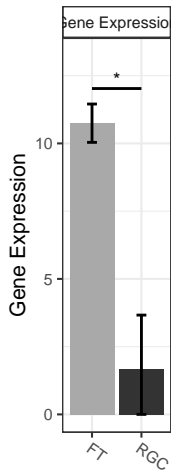
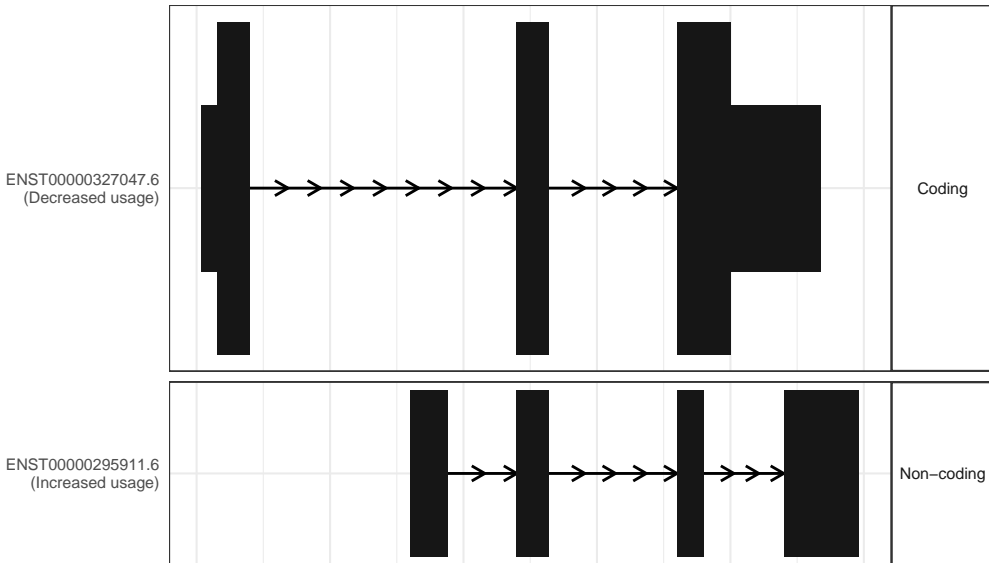


TIR

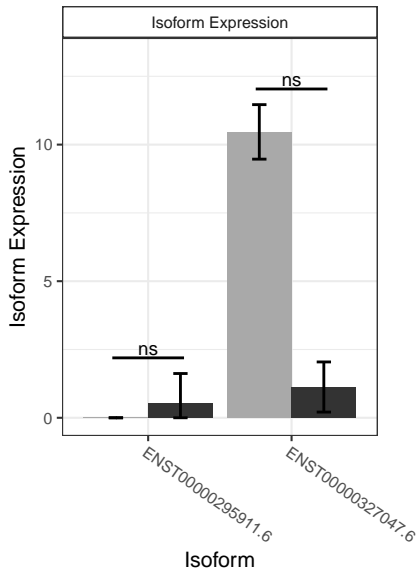
The isoform switch in (FT vs RGC)



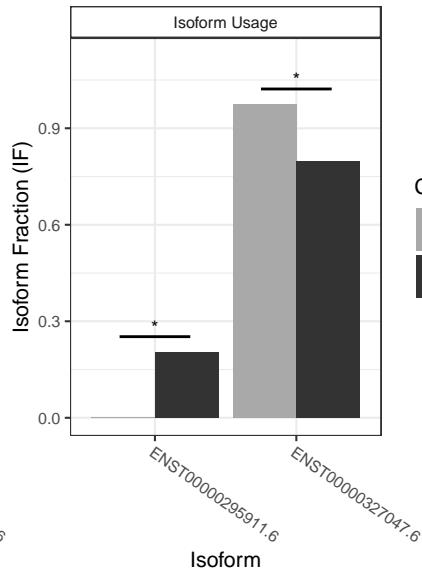
The isoform switch in CLRN1 (FT vs RGC)



Condition



Isoform

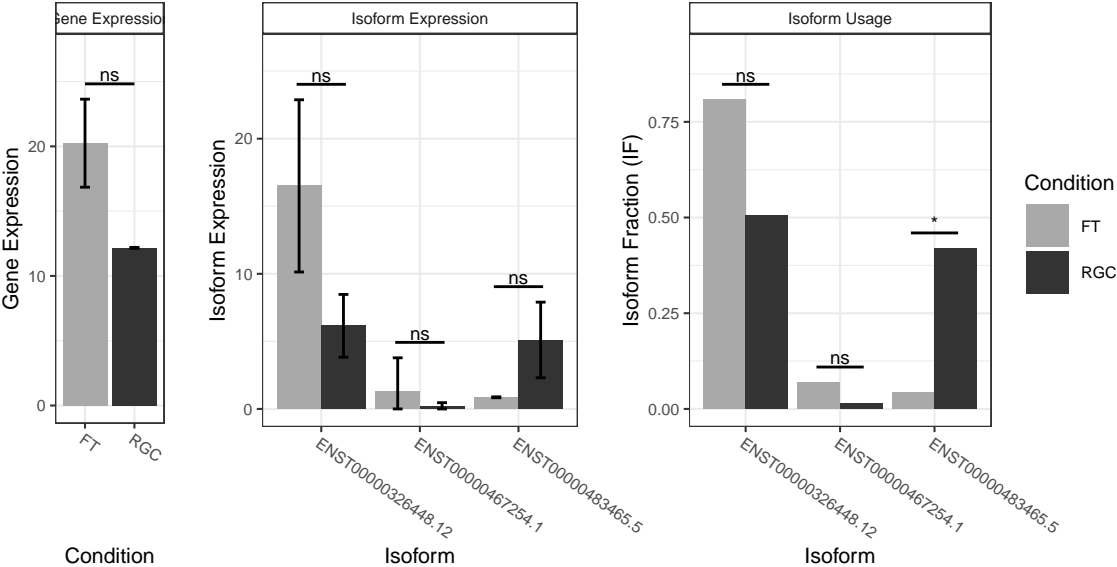
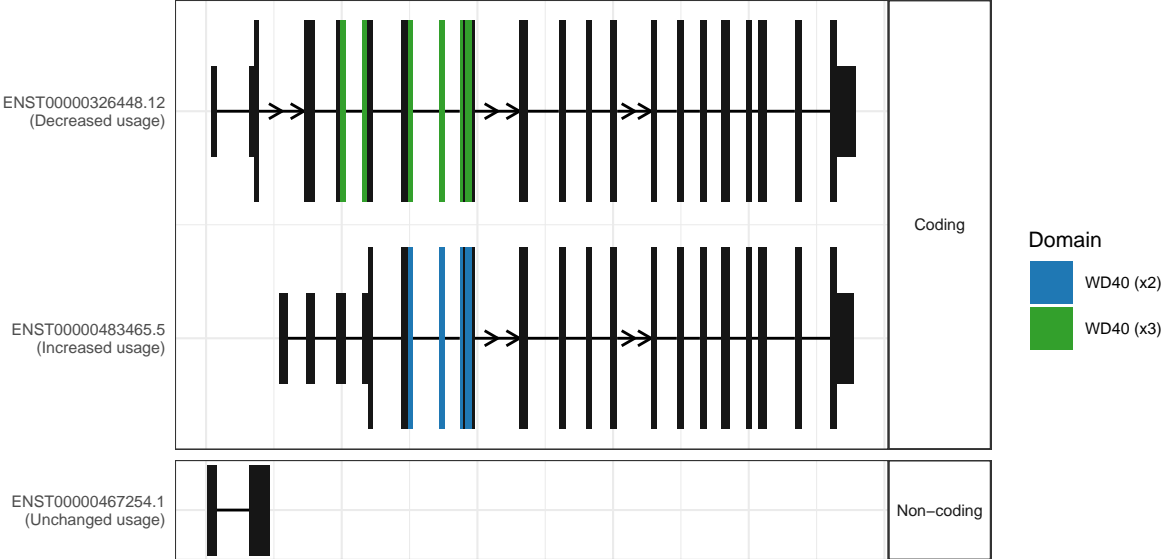


Isoform

Condition



The isoform switch in IFT80 (FT vs RGC)



Domain

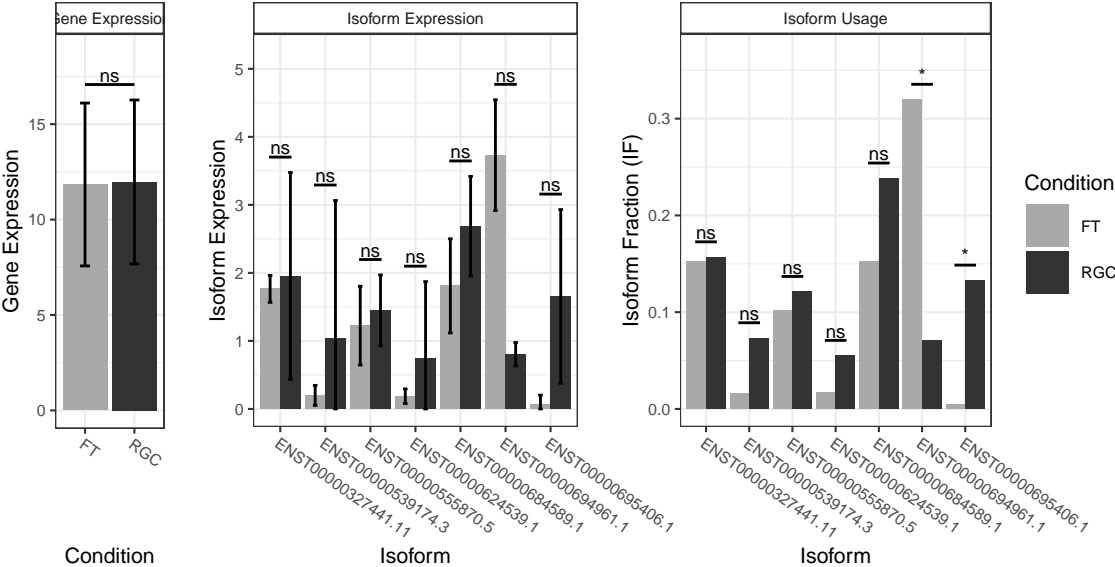
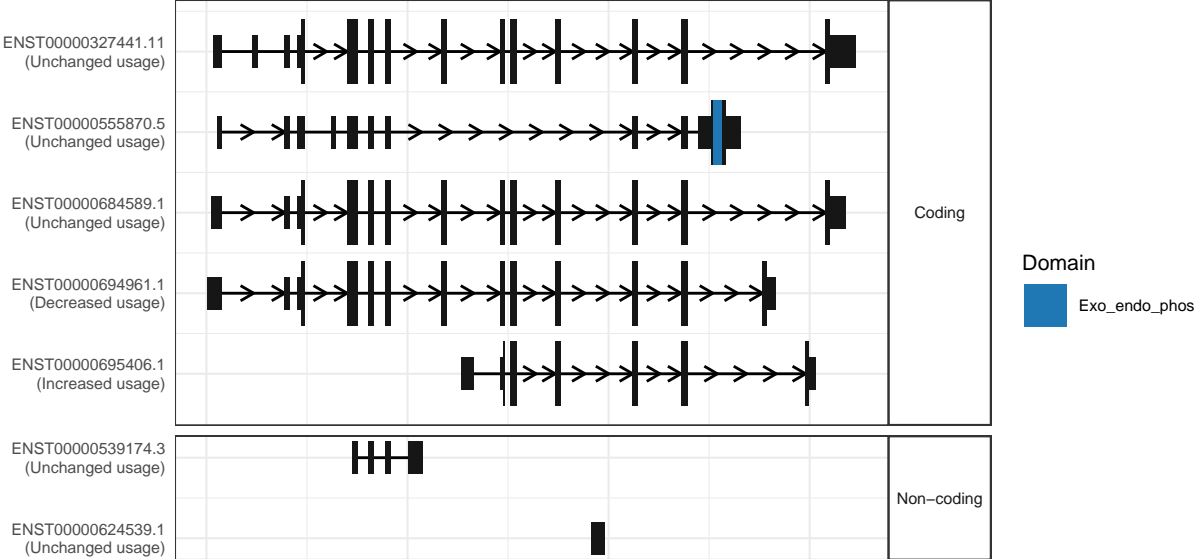


WD40 (x2)



WD40 (x3)

The isoform switch in MIPOL1 (FT vs RGC)

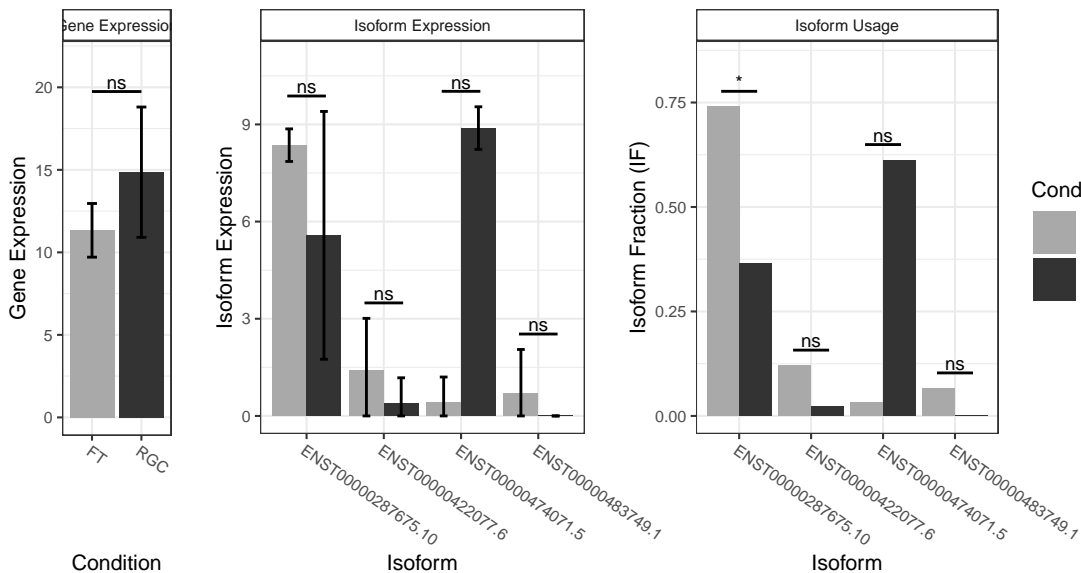
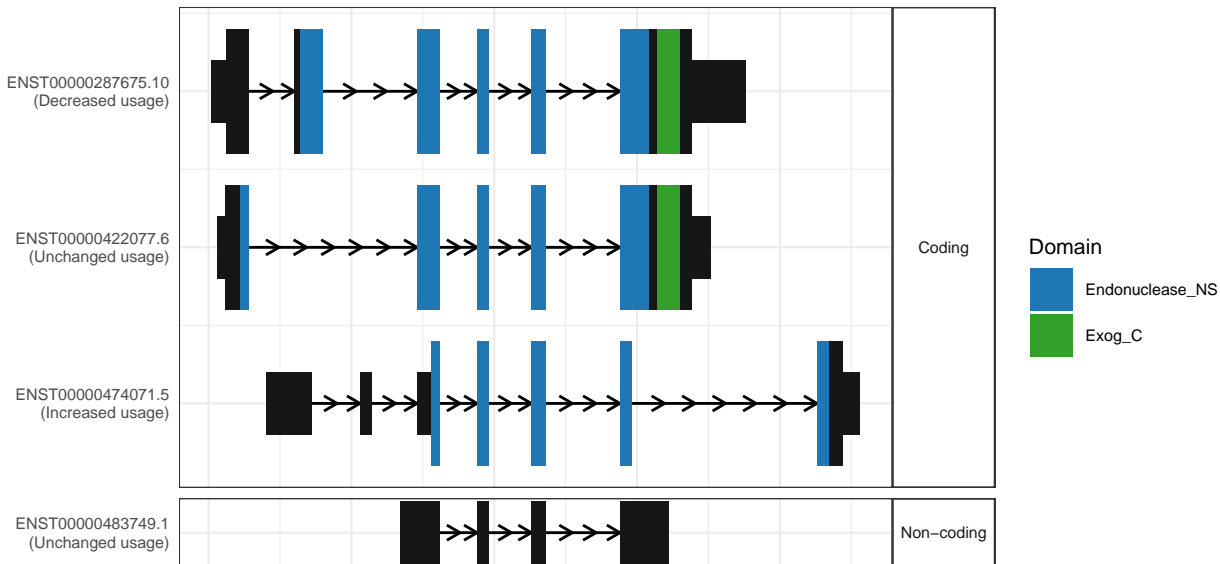


Domain





Exo_endo_phos

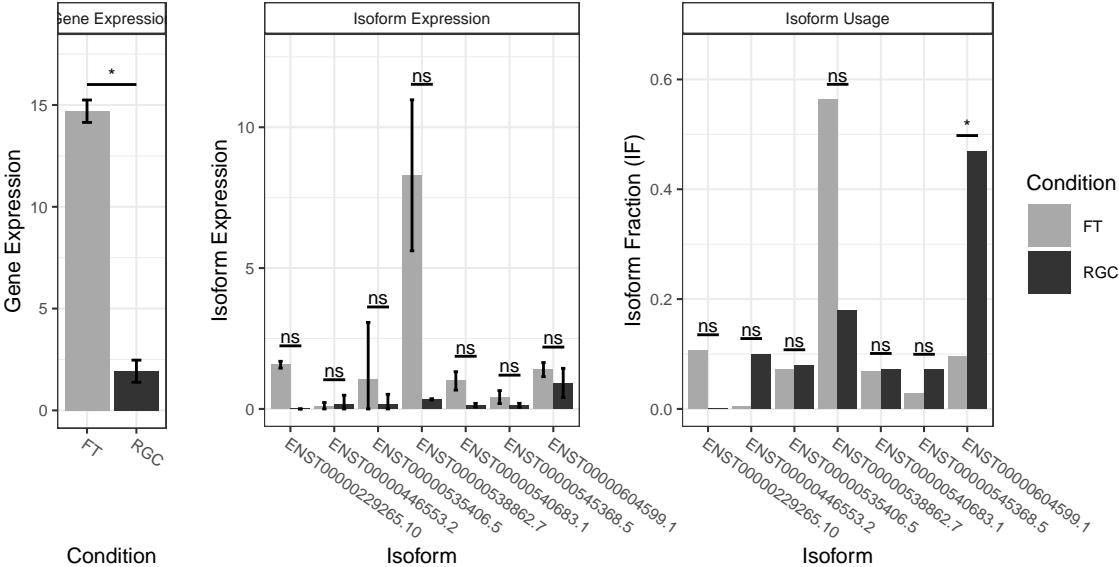
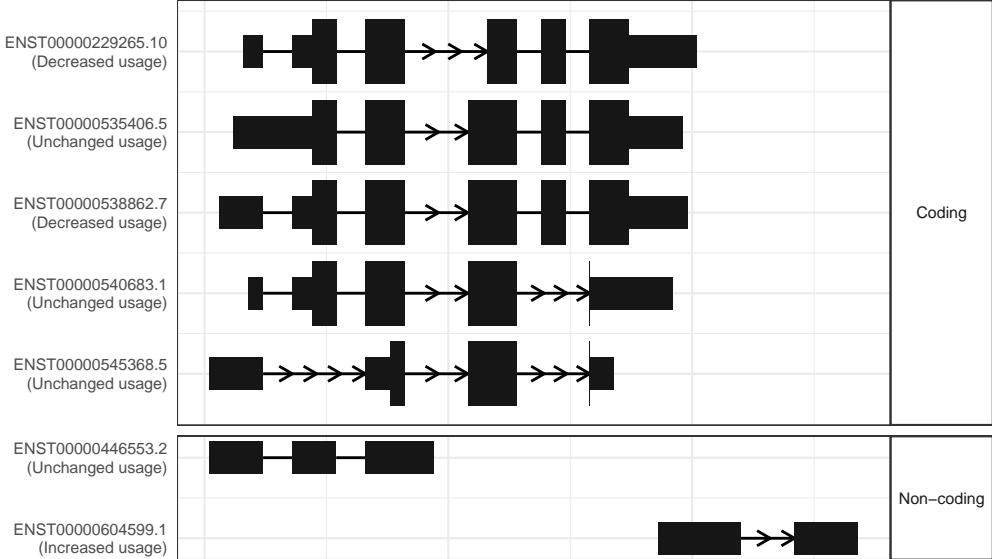
The isoform switch in EXOG (FT vs RGC)



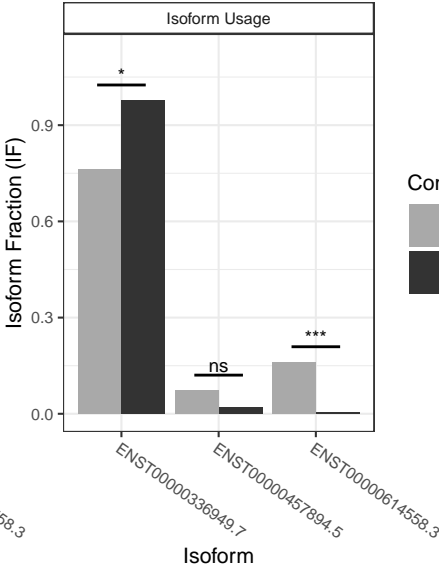
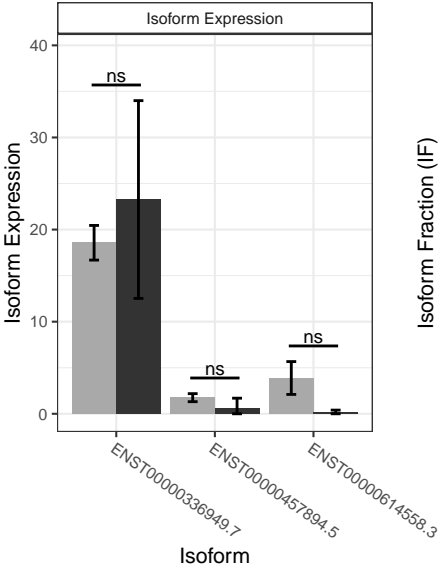
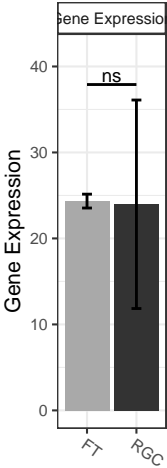
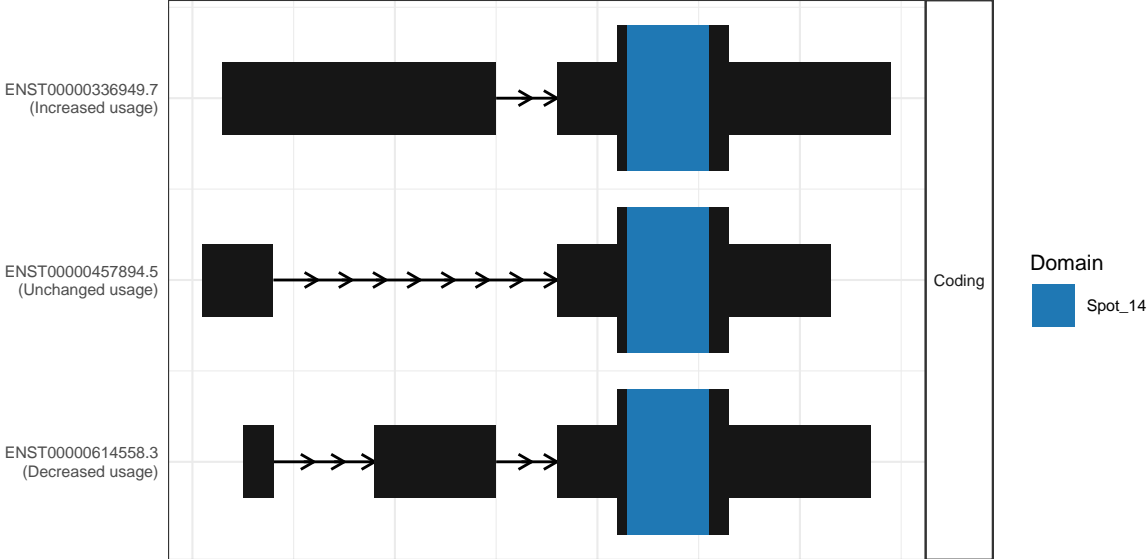
Domain

	Endonuclease_NS
	Exog_C

The isoform switch in CDCA3 (FT vs RGC)



The isoform switch in MID1IP1 (FT vs RGC)

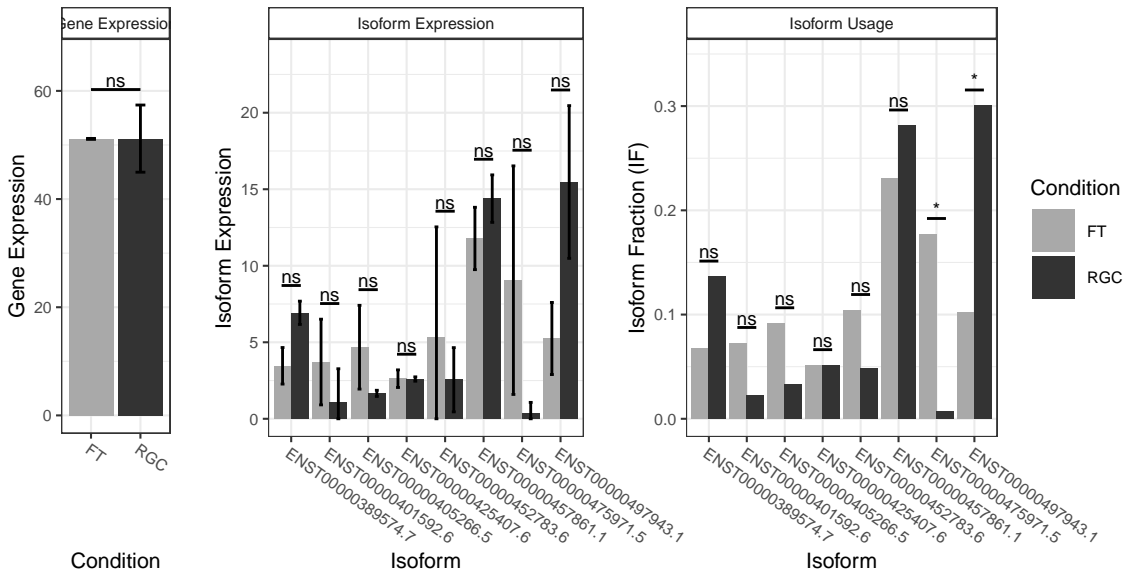
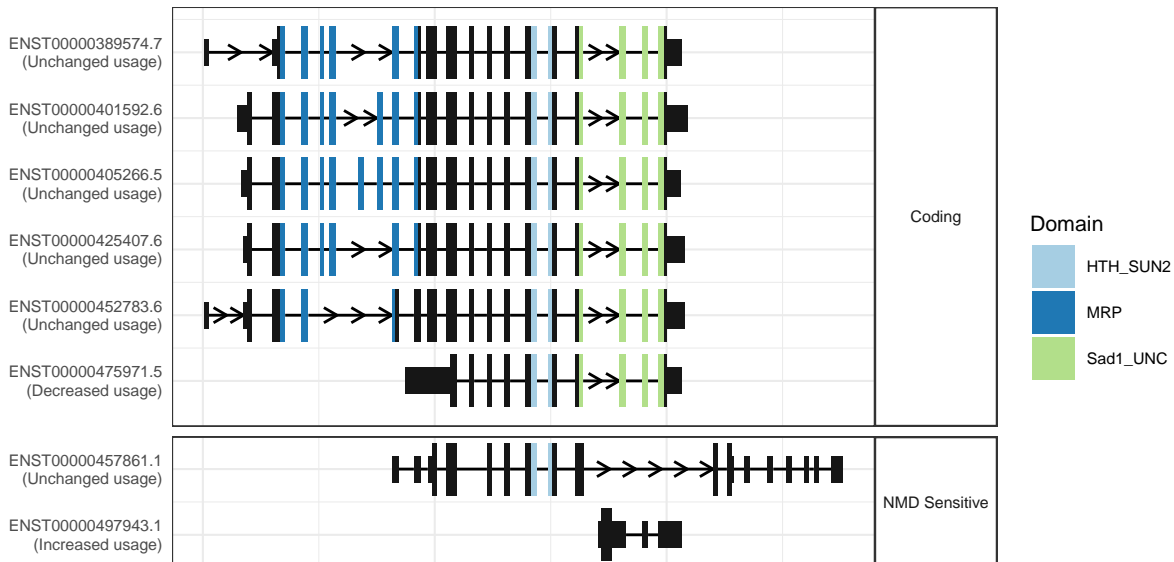


Domain



Spot_14

The isoform switch in SUN1 (FT vs RGC)



Domain



HTH_SUN2

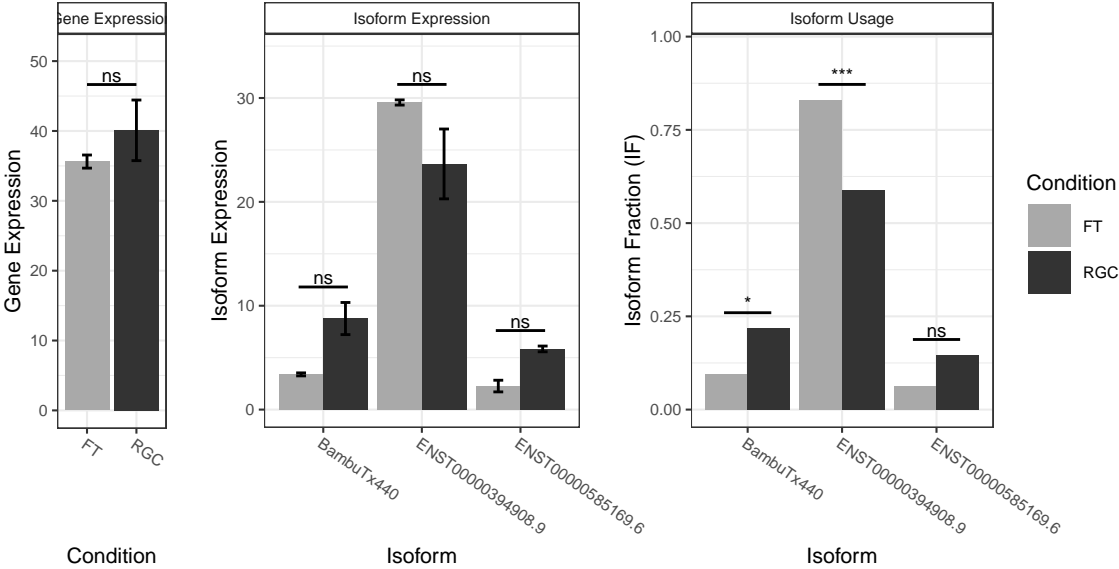
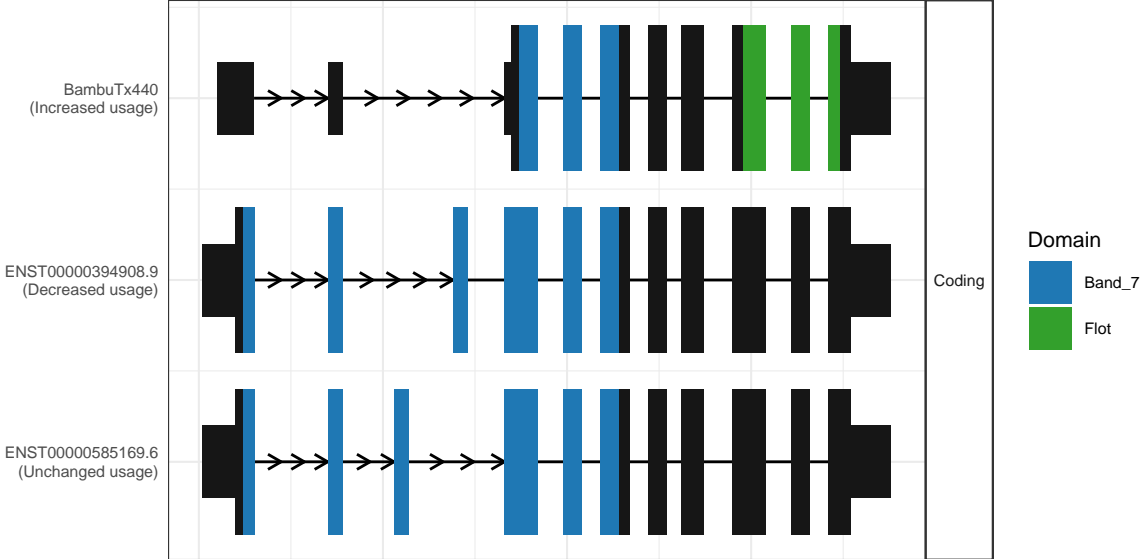


MRP



Sad1_UNC

The isoform switch in FLOT2 (FT vs RGC)



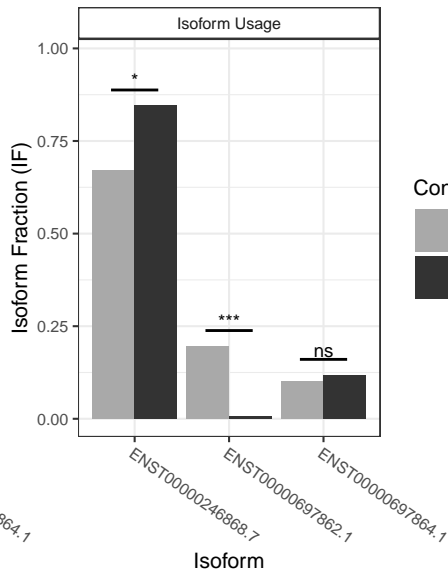
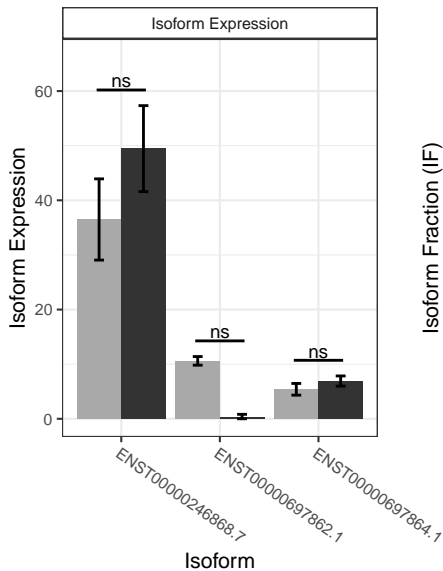
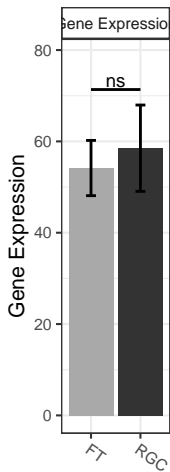
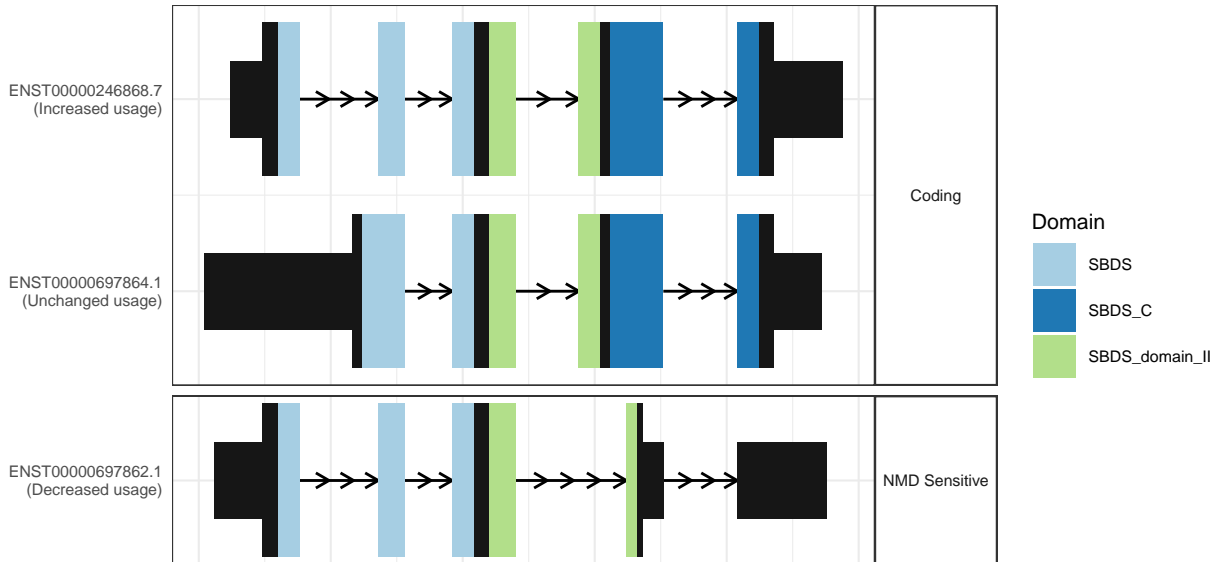
Domain



Band_7

Flot

The isoform switch in SBDS (FT vs RGC)



Domain



SBDS

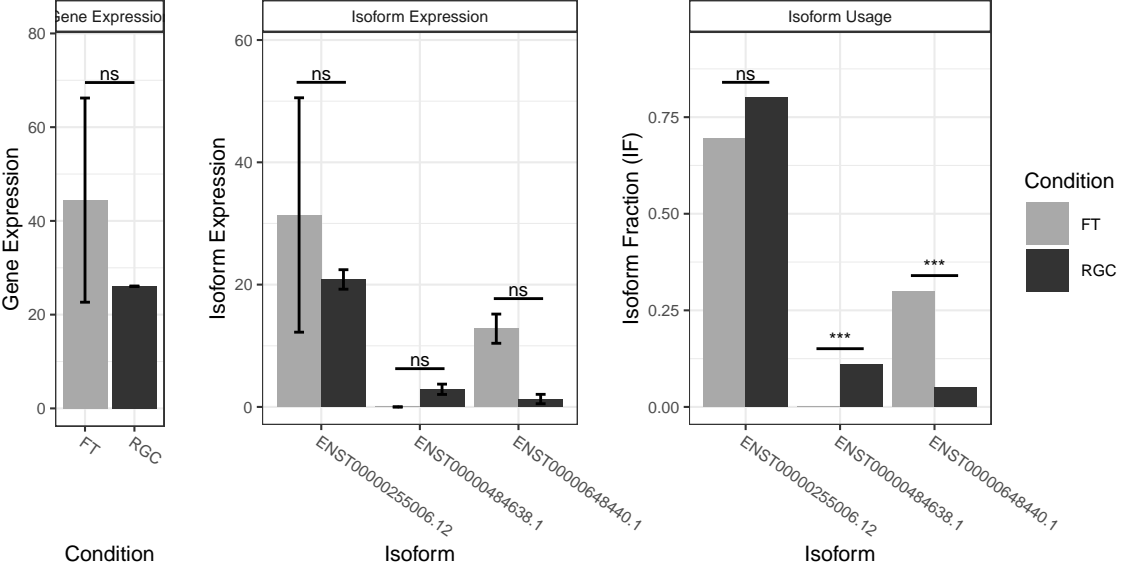
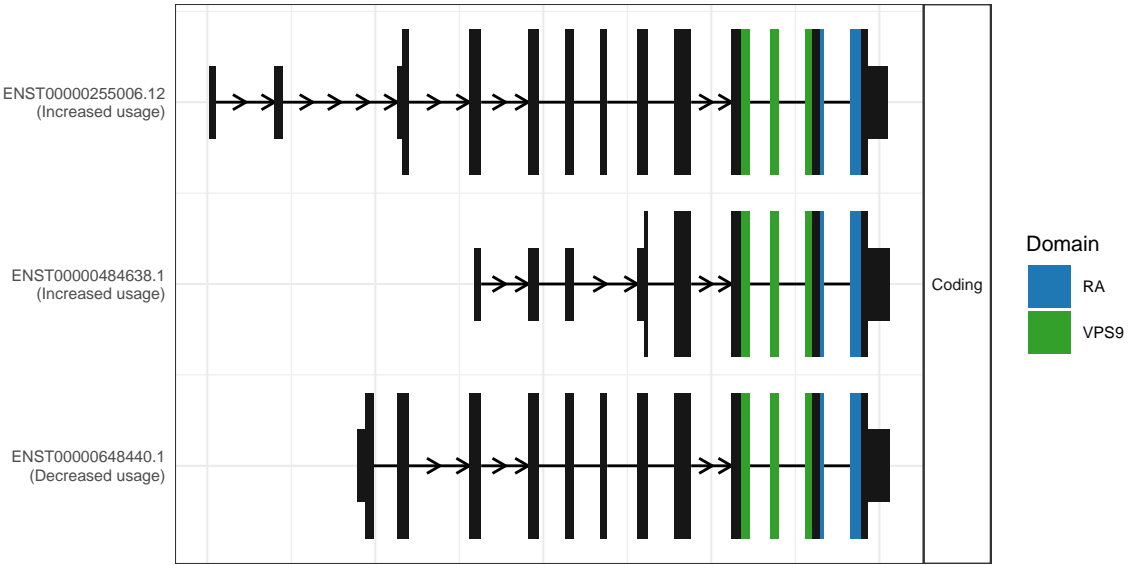


SBDS_C



SBDS_domain_II

The isoform switch in RIN2 (FT vs RGC)



Domain

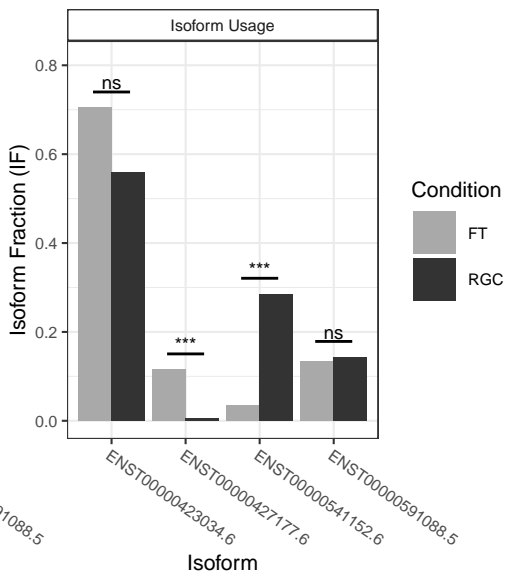
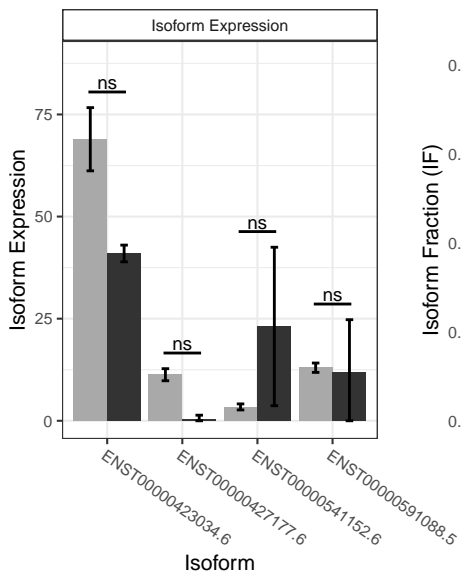
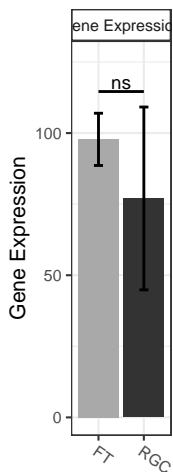
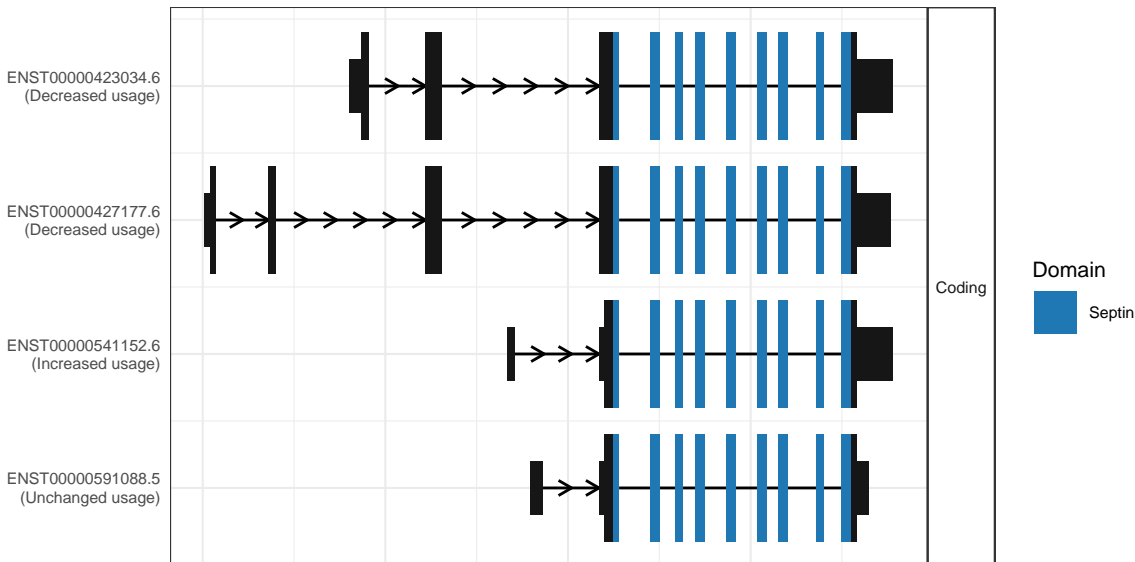


RA



VPS9

The isoform switch in SEPTIN9 (FT vs RGC)

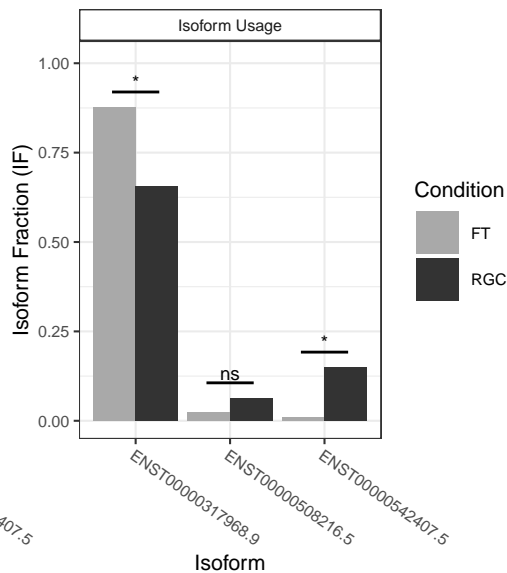
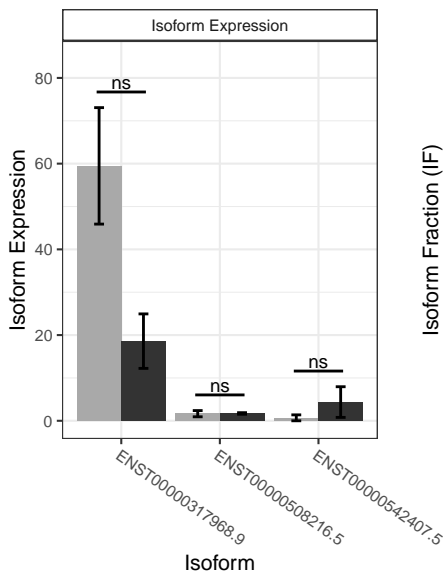
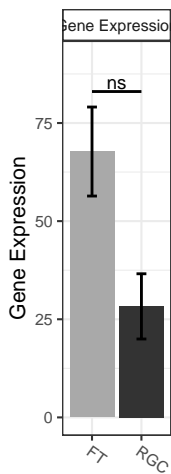
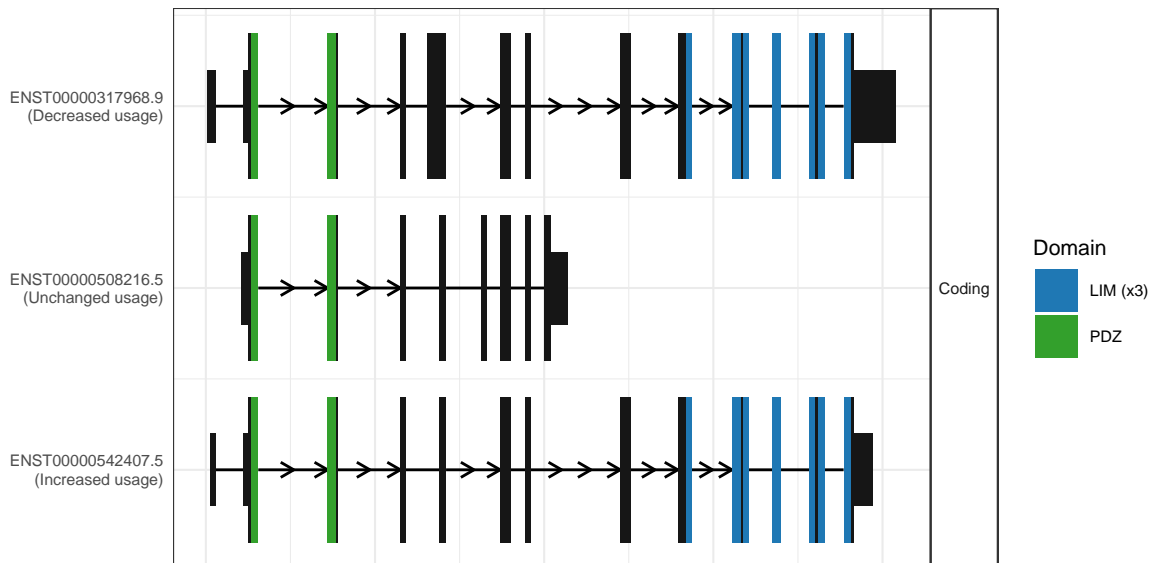


Domain



Septin

The isoform switch in PDLIM5 (FT vs RGC)



Domain

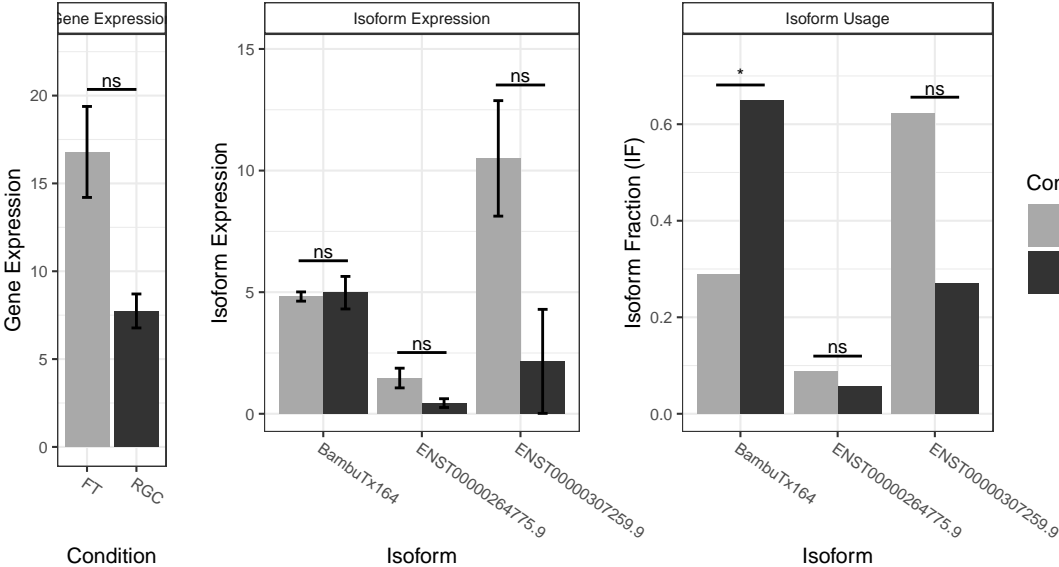
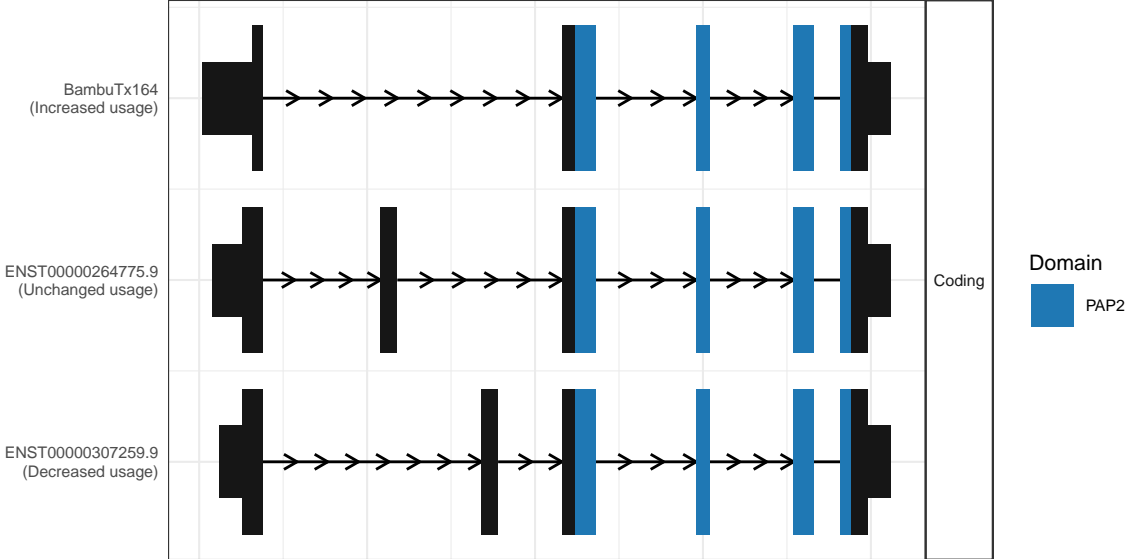


LIM (x3)



PDZ

The isoform switch in PLPP1 (FT vs RGC)

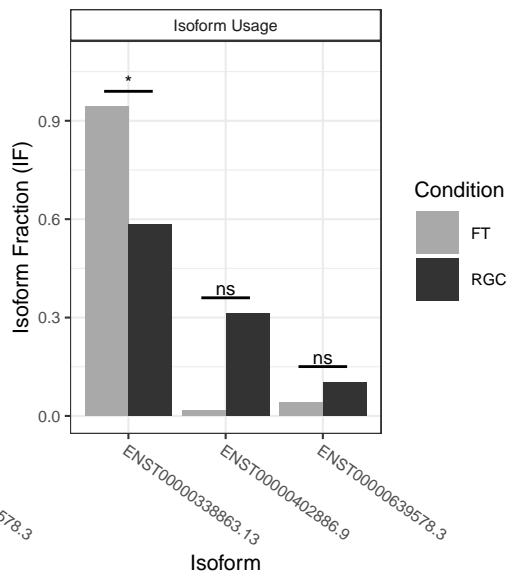
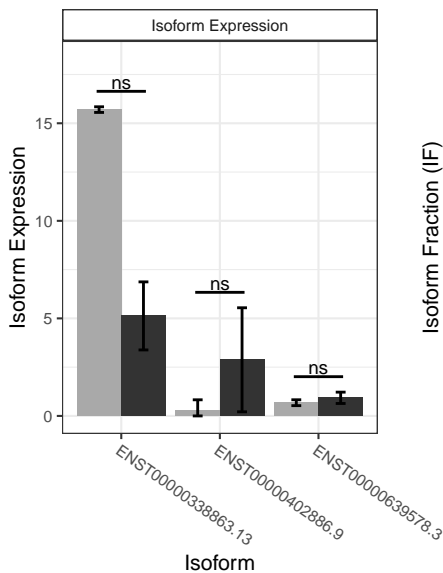
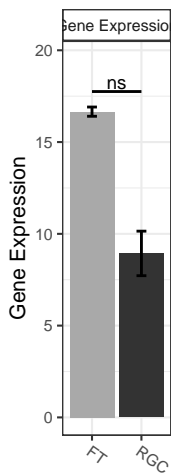
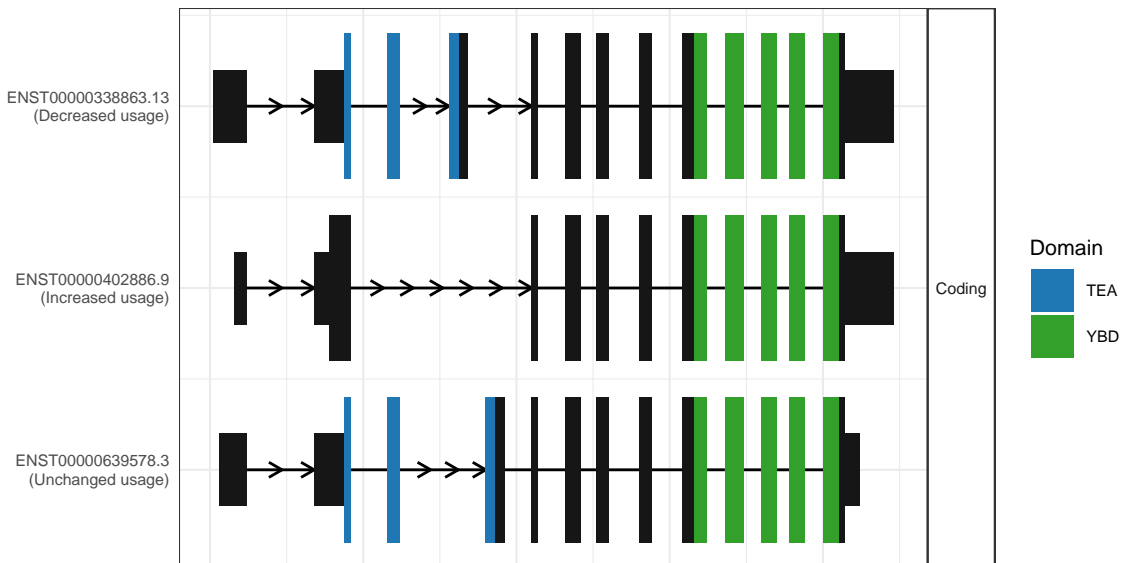


Domain



PAP2

The isoform switch in TEAD3 (FT vs RGC)



Domain

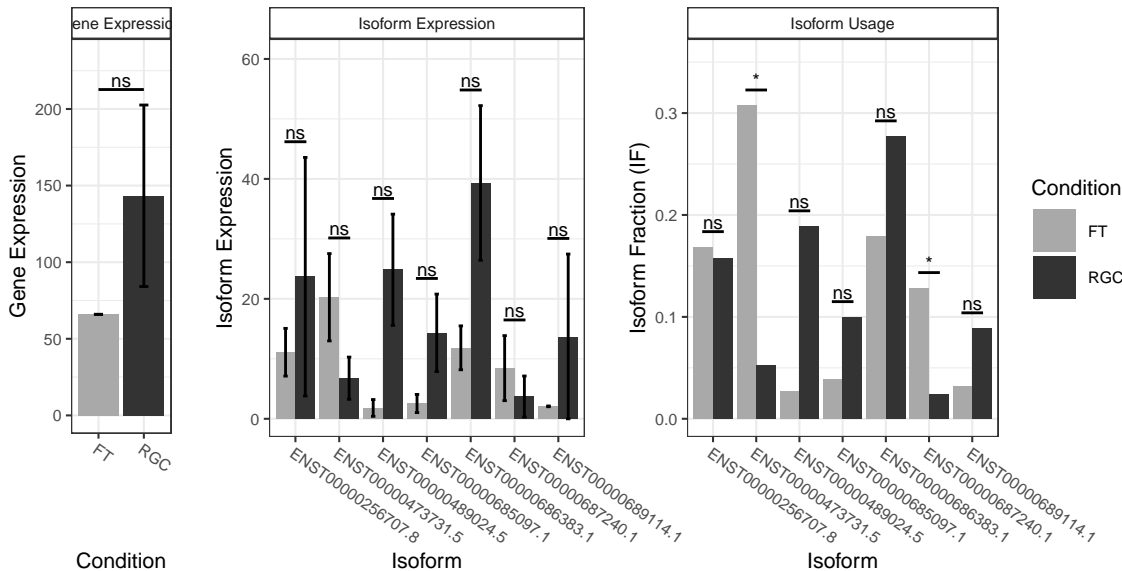
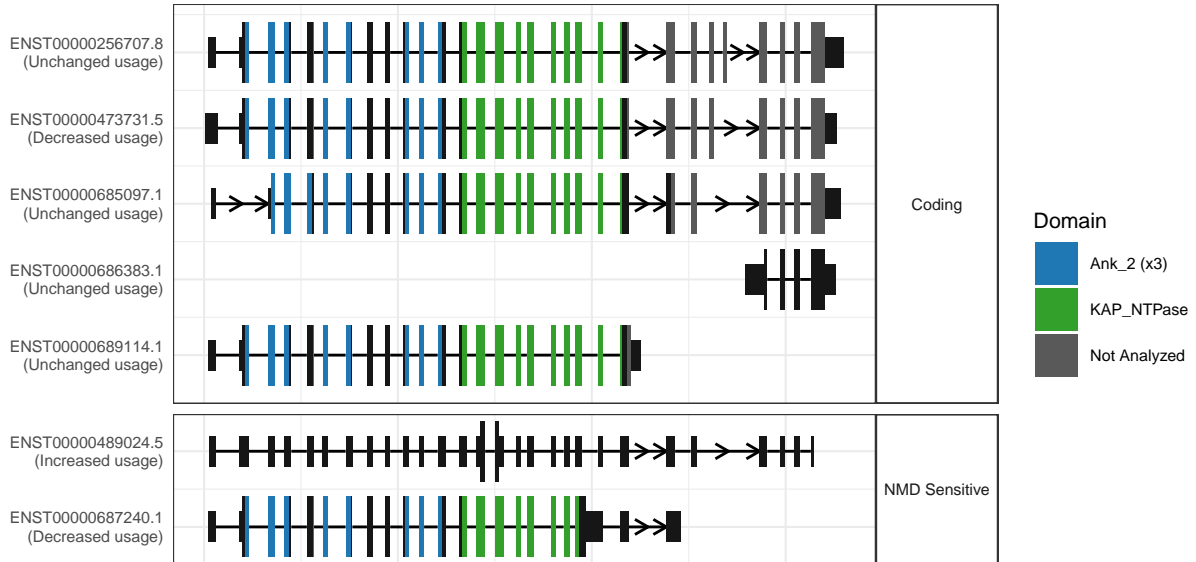


TEA






YBD

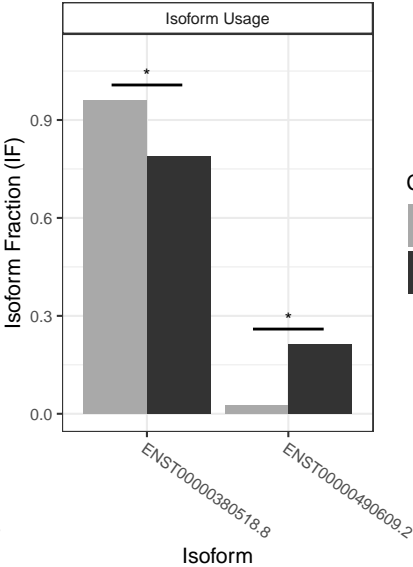
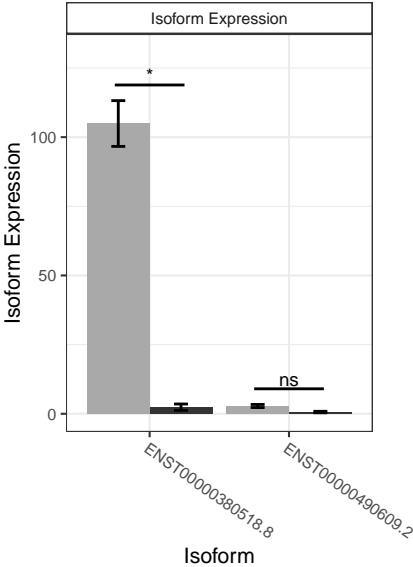
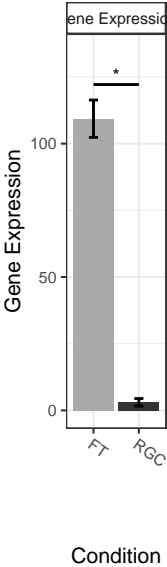
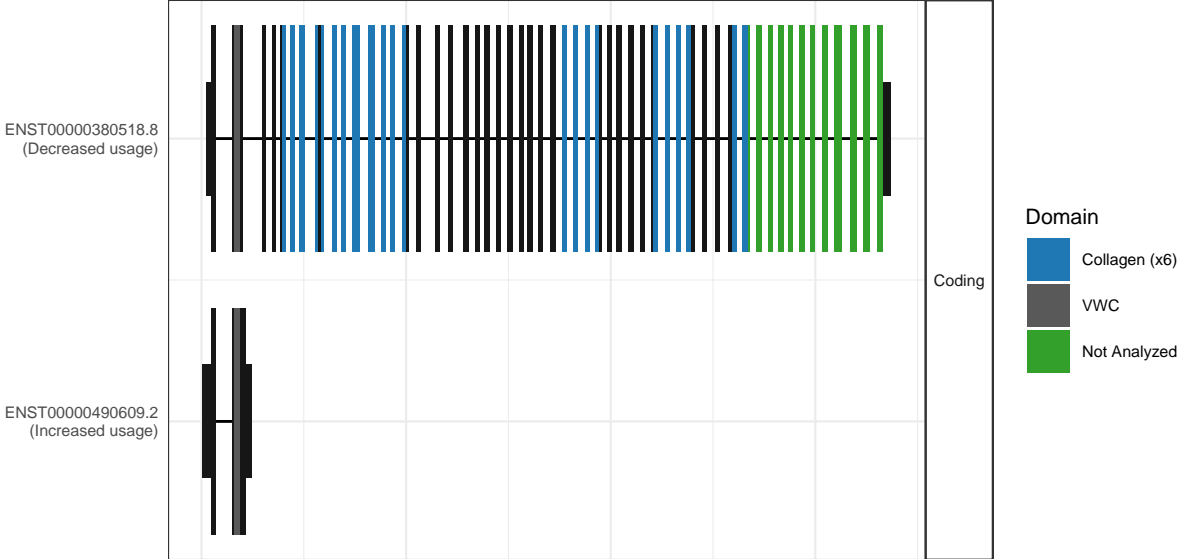
The isoform switch in KIDINS220 (FT vs RGC)



Domain

	Ank_2 (x3)
	KAP_NTPase
	Not Analyzed

The isoform switch in COL2A1 (FT vs RGC)



Domain



Collagen (x6)

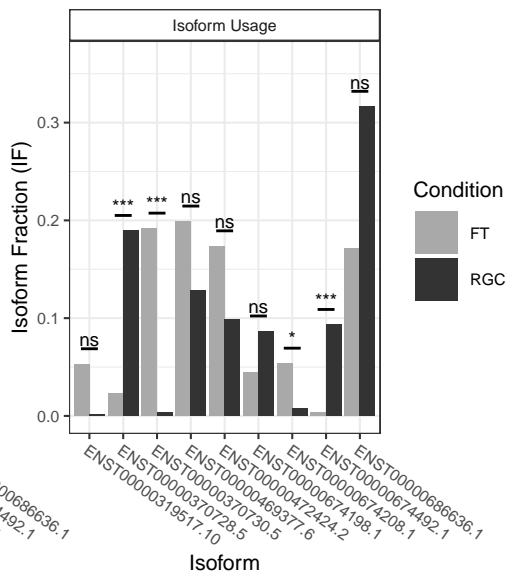
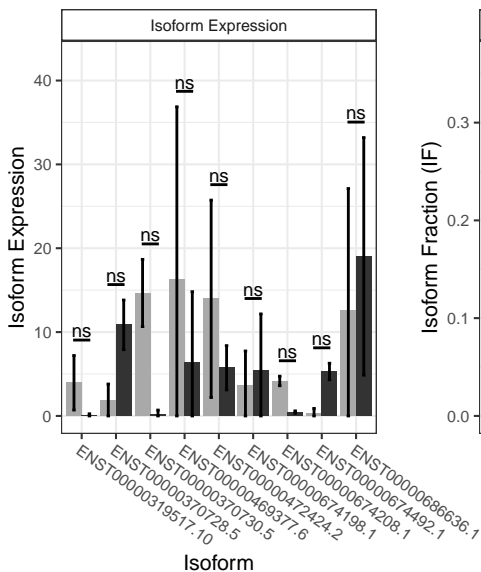
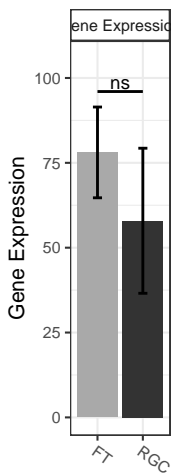
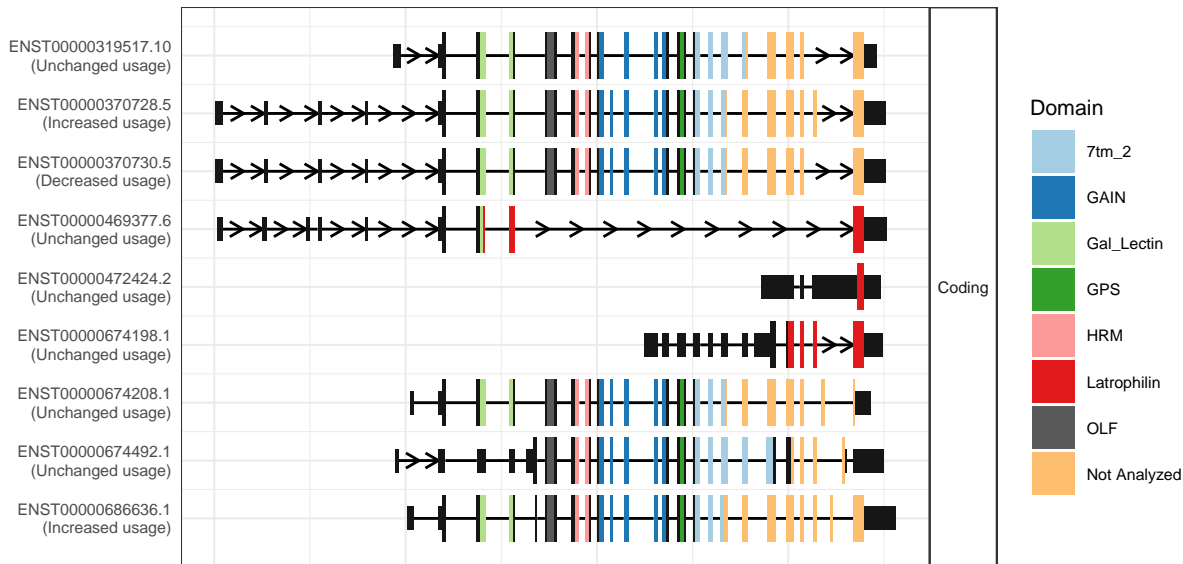


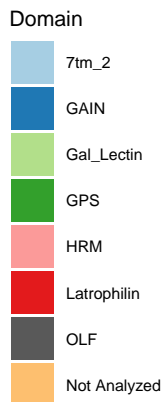
VWC



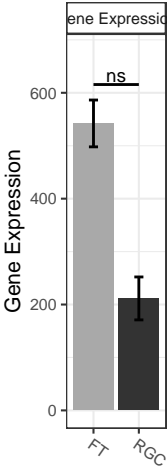
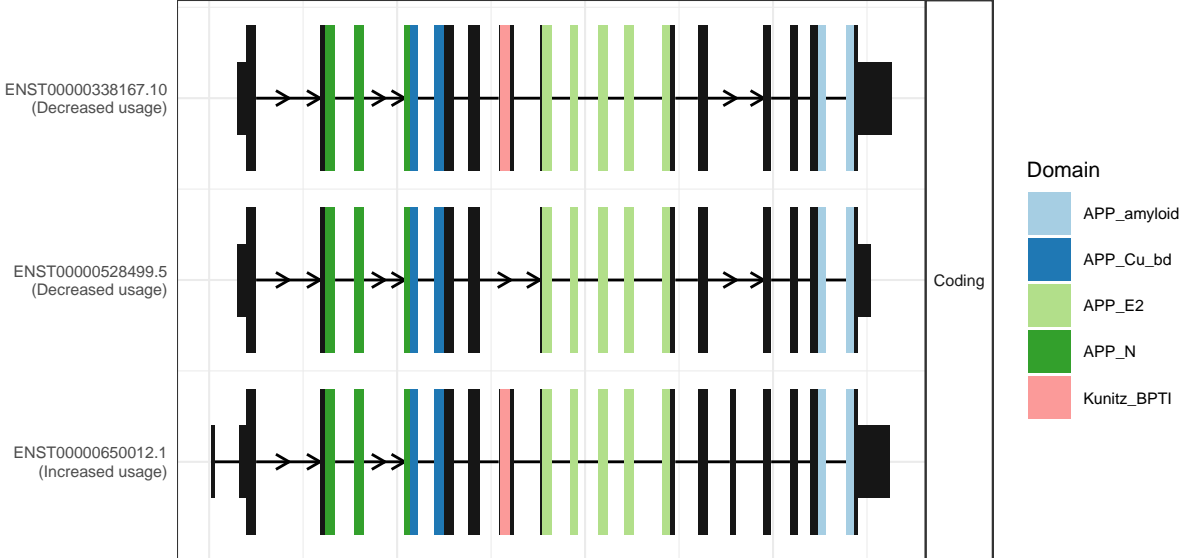
Not Analyzed

The isoform switch in ADGRL2 (FT vs RGC)

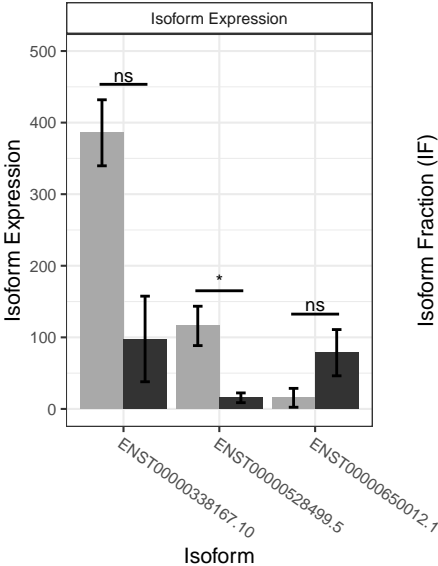




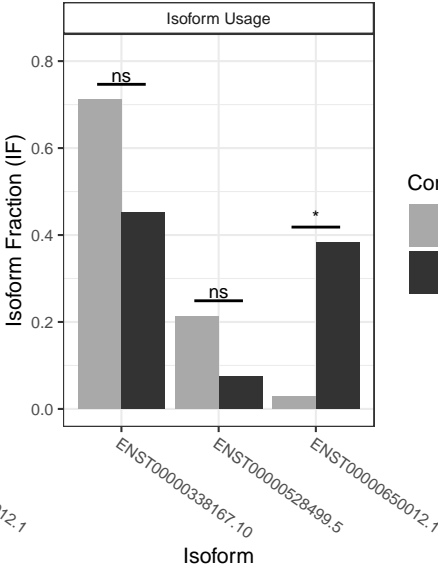
The isoform switch in APLP2 (FT vs RGC)



Condition



Isoform








Isoform

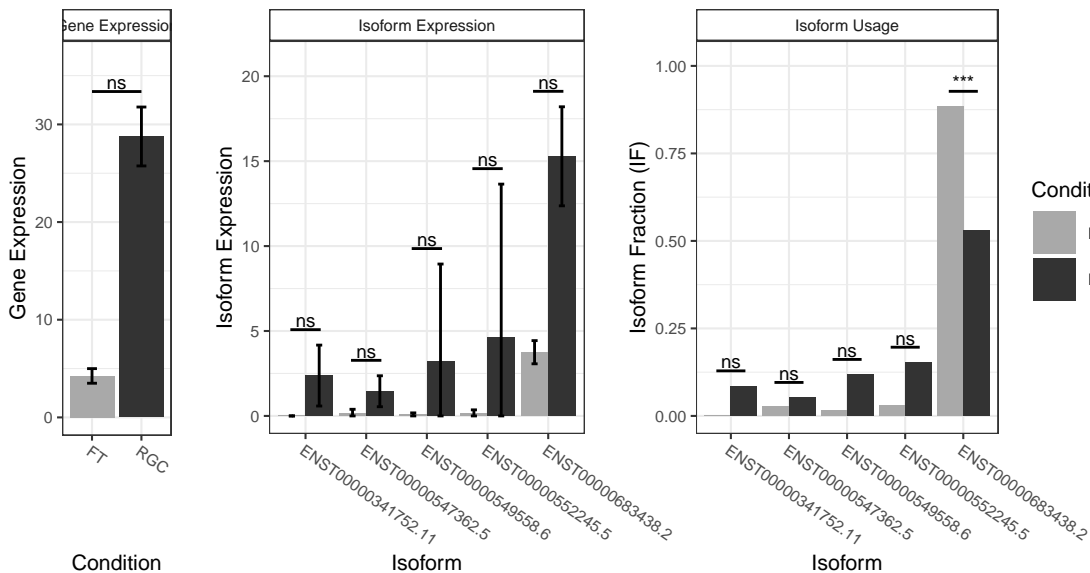
Condition



Domain

	APP_amyloid
	APP_Cu_bd
	APP_E2
	APP_N
	Kunitz_BPTI

The diagram illustrates a protein-coding gene structure. It features a horizontal line representing the DNA sequence, with vertical bars indicating exons and introns. The gene is divided into several segments: a long initial segment with a black bar at the start, followed by a series of blue bars (exons) and black bars (introns). The final segment is a long black bar. The gene is flanked by two vertical lines, and the word "Coding" is written to the right of the final segment.



Domain



Ank_2 (x2)



PID

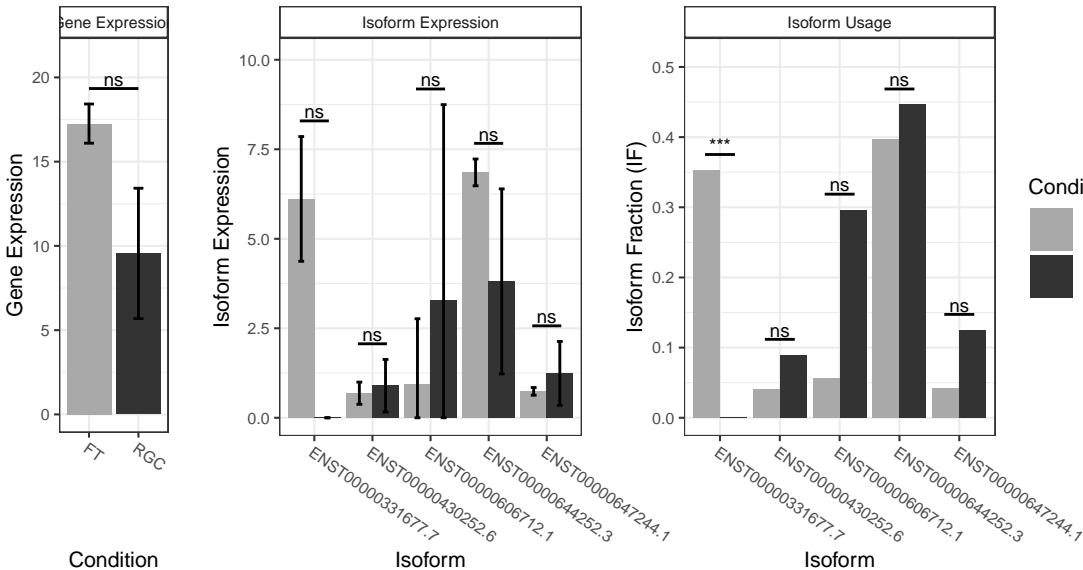
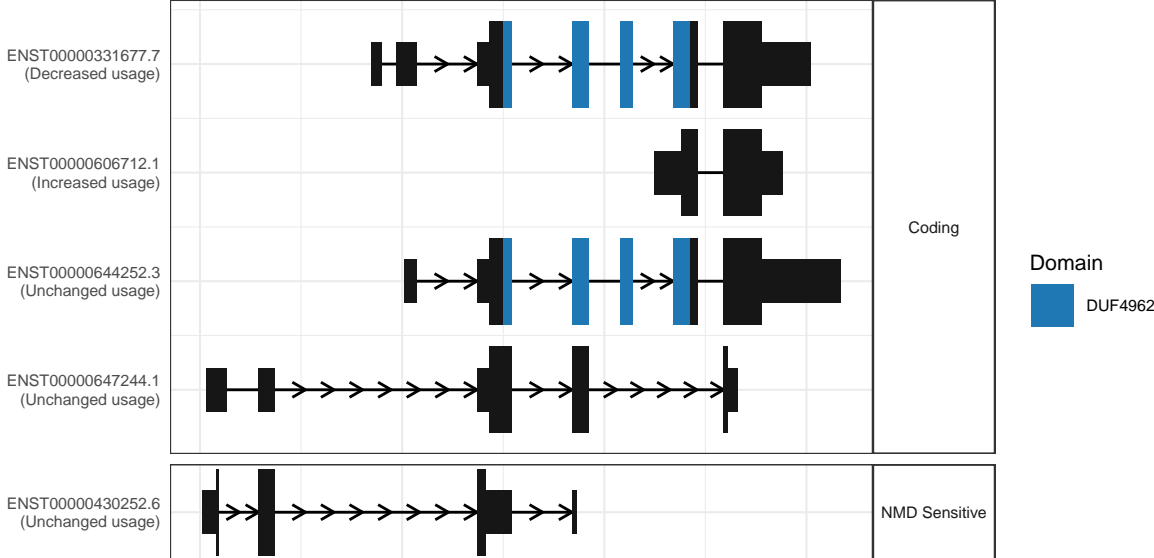


SAM_1 (x2)



Not Analyzed

The isoform switch in DSE (FT vs RGC)

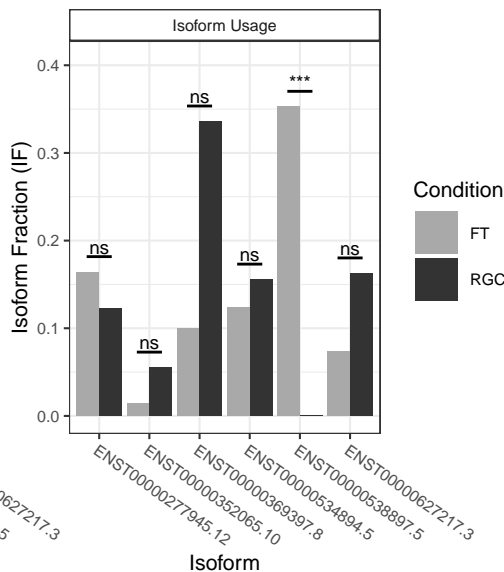
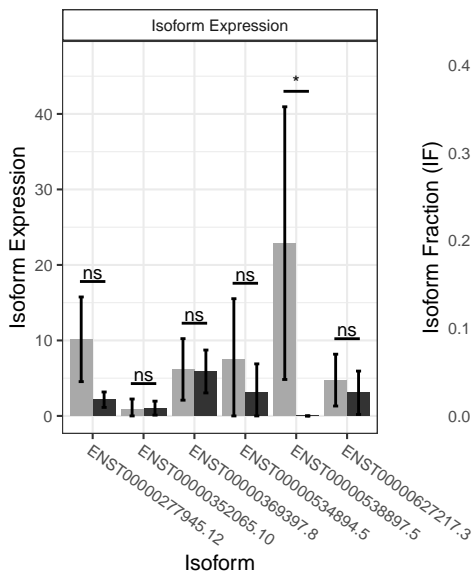
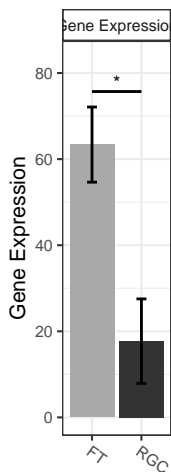
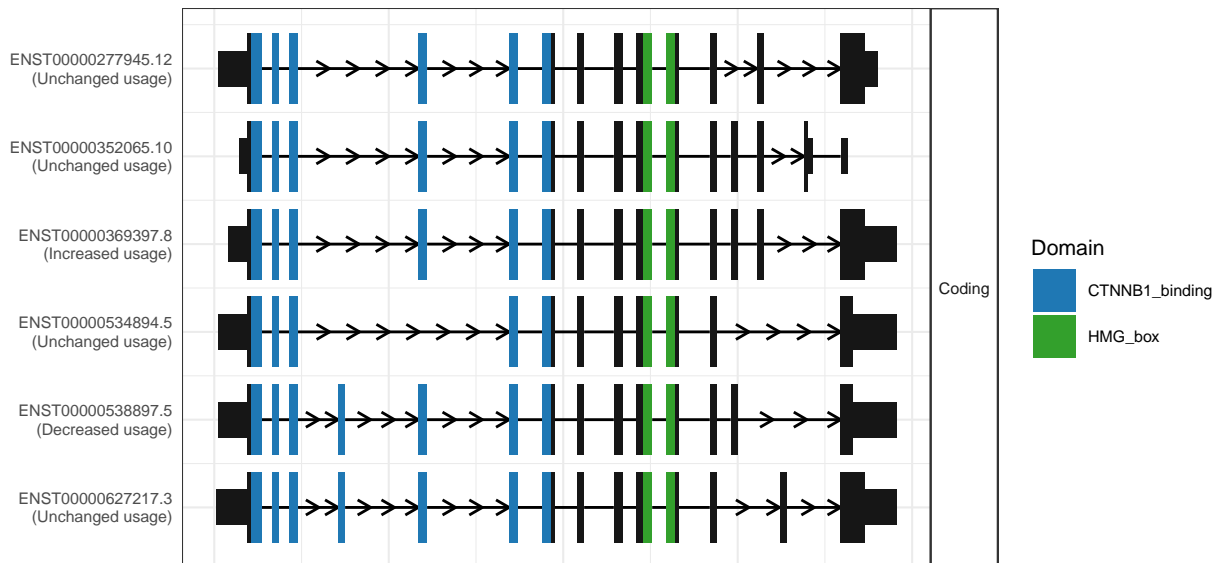


Domain



DUF4962

The isoform switch in TCF7L2 (FT vs RGC)



Condition

Isoform

Isoform

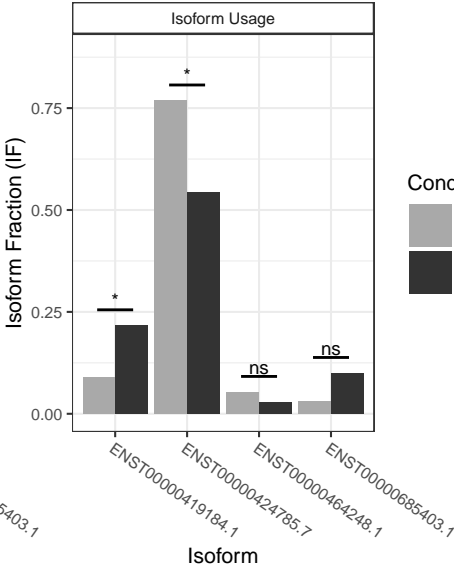
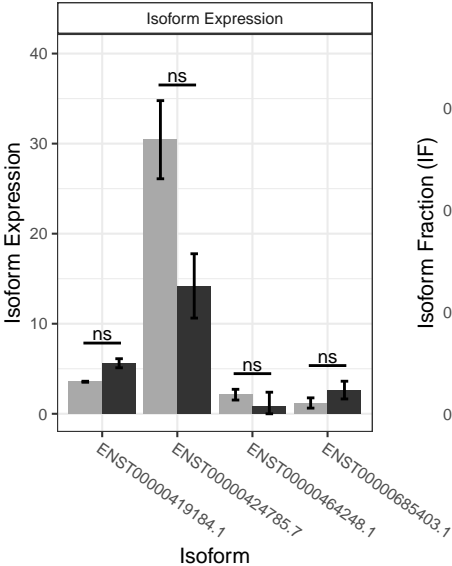
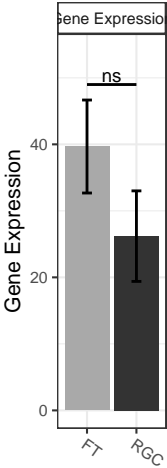
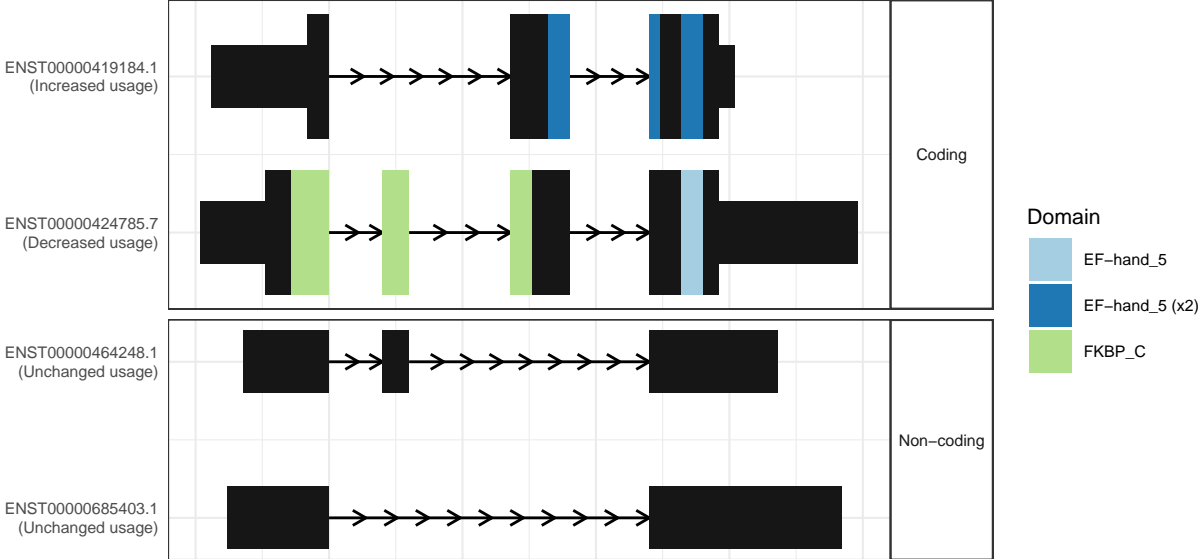
Domain






CTNNB1_binding

HMG_box

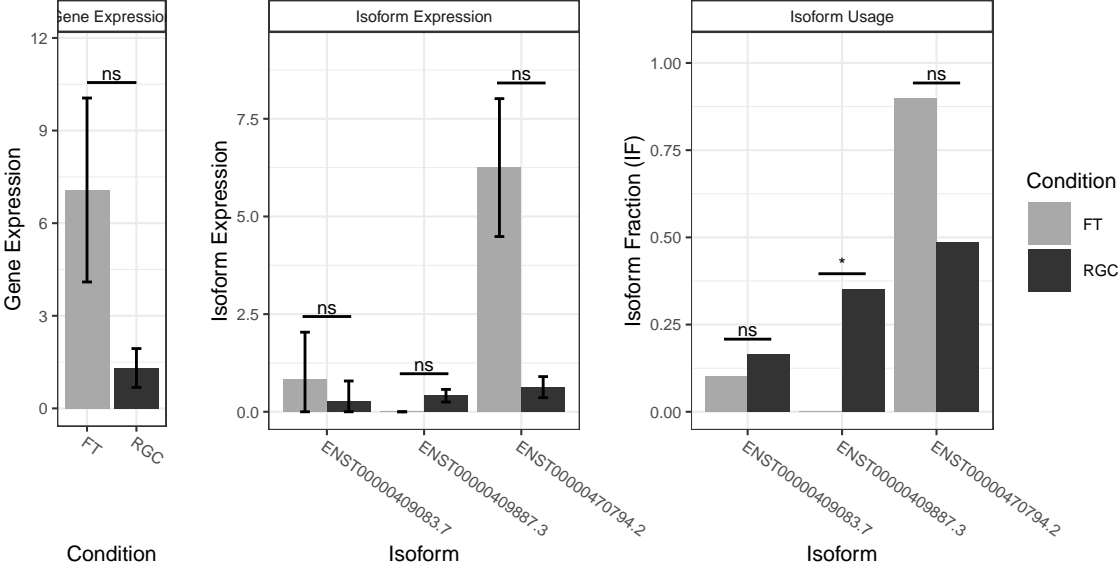
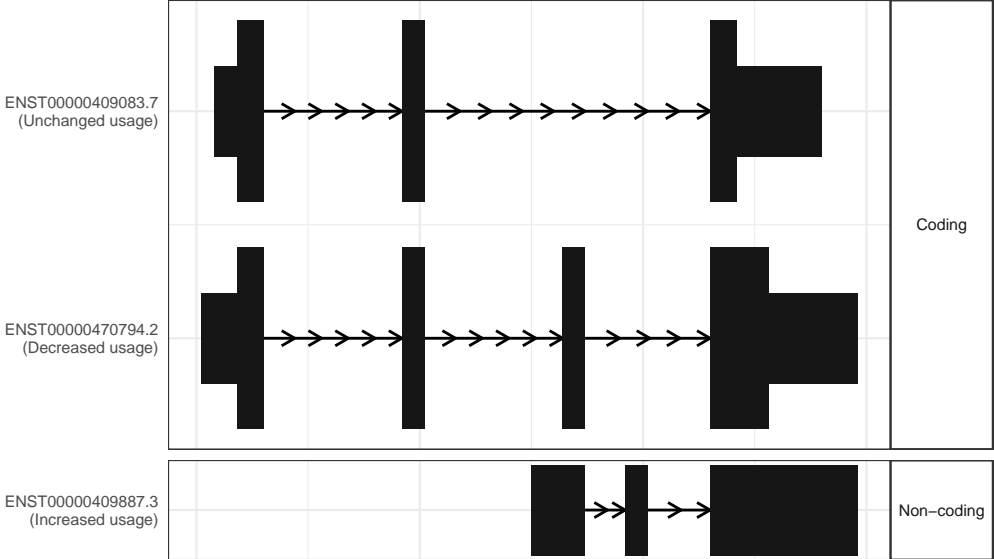
The isoform switch in FKBP7 (FT vs RGC)



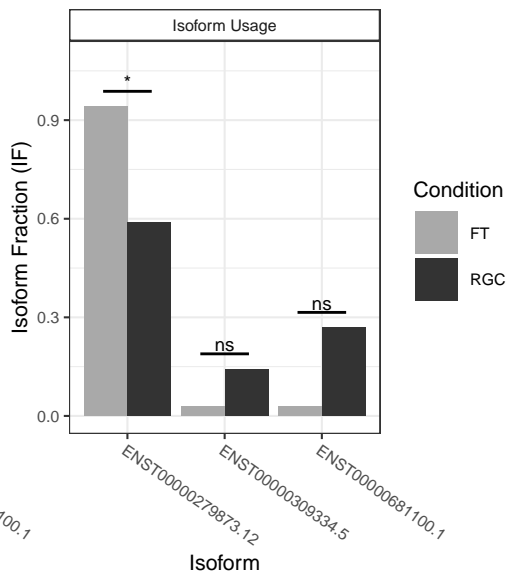
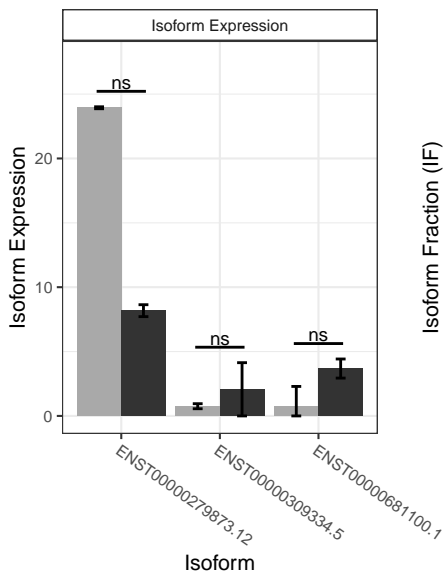
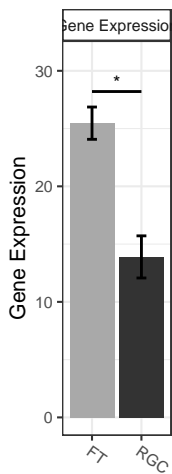
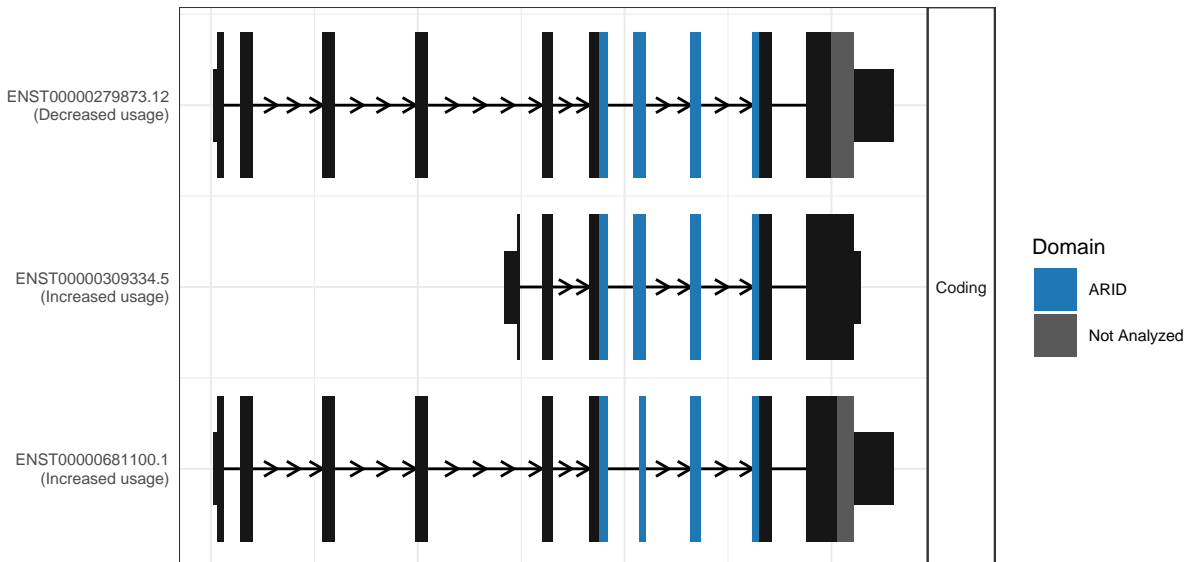
Domain

	EF-hand_5
	EF-hand_5 (x2)
	FKBP_C

The isoform switch in LRRC75A (FT vs RGC)



The isoform switch in ARID5B (FT vs RGC)



Domain

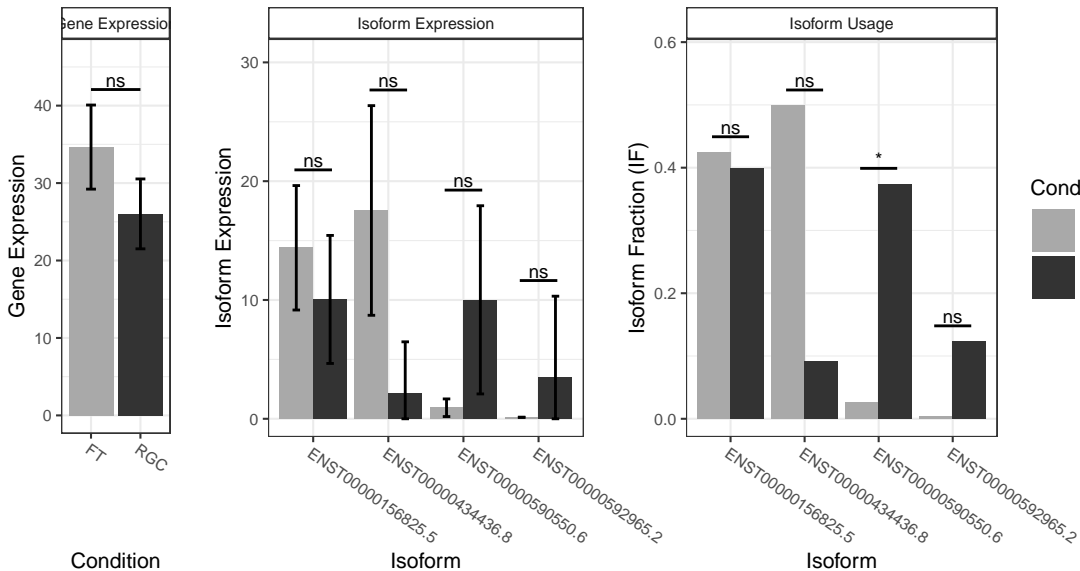
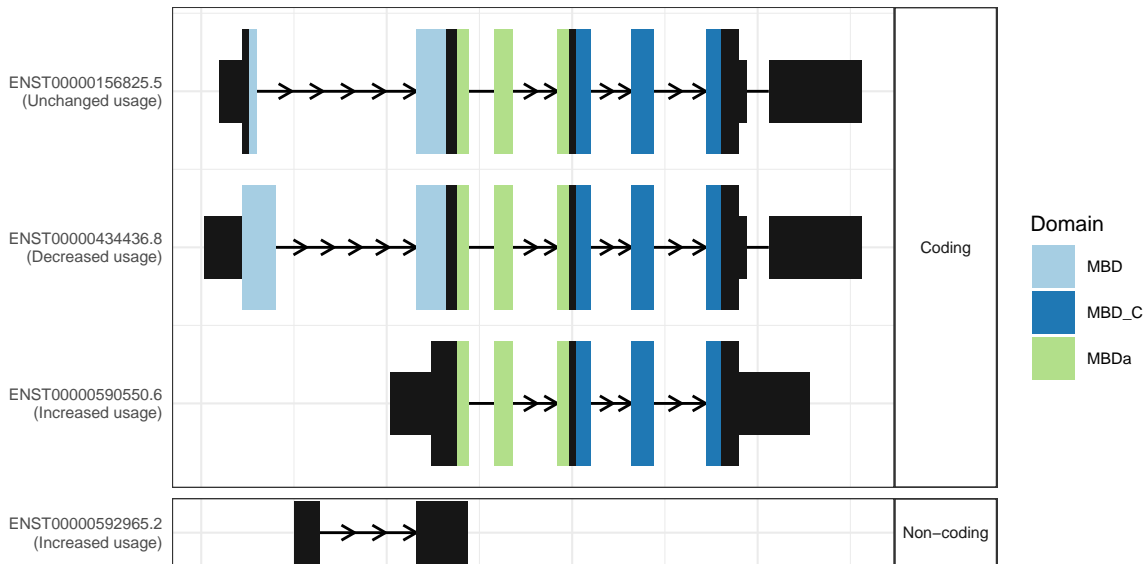


ARID



Not Analyzed

The isoform switch in MBD3 (FT vs RGC)



Domain



MBD

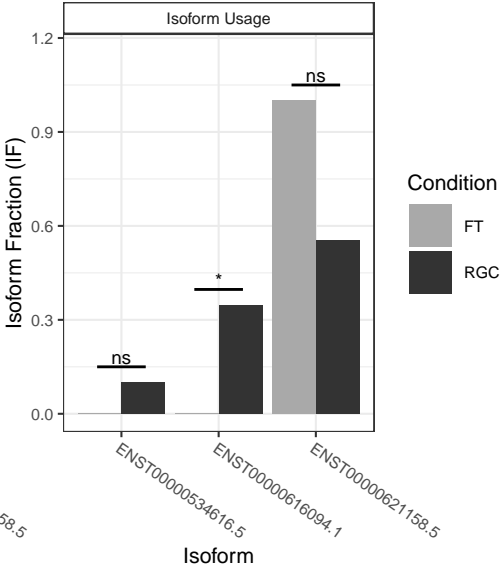
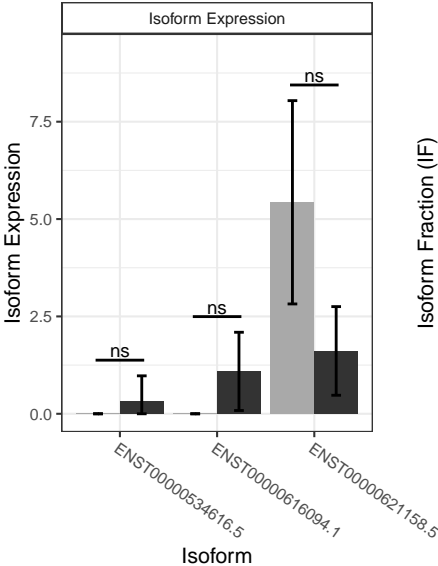
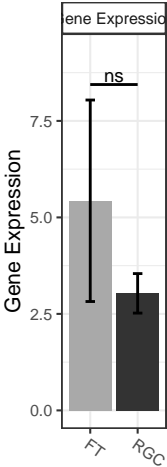
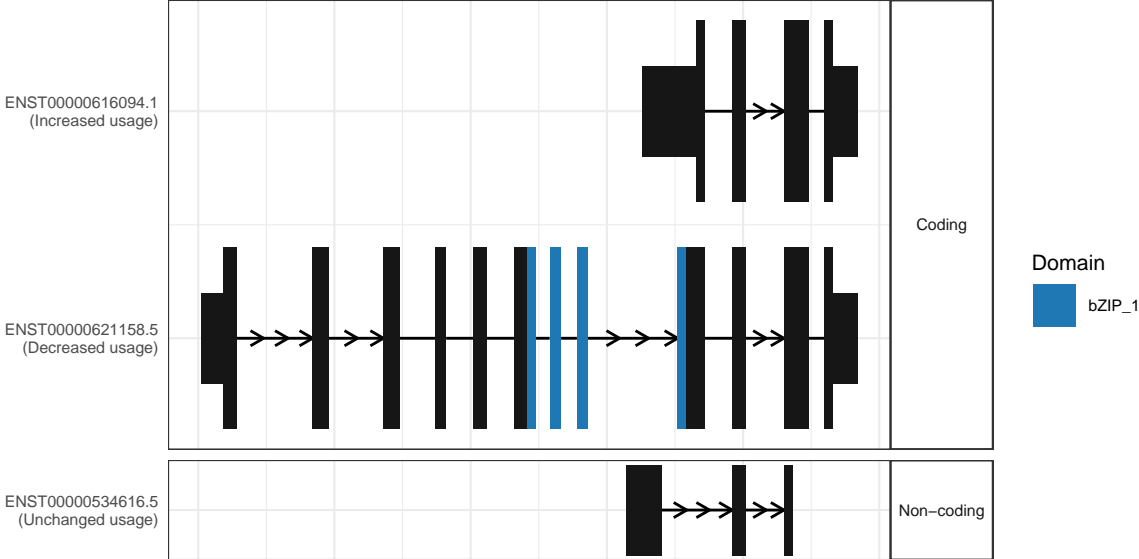


MBD_C



MBDa

The isoform switch in CREB3L1 (FT vs RGC)

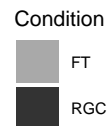
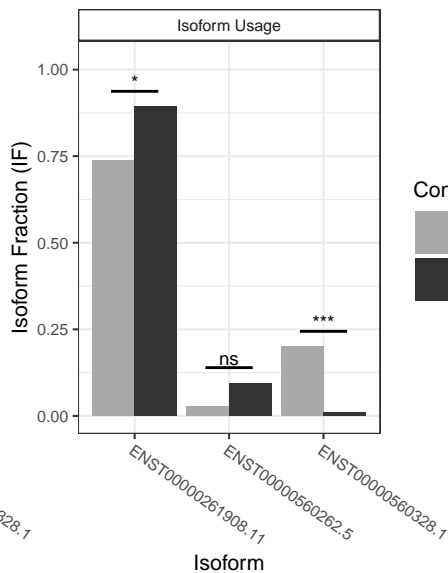
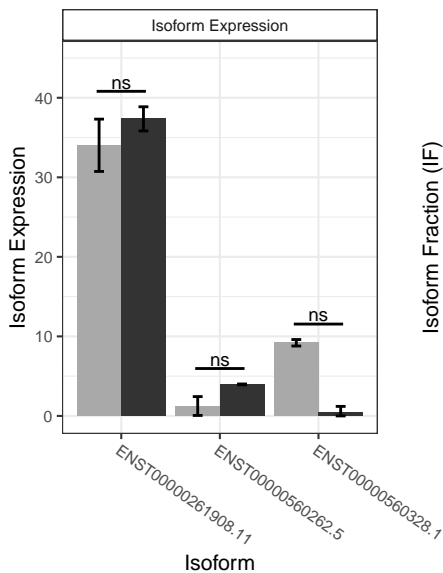
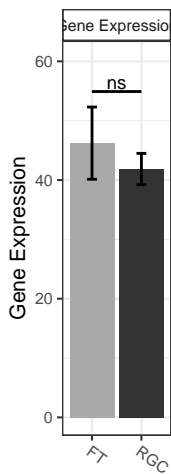
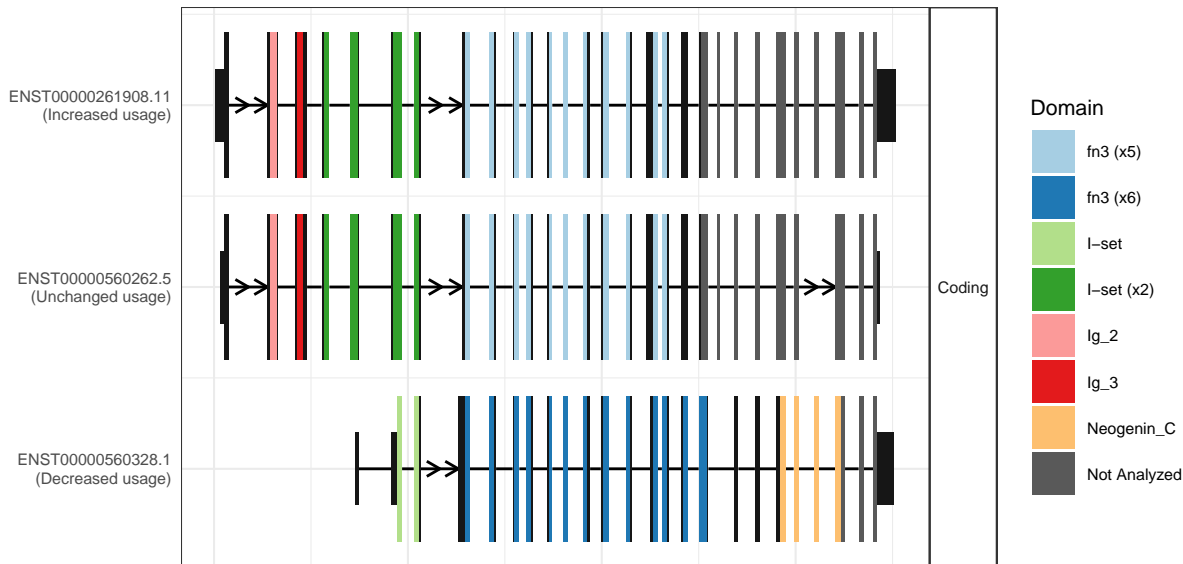


Domain











bZIP_1

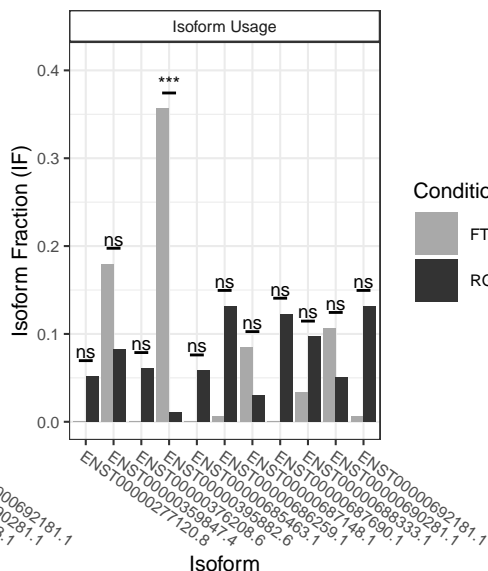
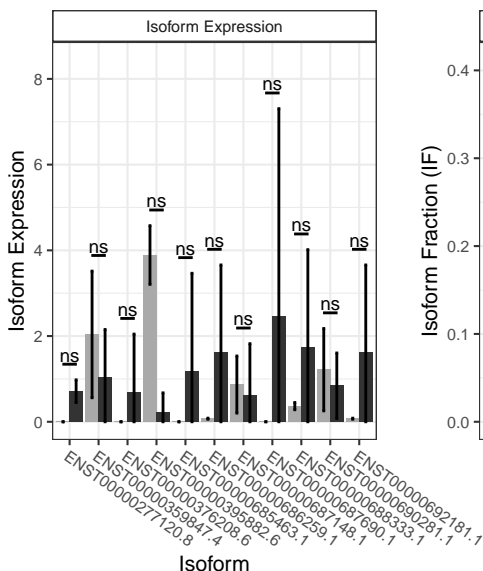
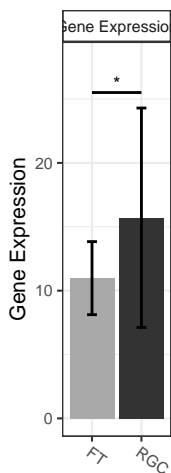
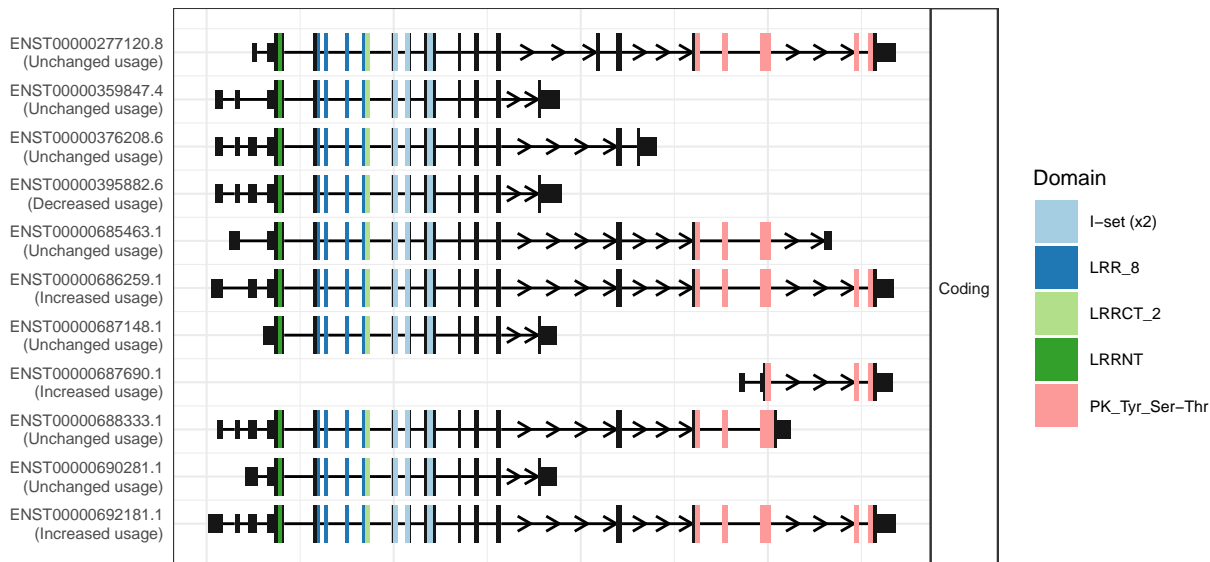
The isoform switch in NEO1 (FT vs RGC)



Domain

	fn3 (x5)
	fn3 (x6)
	I-set
	I-set (x2)
	Ig_2
	Ig_3
	Neogenin_C
	Not Analyzed

The isoform switch in NTRK2 (FT vs RGC)








Condition

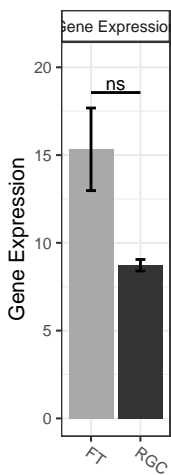
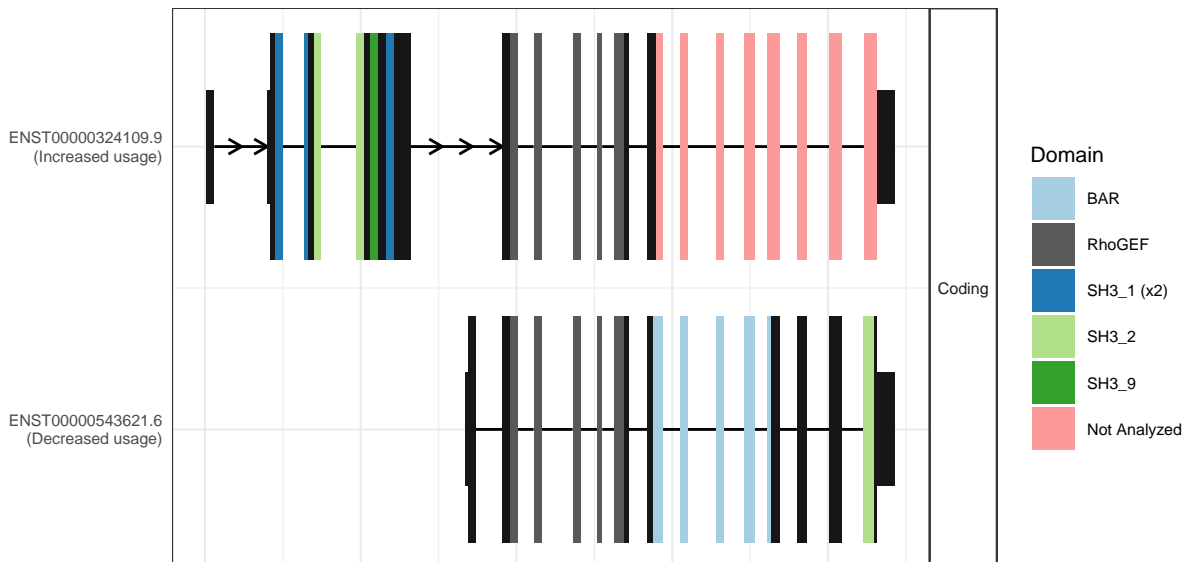
Isoform

Isoform

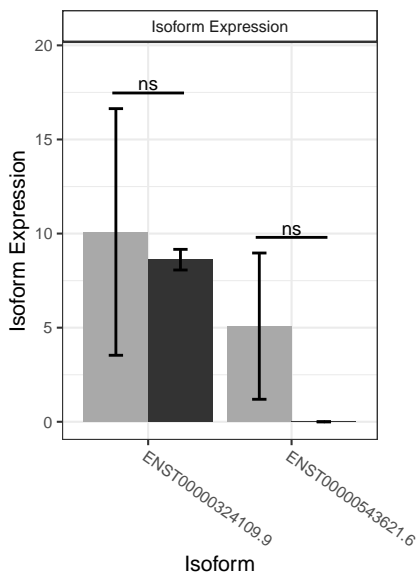
Domain

	I-set (x2)
	LRR_8
	LRRCT_2
	LRRNT
	PK_Tyr_Ser-Thr

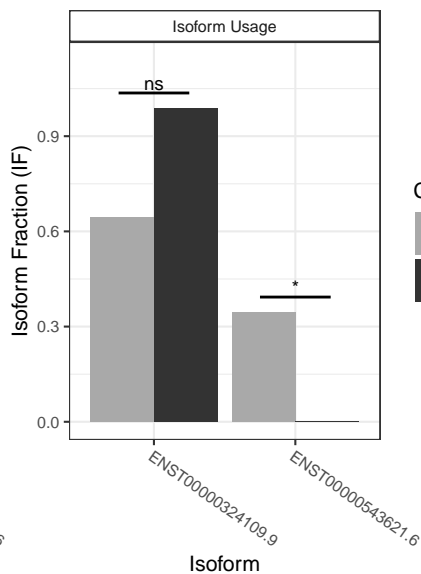
The isoform switch in DNMBP (FT vs RGC)



Condition









Isoform

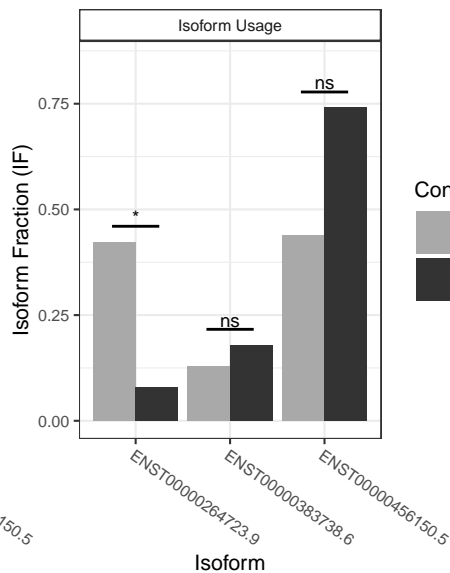
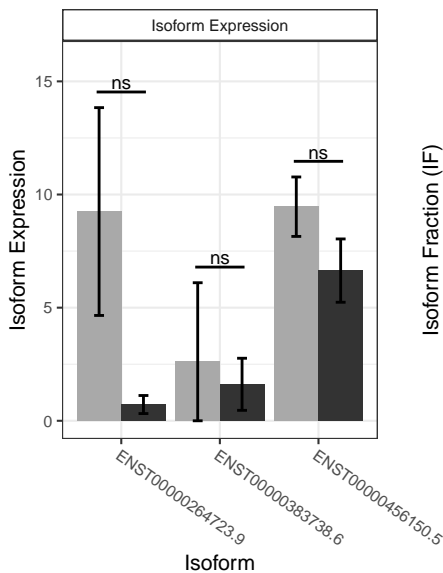
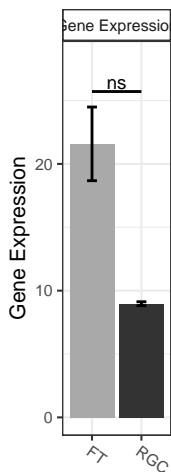
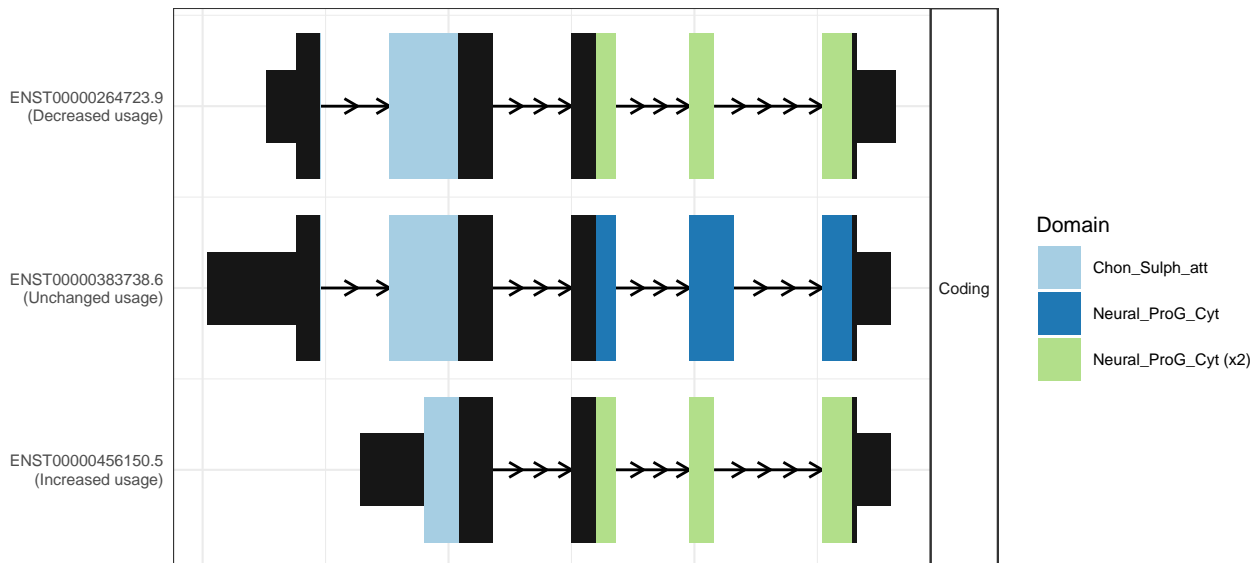


Isoform




Domain

	BAR
	RhoGEF
	SH3_1 (x2)
	SH3_2
	SH3_9
	Not Analyzed

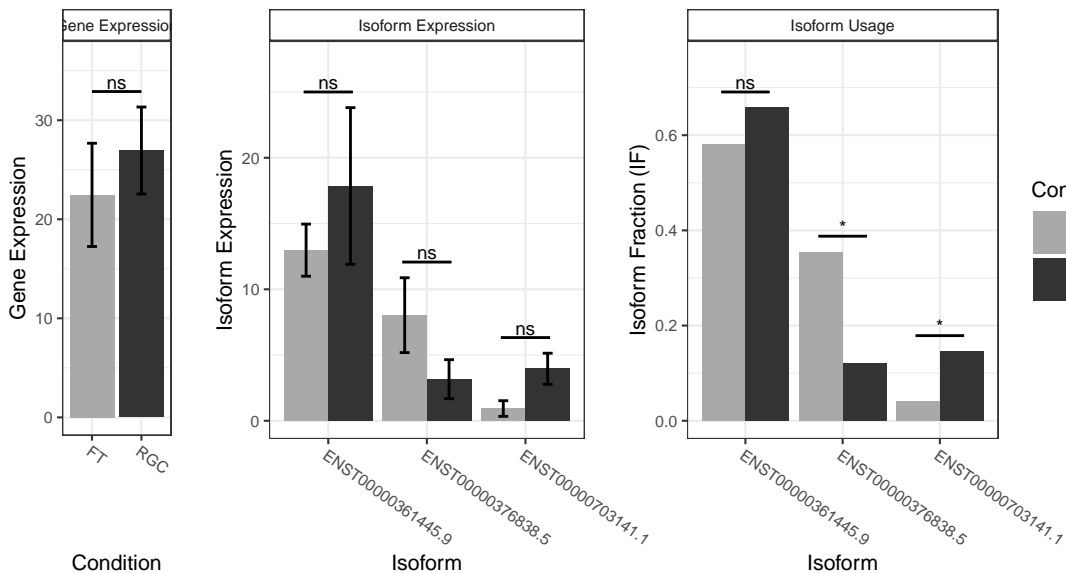
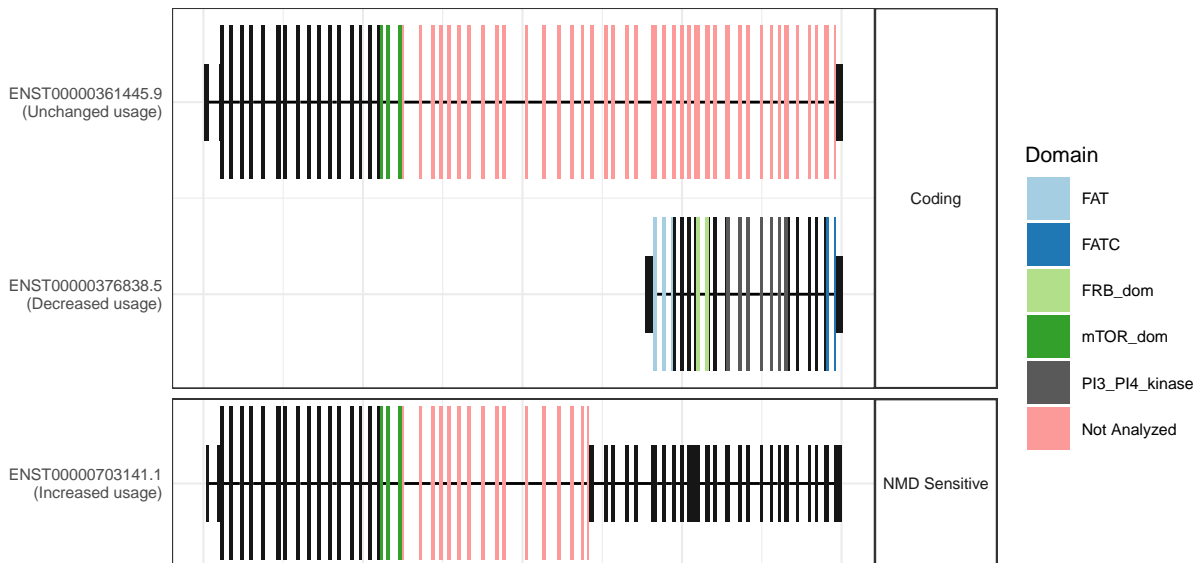
The isoform switch in CSPG5 (FT vs RGC)



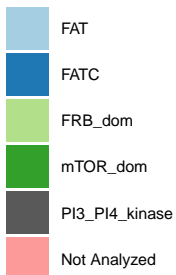
Domain

	Chon_Sulph_att
	Neural_ProG_Cyt
	Neural_ProG_Cyt (x2)

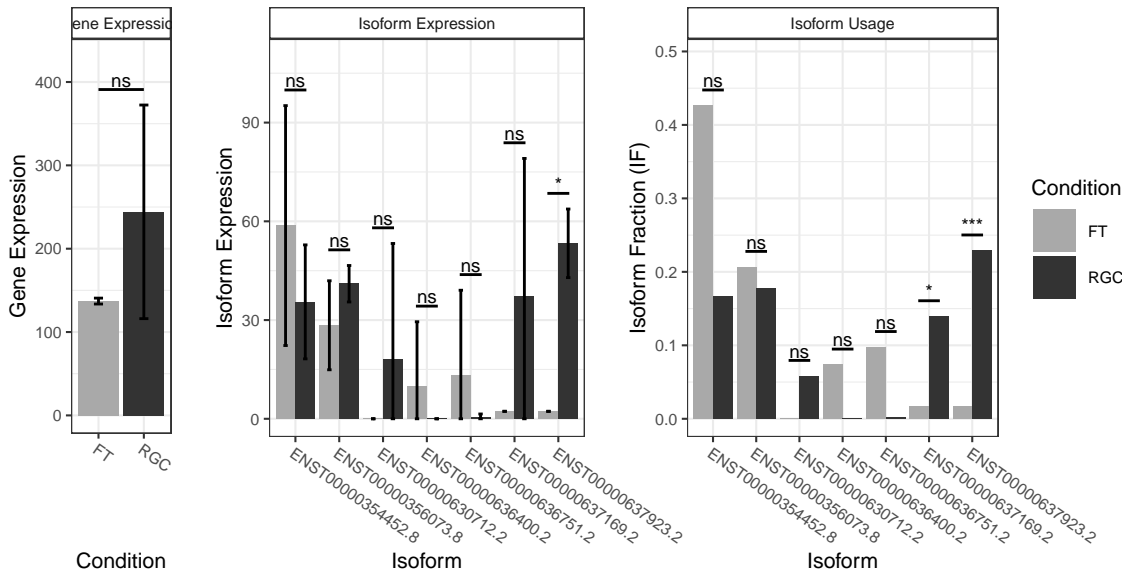
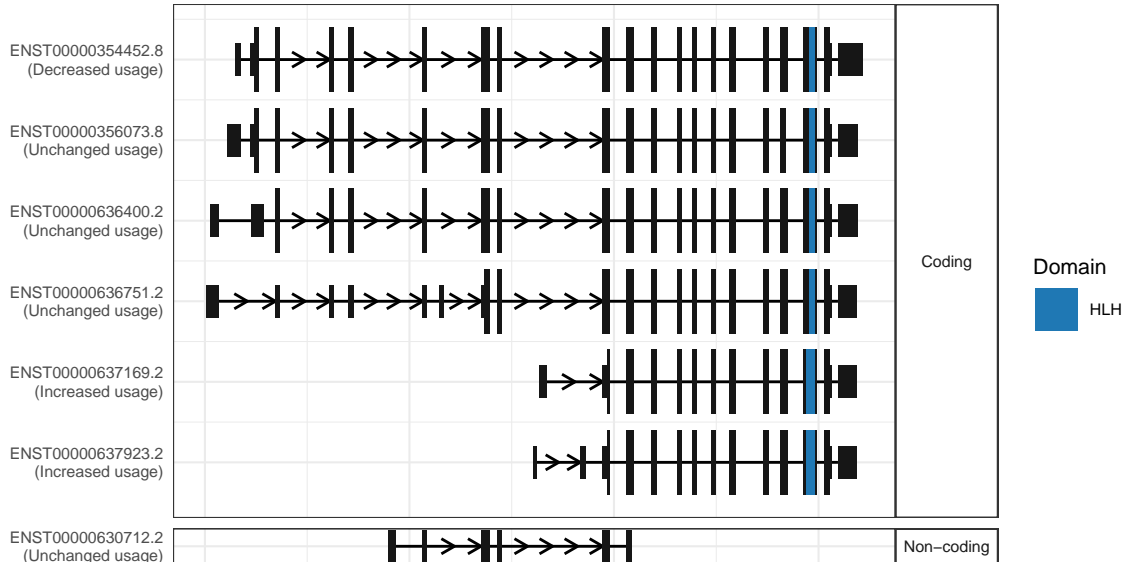
The isoform switch in MTOR (FT vs RGC)



Domain



The isoform switch in TCF4 (FT vs RGC)

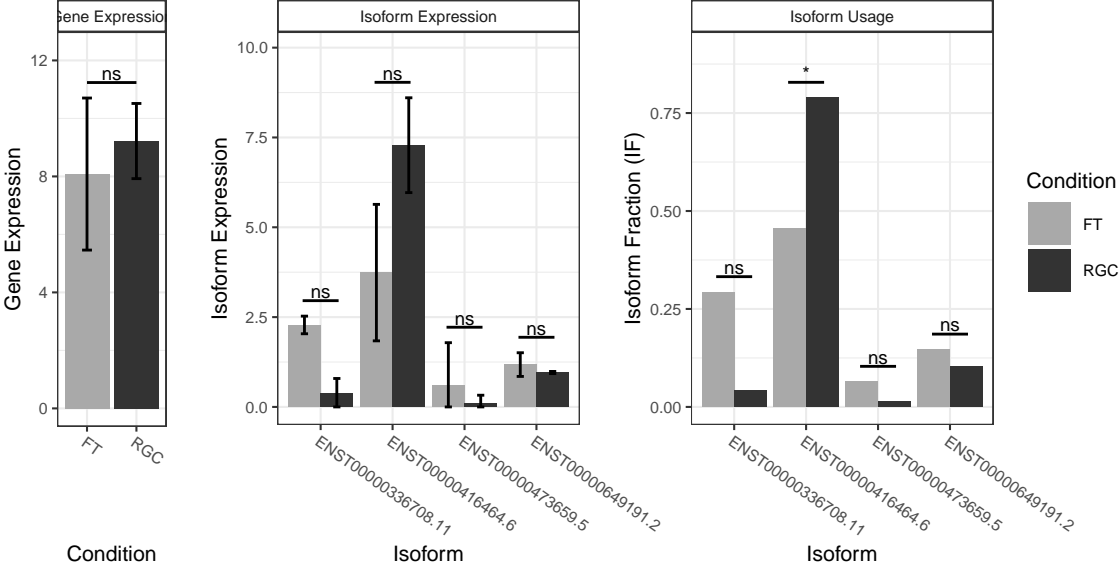
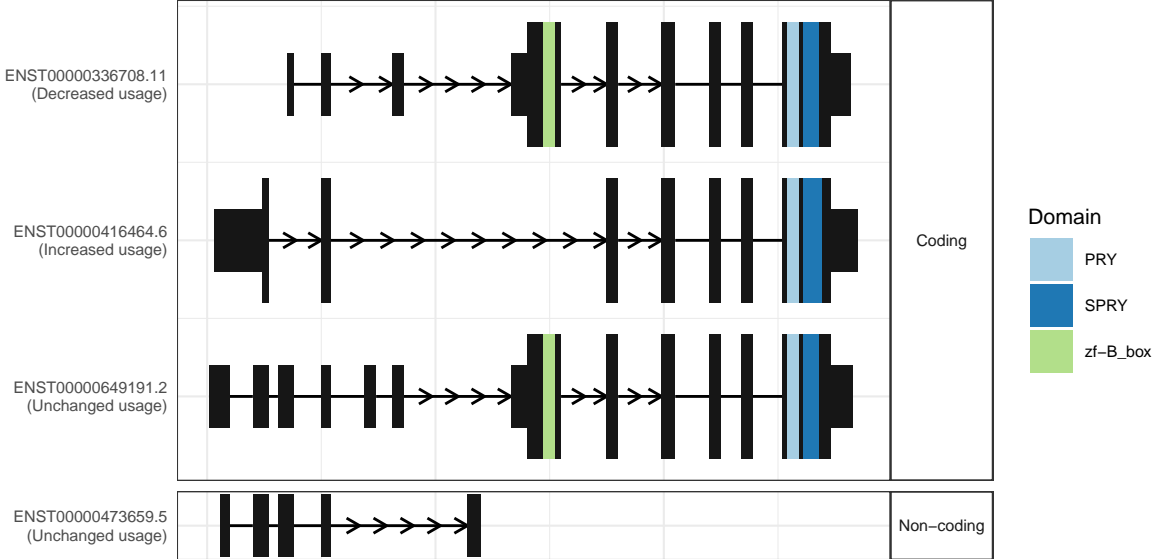


Domain



HLH

The isoform switch in TRIM16 (FT vs RGC)



Domain



PRY

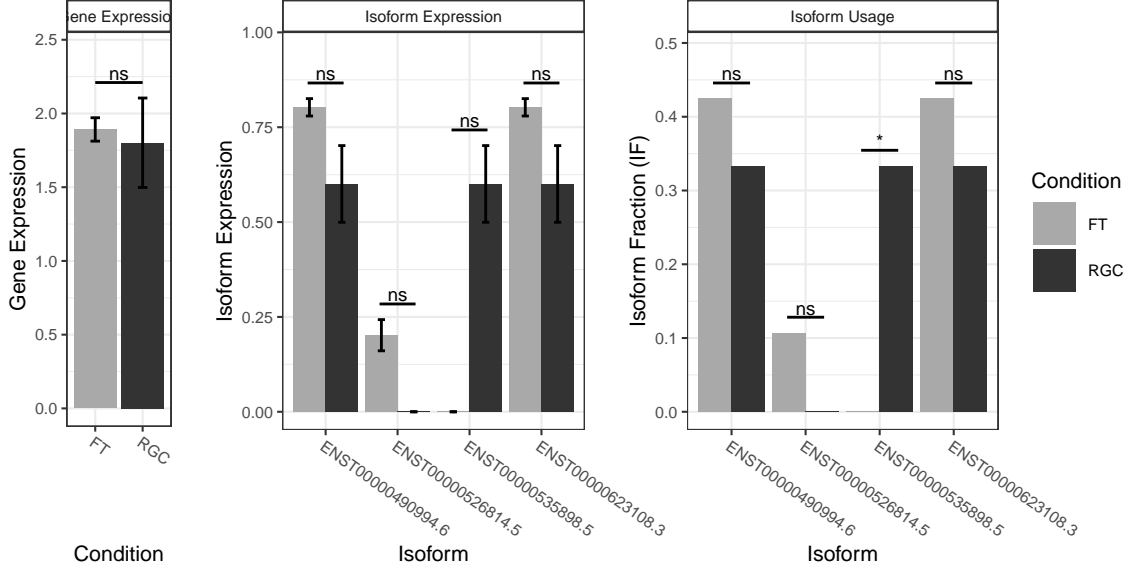
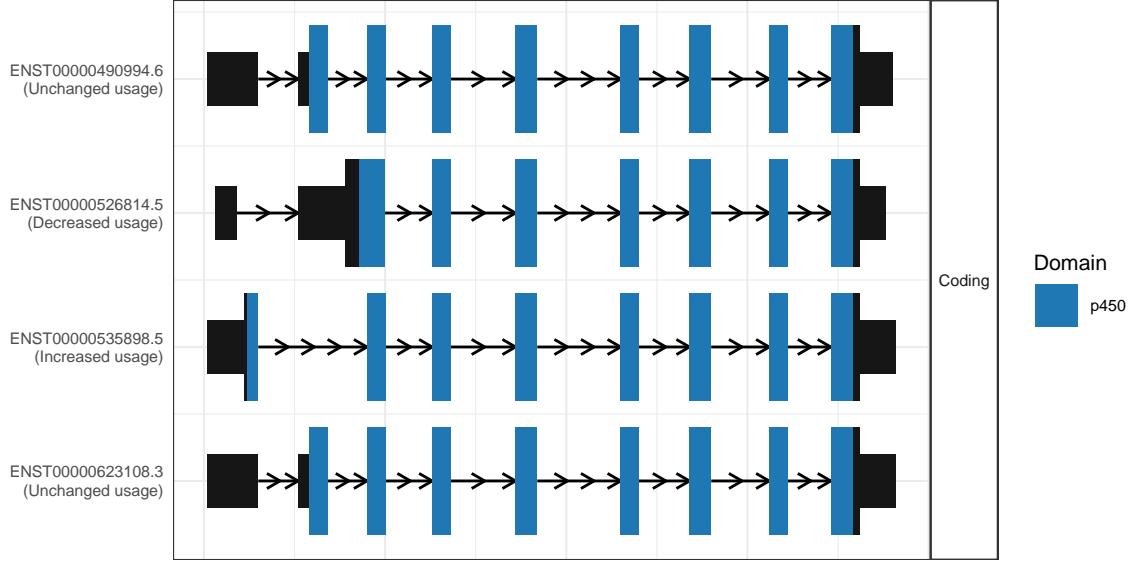


SPRY



zf-B_box

The isoform switch in CYP2C8 (FT vs RGC)

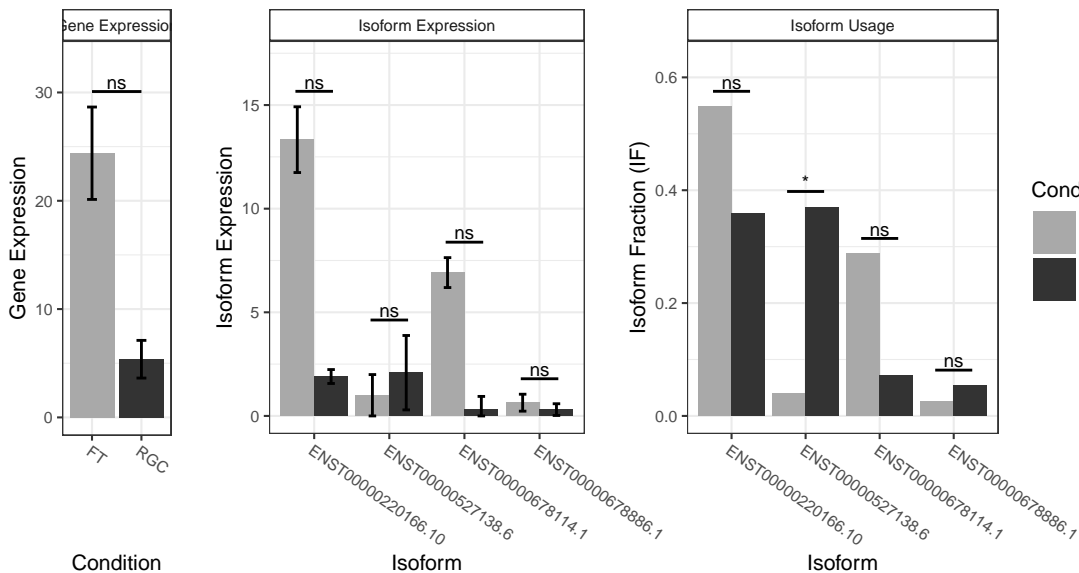
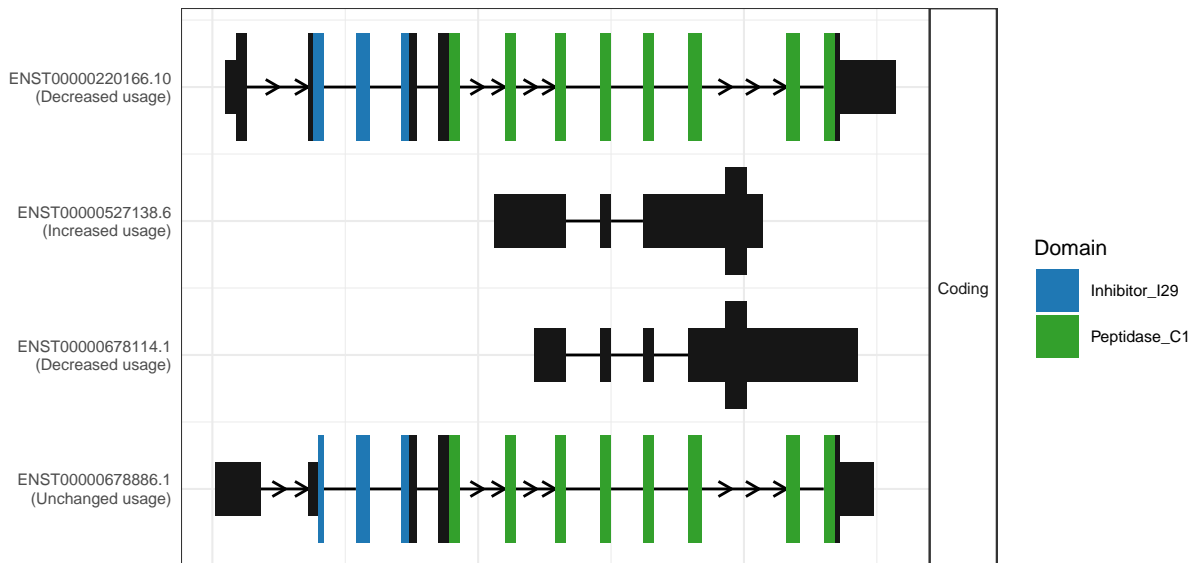


Domain



p450

The isoform switch in CTSH (FT vs RGC)



Domain

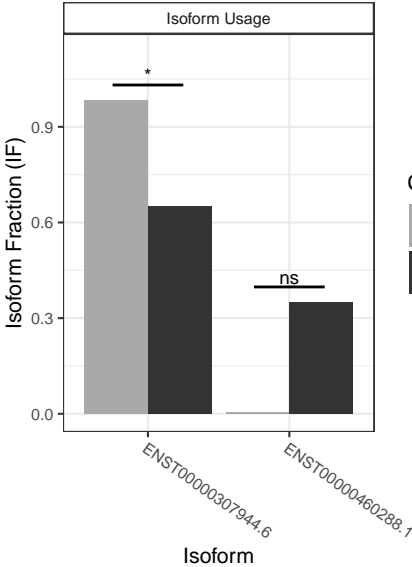
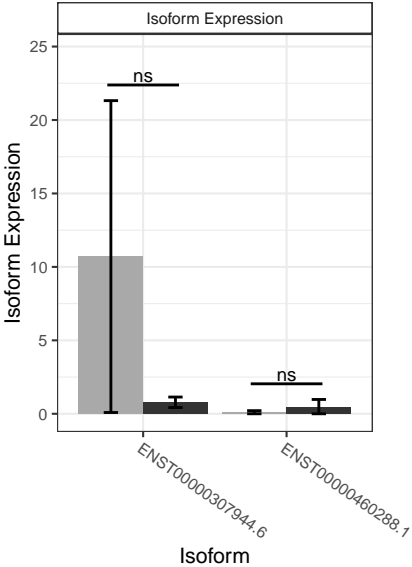
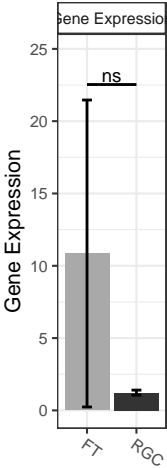
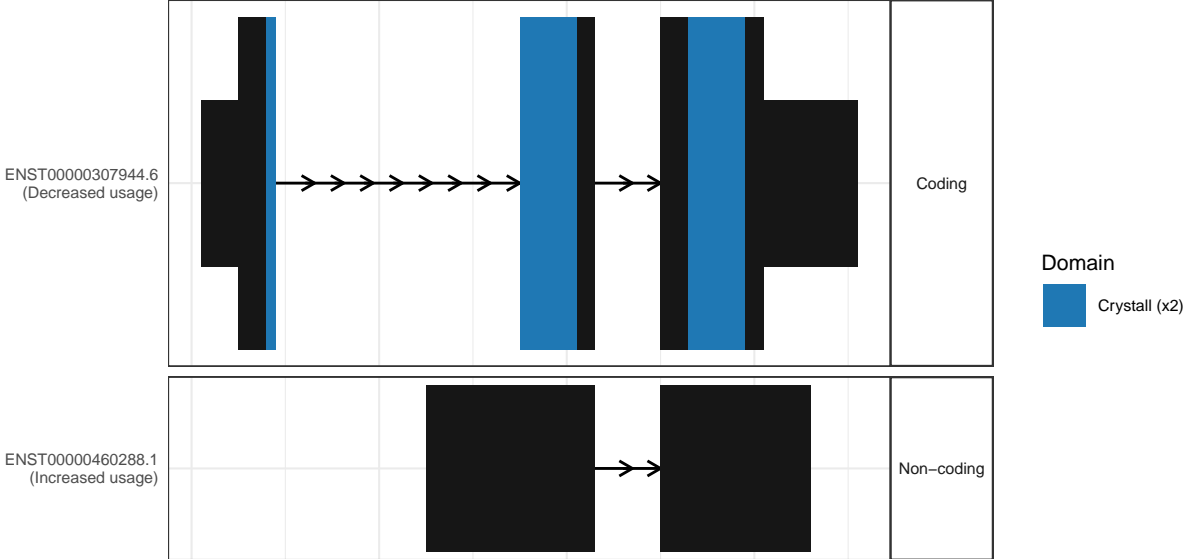


Inhibitor_I29



Peptidase_C1

The isoform switch in CRYGS (FT vs RGC)

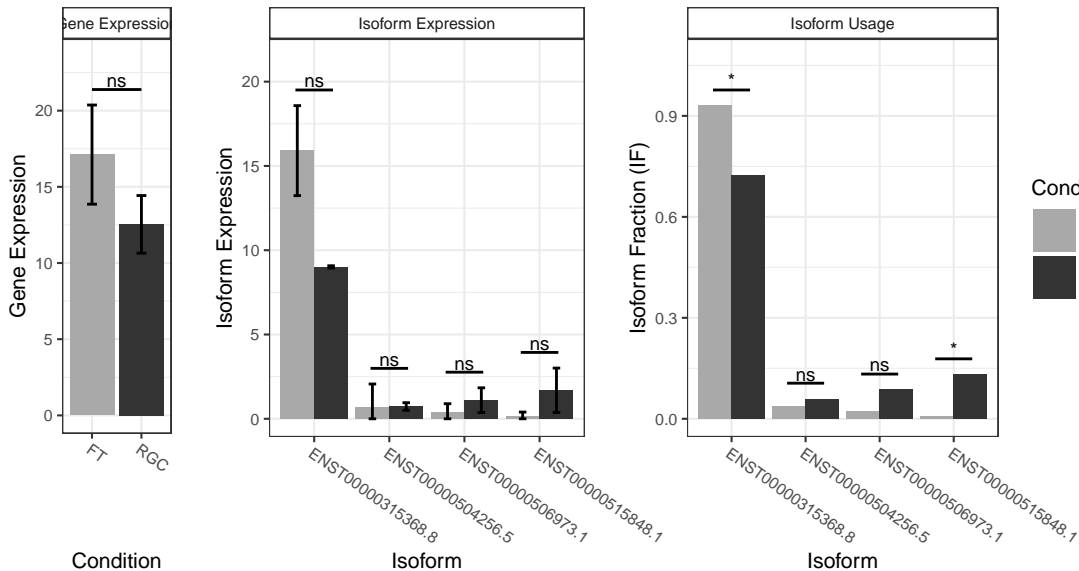
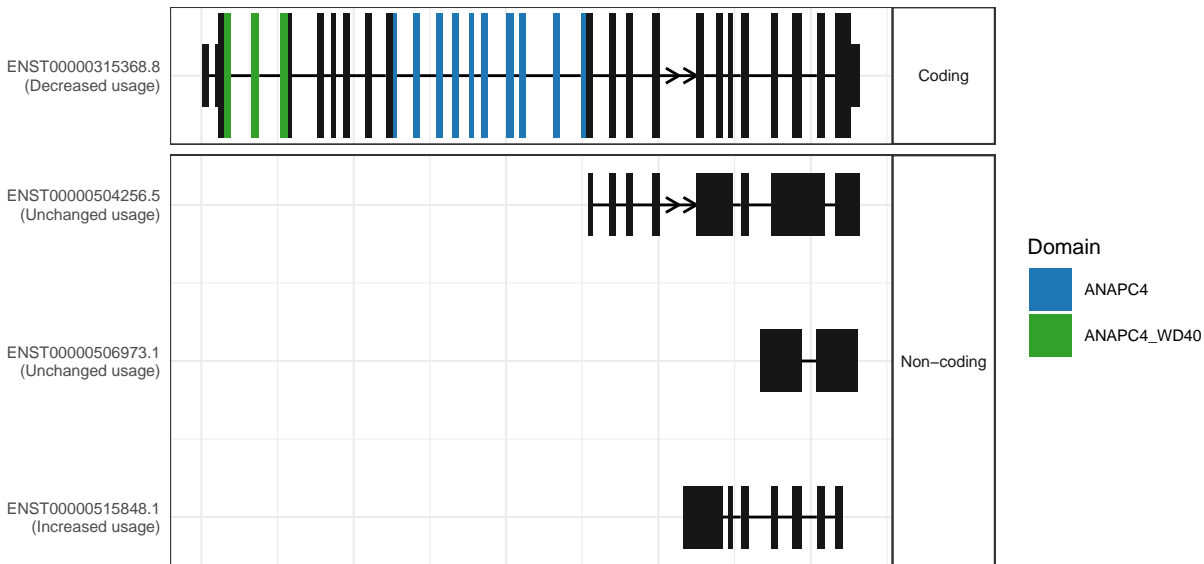


Domain



Crystall (x2)

The isoform switch in ANAPC4 (FT vs RGC)



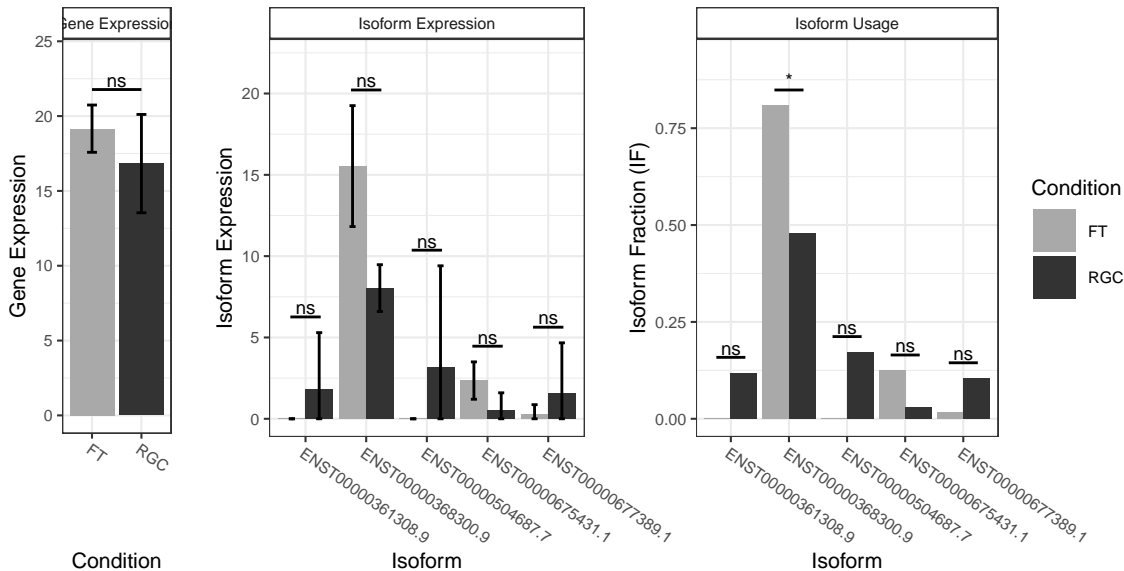
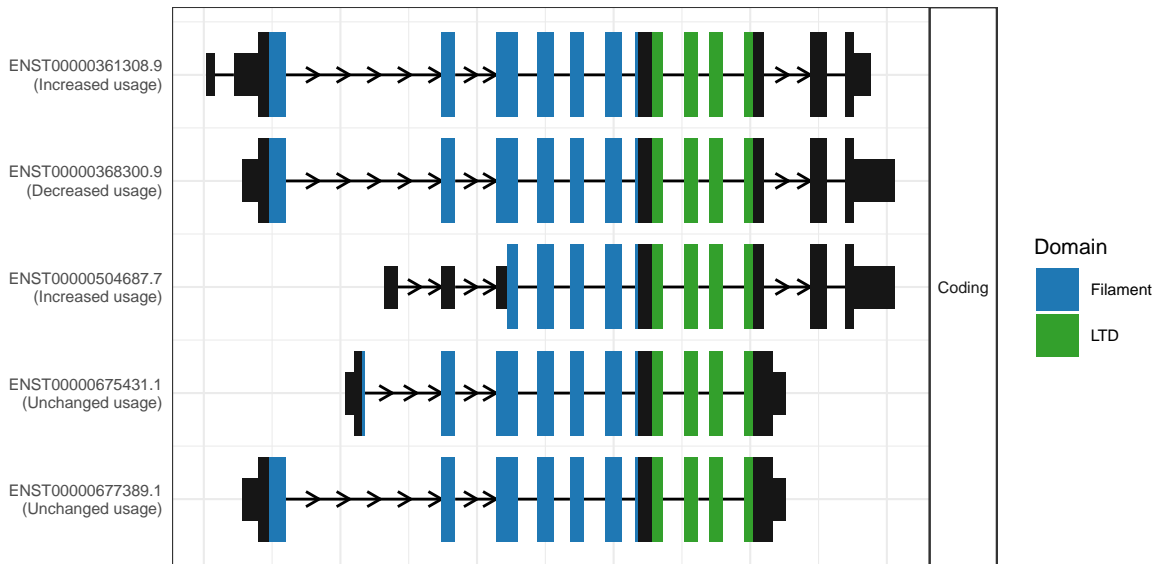
Domain



ANAPC4

ANAPC4_WD40

The isoform switch in LMNA (FT vs RGC)



Domain

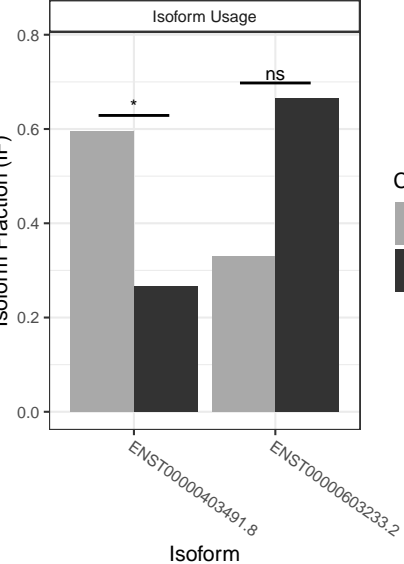
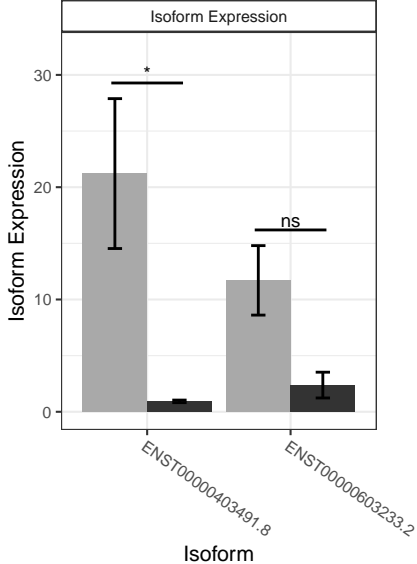
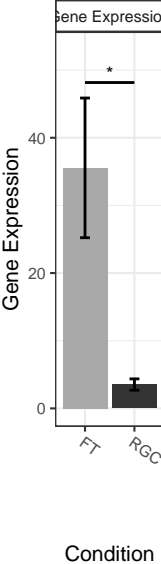
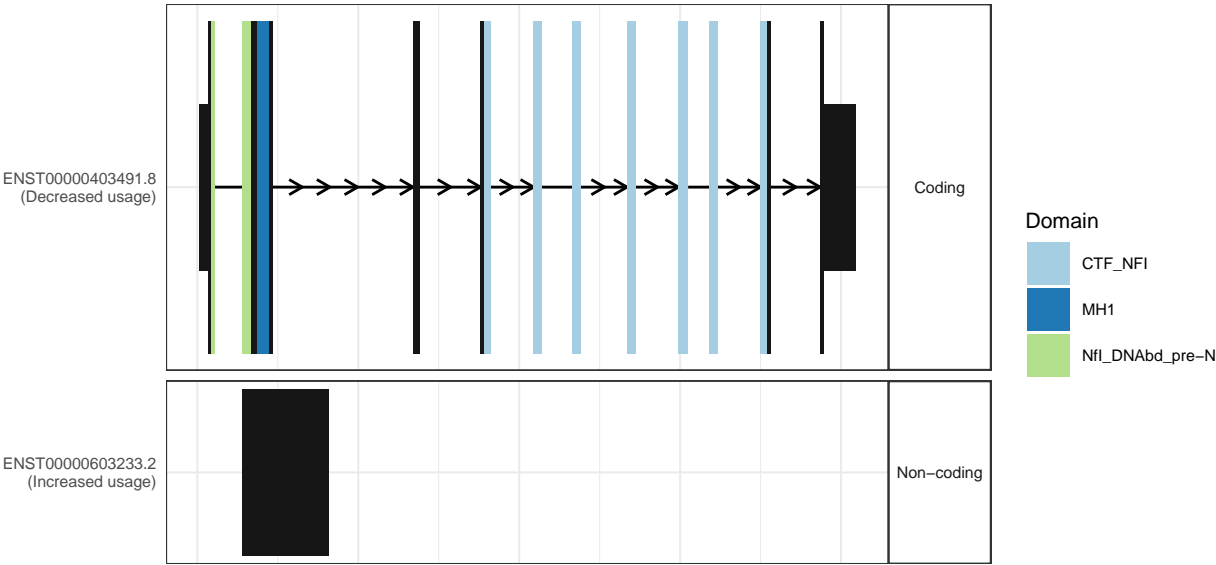


Filament



LTD

The isoform switch in NFIA (FT vs RGC)



Domain



CTF_NFI

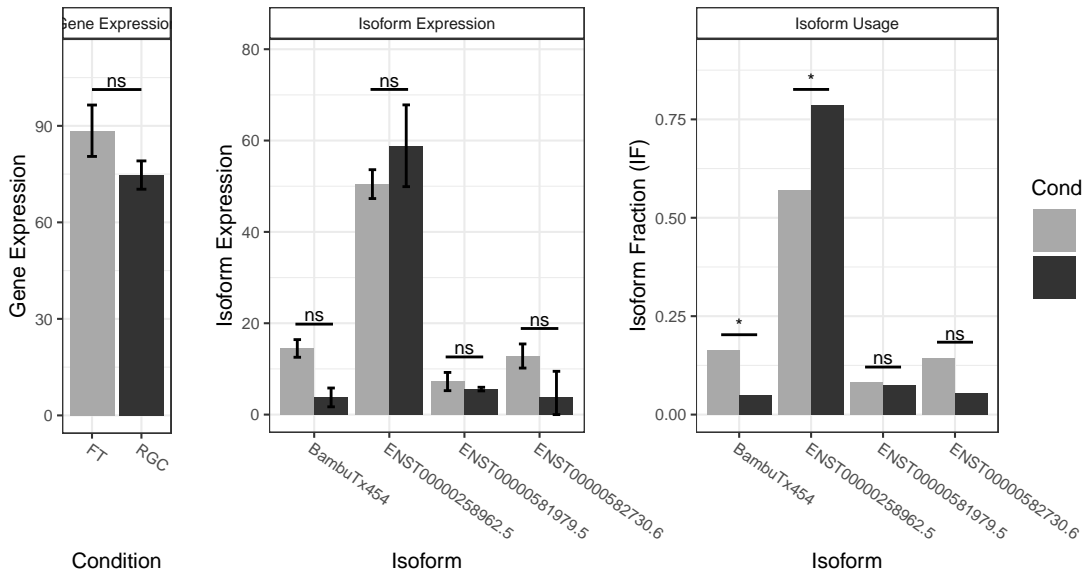
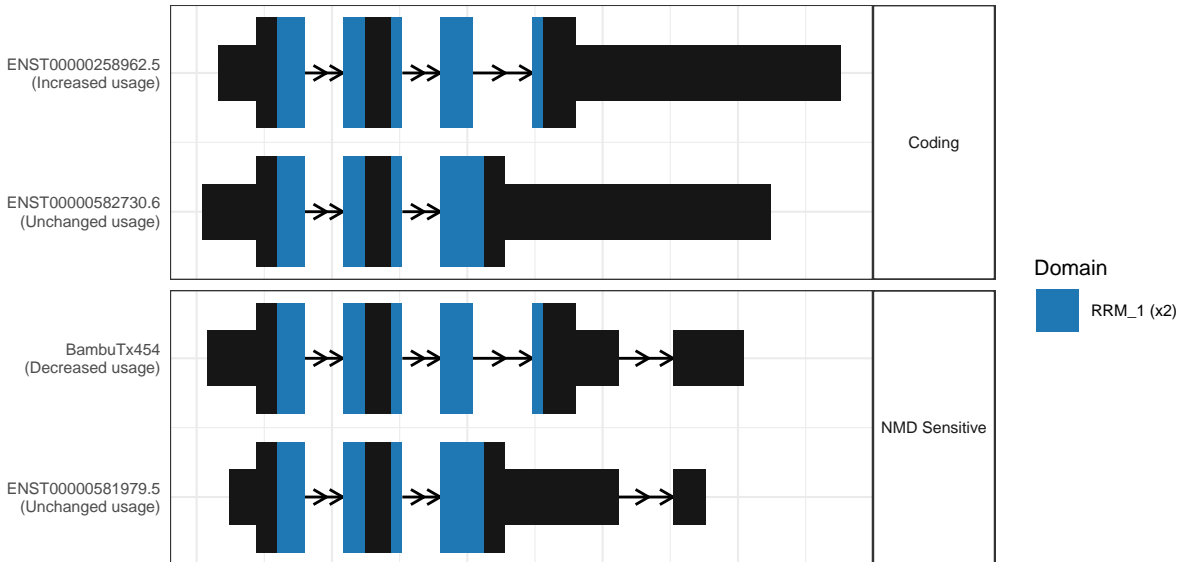


MH1




Nfil_DNAAbd_pre-N

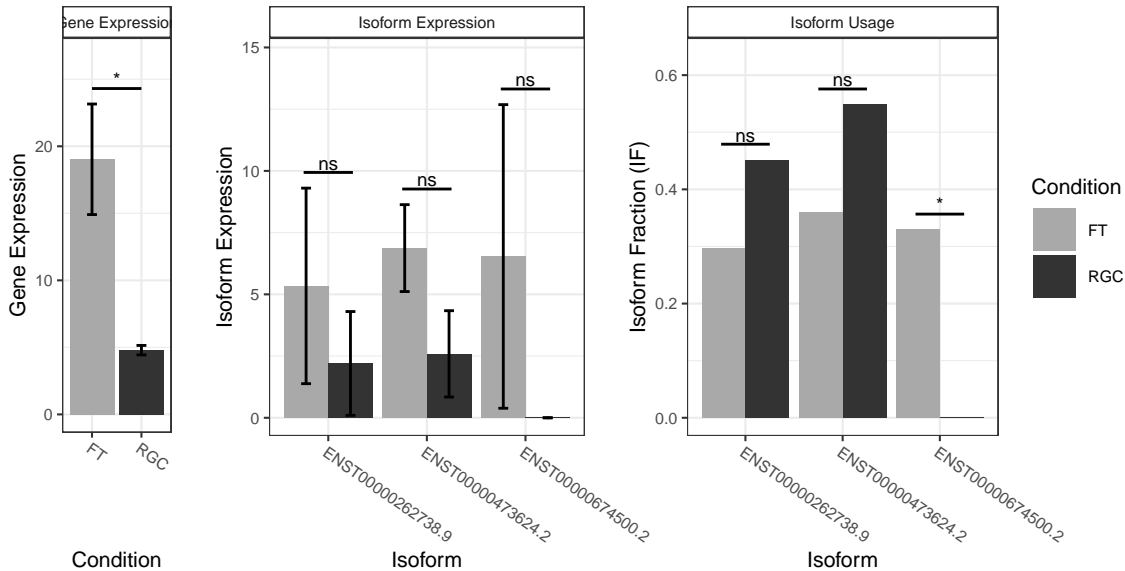
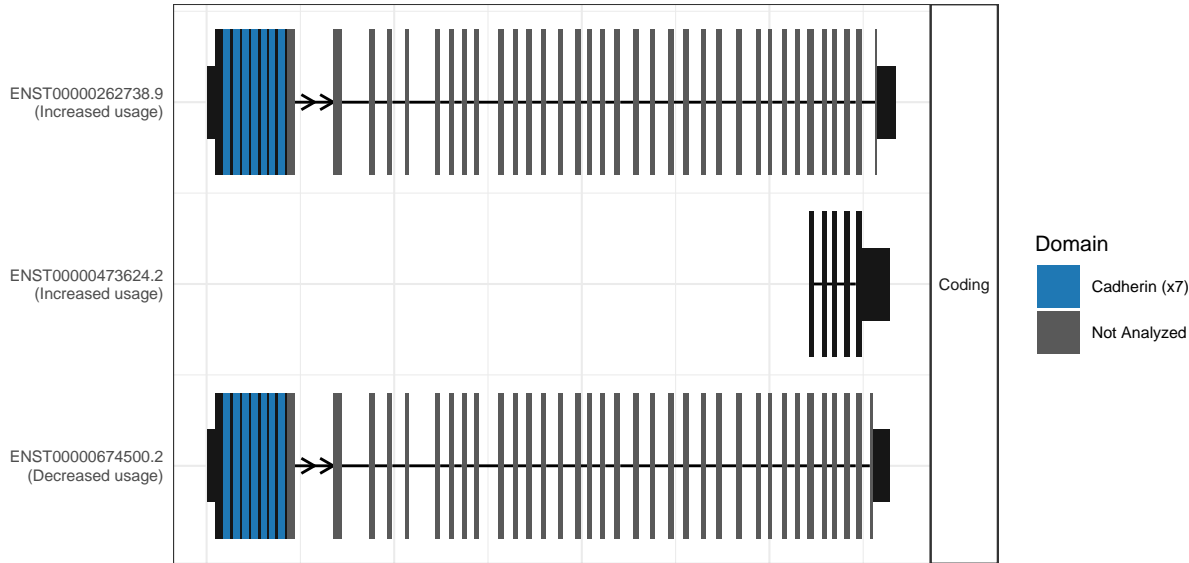
The isoform switch in SRSF1 (FT vs RGC)



Domain

 RRM_1 (x2)

The isoform switch in CELSR1 (FT vs RGC)



Domain

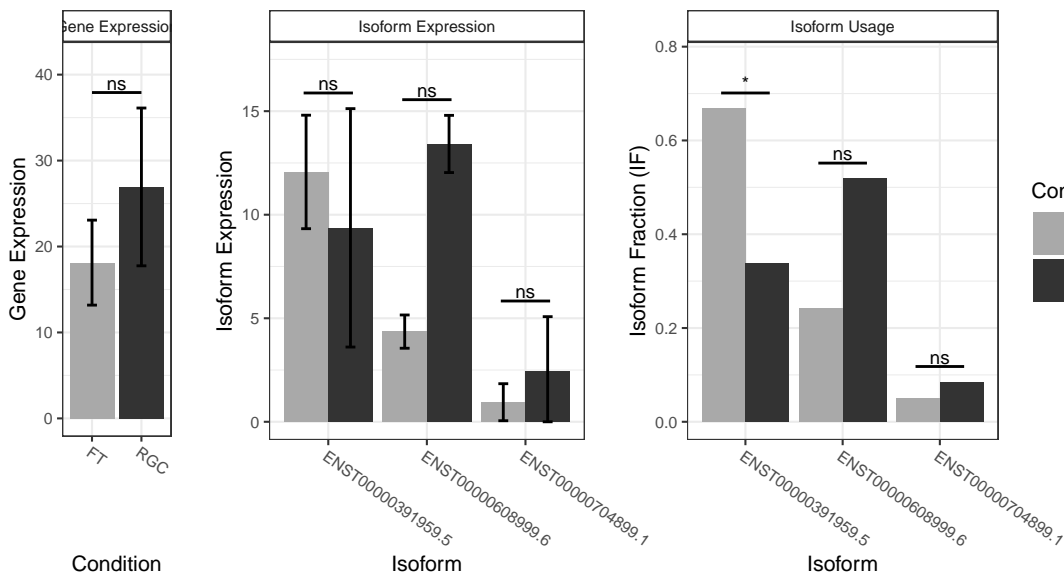
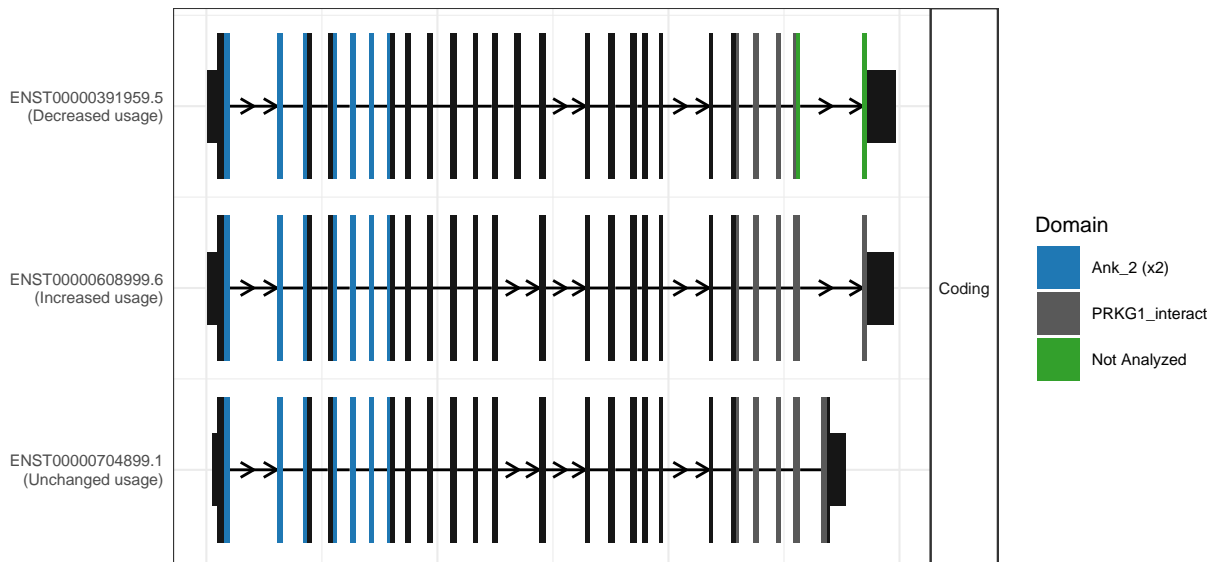


Cadherin (x7)

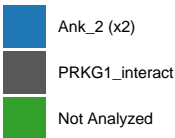


Not Analyzed

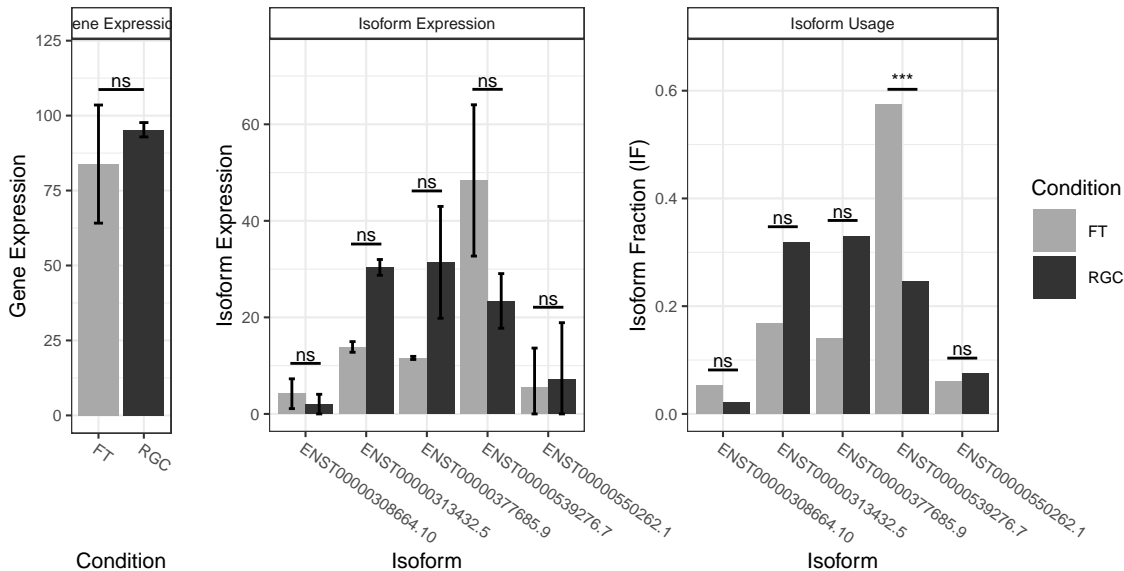
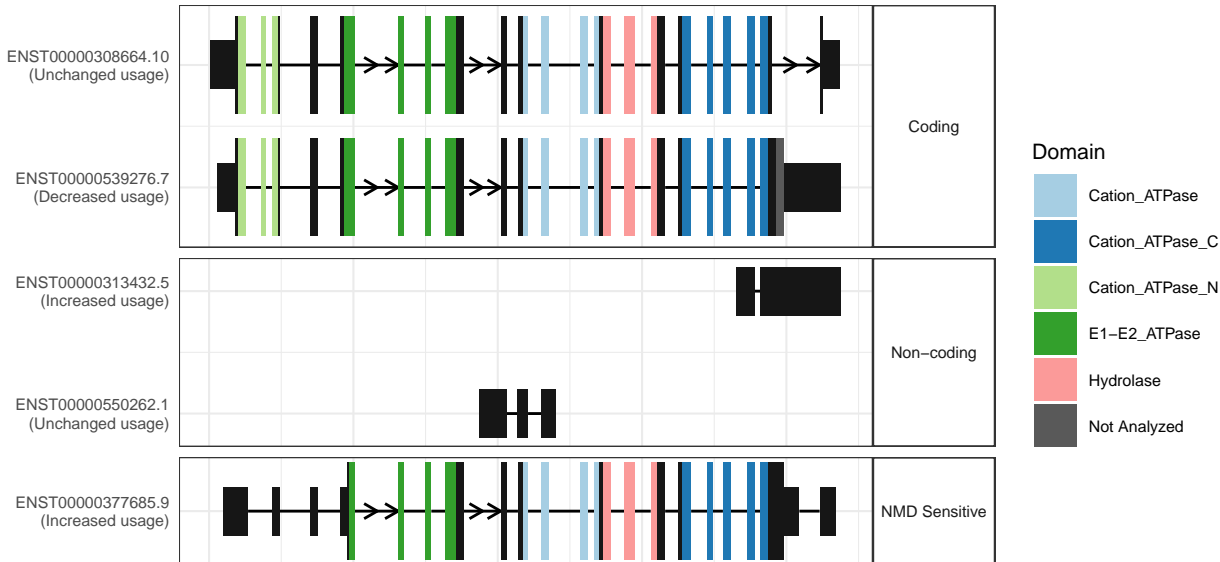
The isoform switch in PPP1R12B (FT vs RGC)



Domain



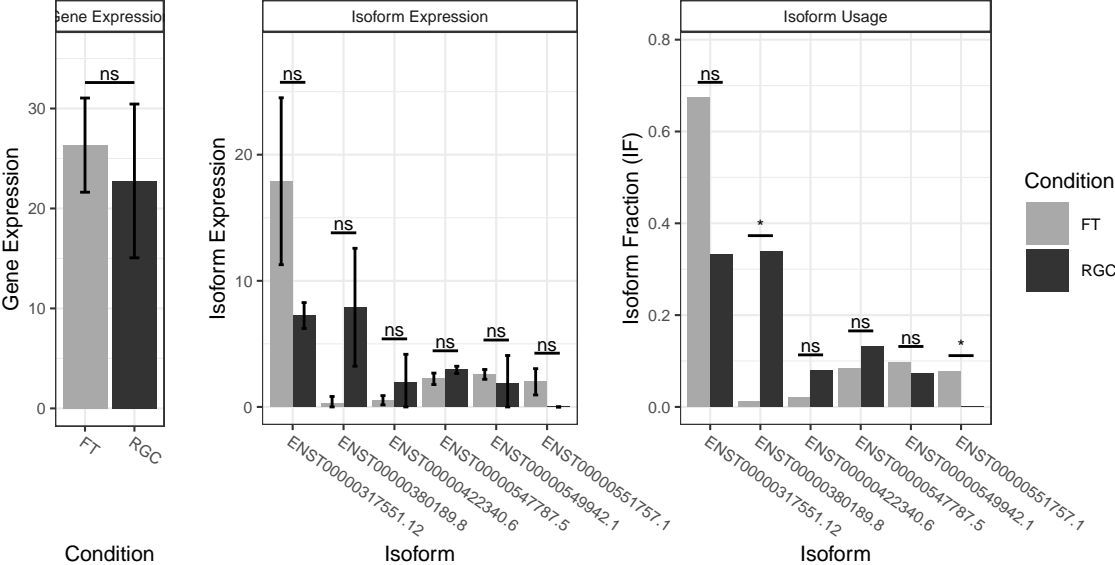
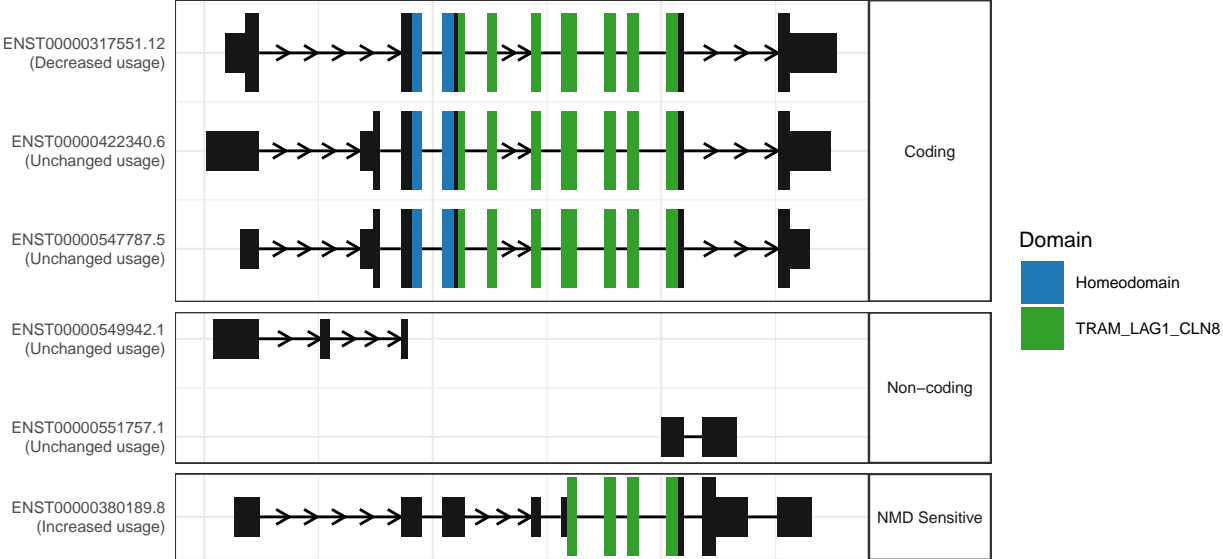
The isoform switch in ATP2A2 (FT vs RGC)



Domain

	Cation_ATPase
	Cation_ATPase_C
	Cation_ATPase_N
	E1-E2_ATPase
	Hydrolase
	Not Analyzed

The isoform switch in CERS5 (FT vs RGC)



Domain

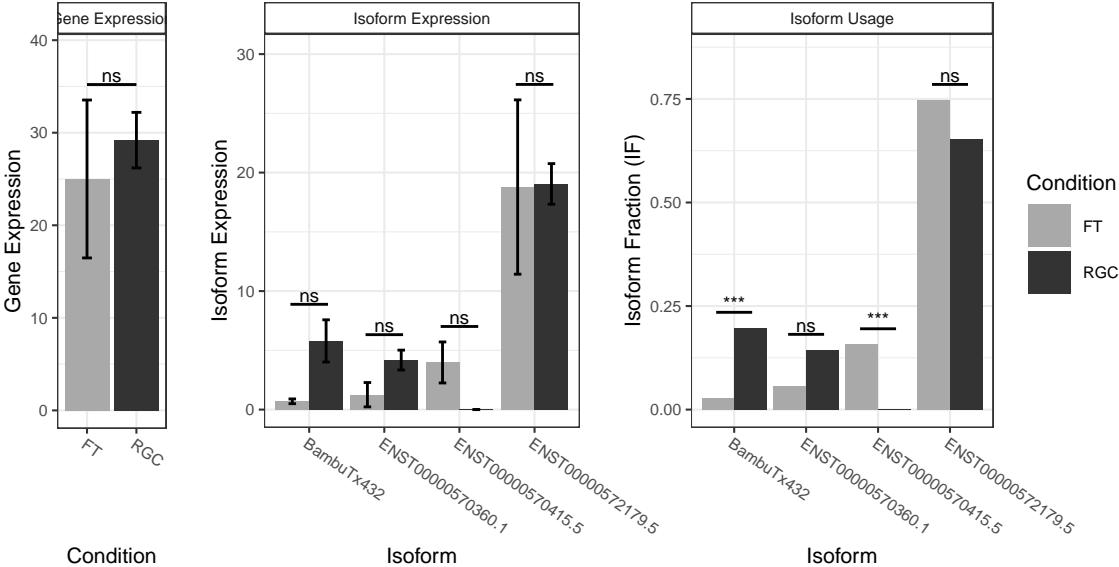
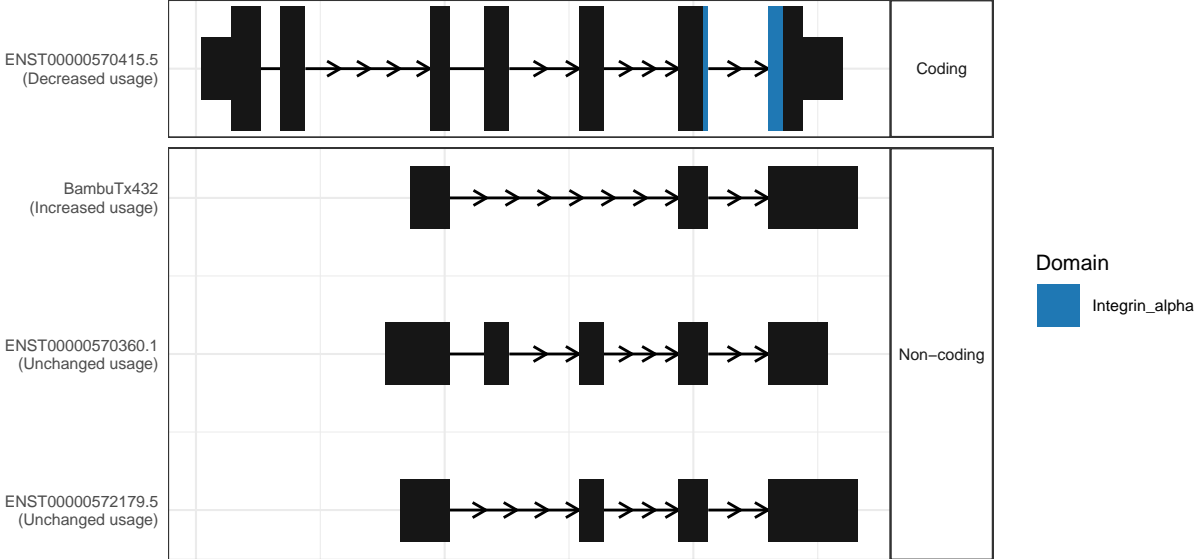


Homeodomain



TRAM_LAG1_CLN8

The isoform switch in ITGAE (FT vs RGC)

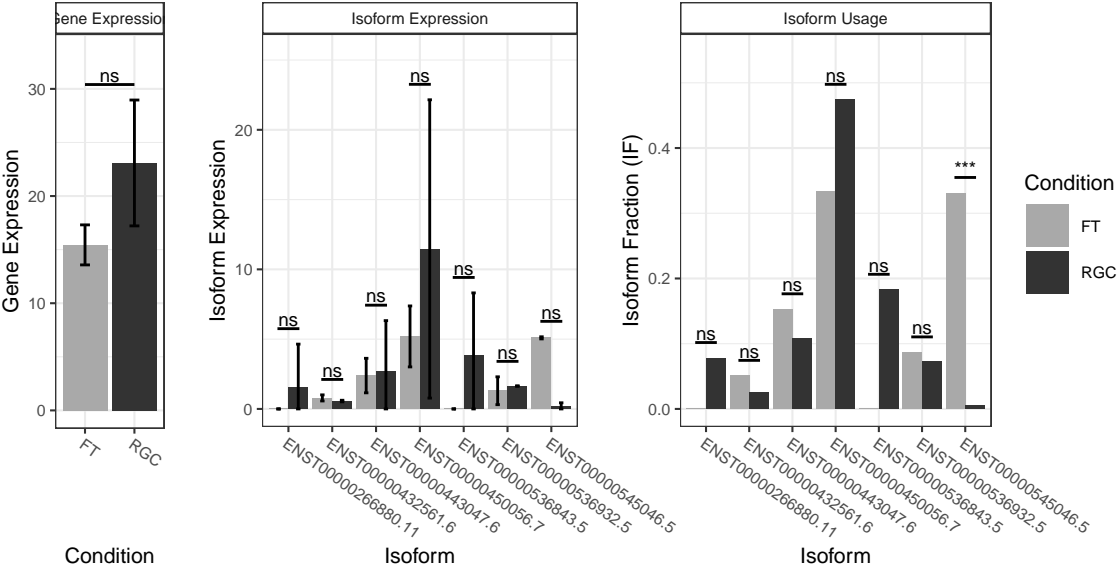
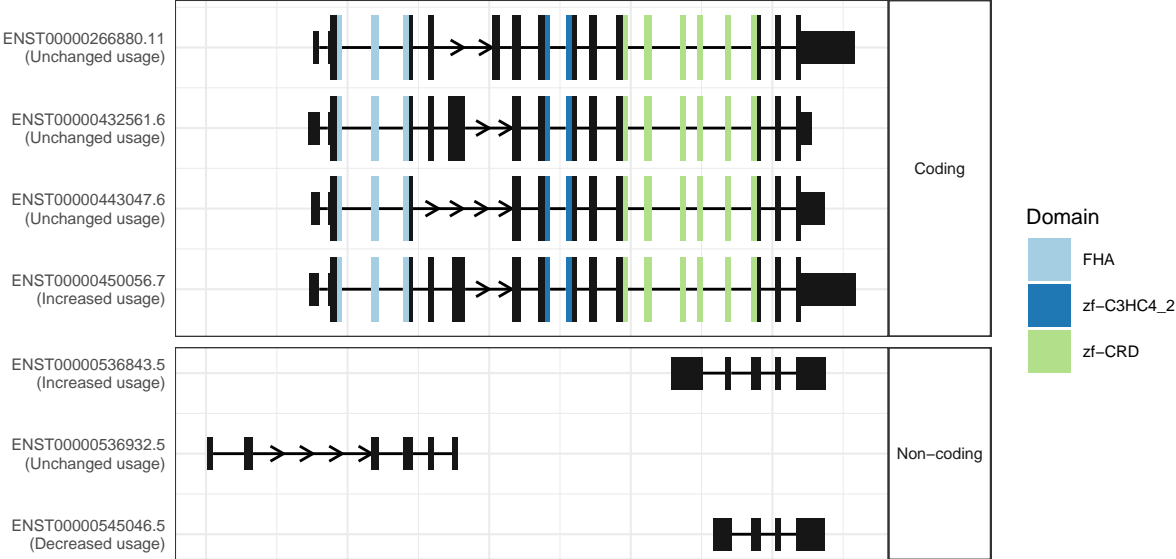


Domain






Integrin_alpha

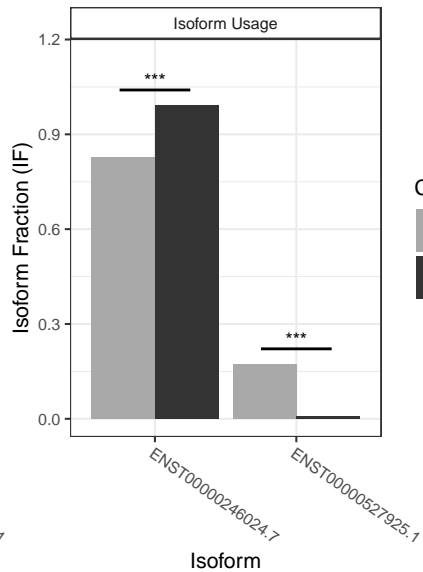
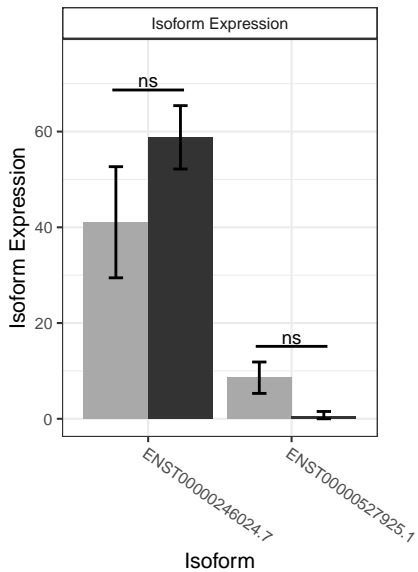
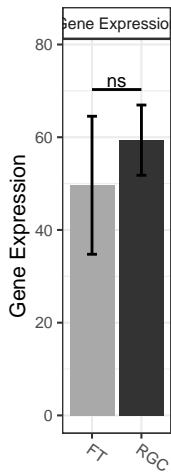
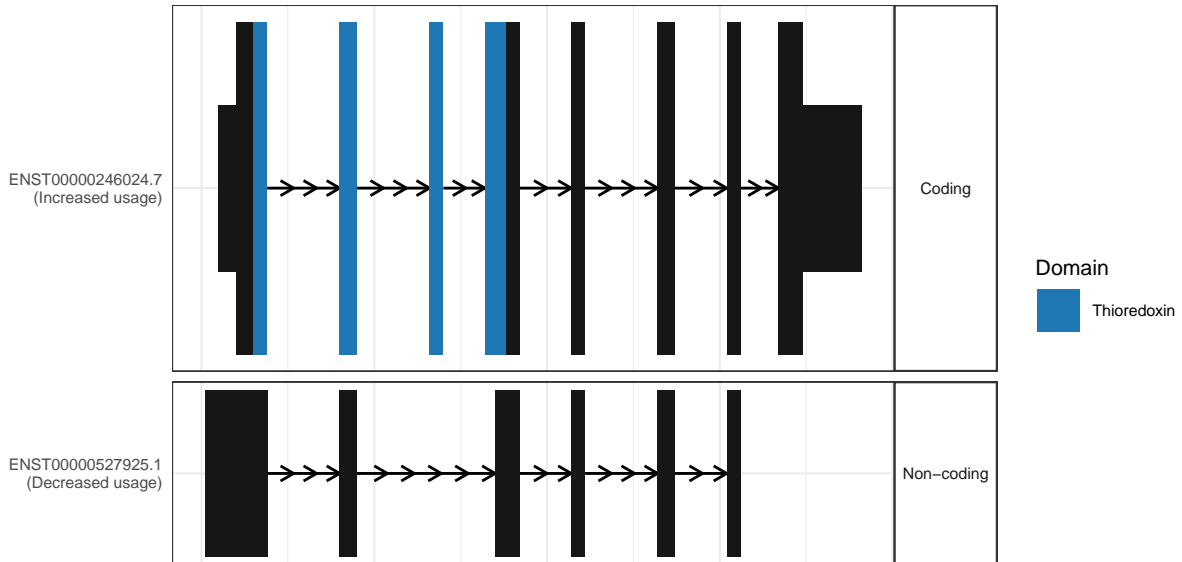
The isoform switch in CHFR (FT vs RGC)



Domain

	FHA
	zf-C3HC4_2
	zf-CRD

The isoform switch in TMX4 (FT vs RGC)

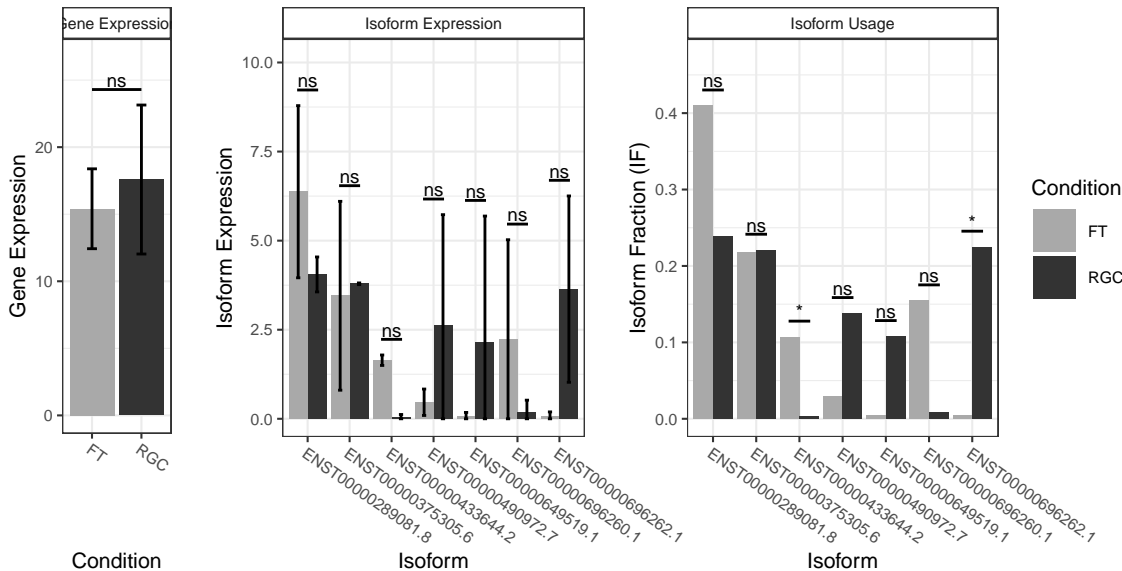
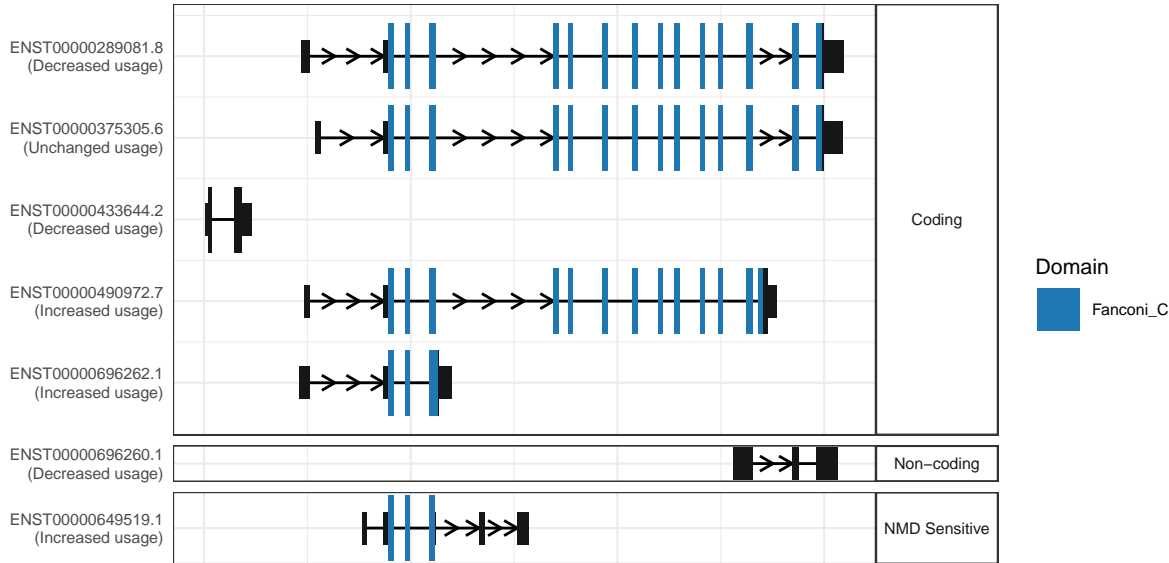


Domain



Thioredoxin

The isoform switch in FANCC (FT vs RGC)

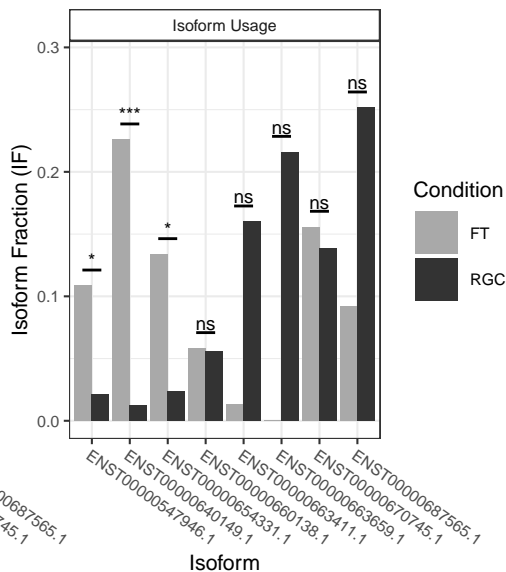
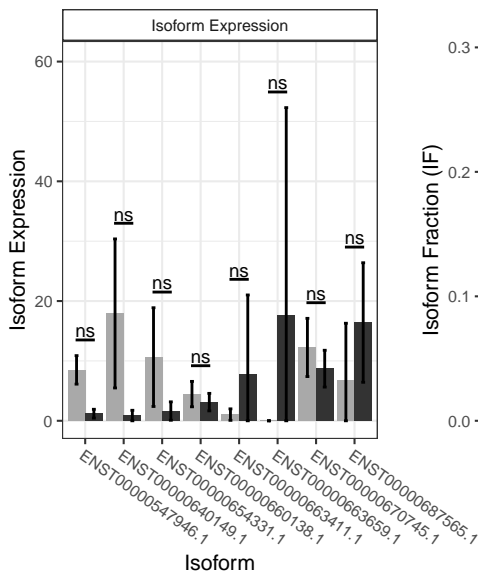
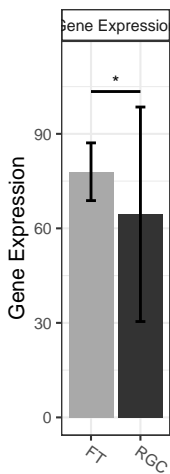
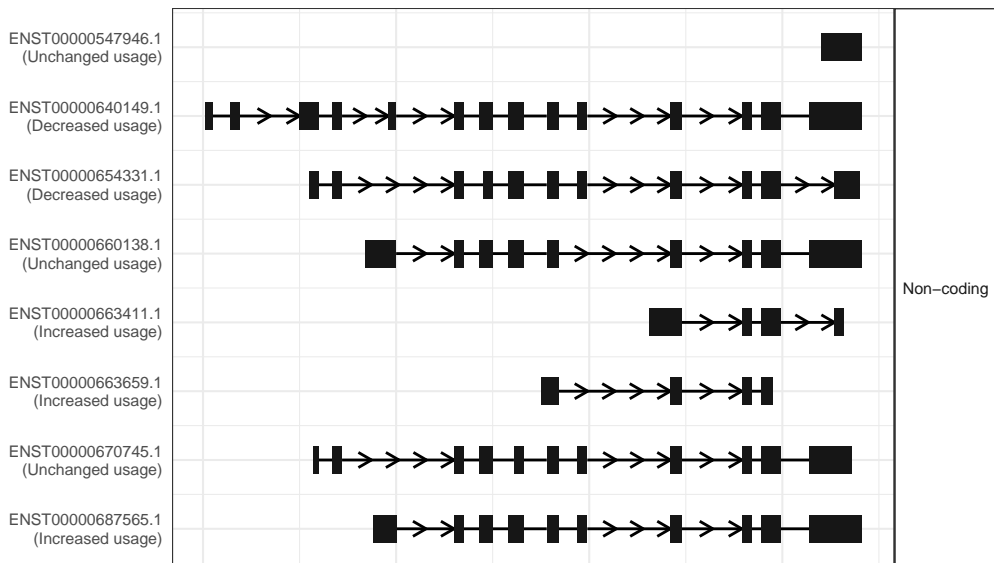


Domain

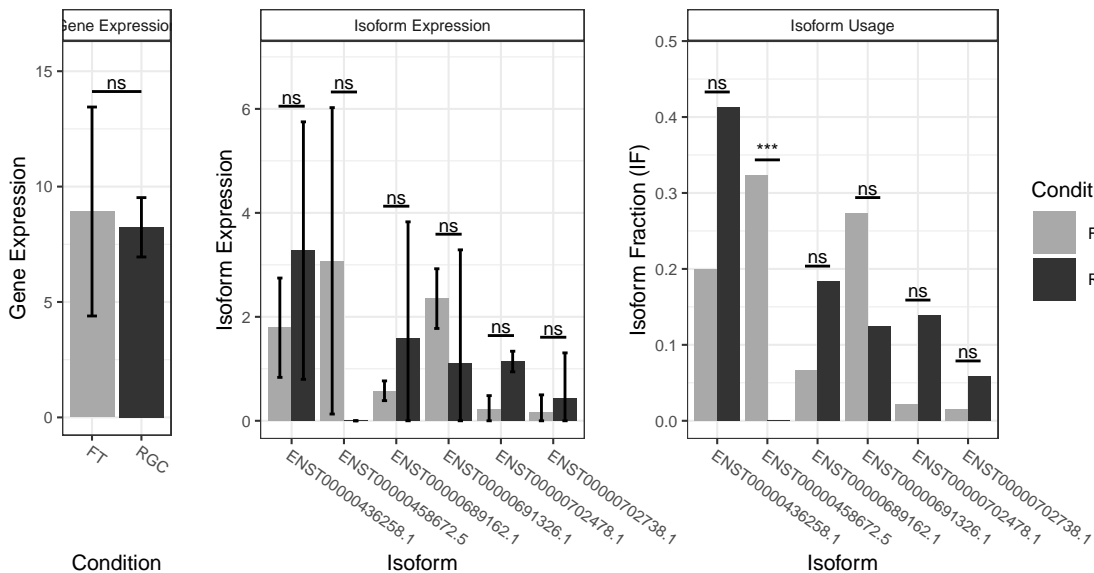
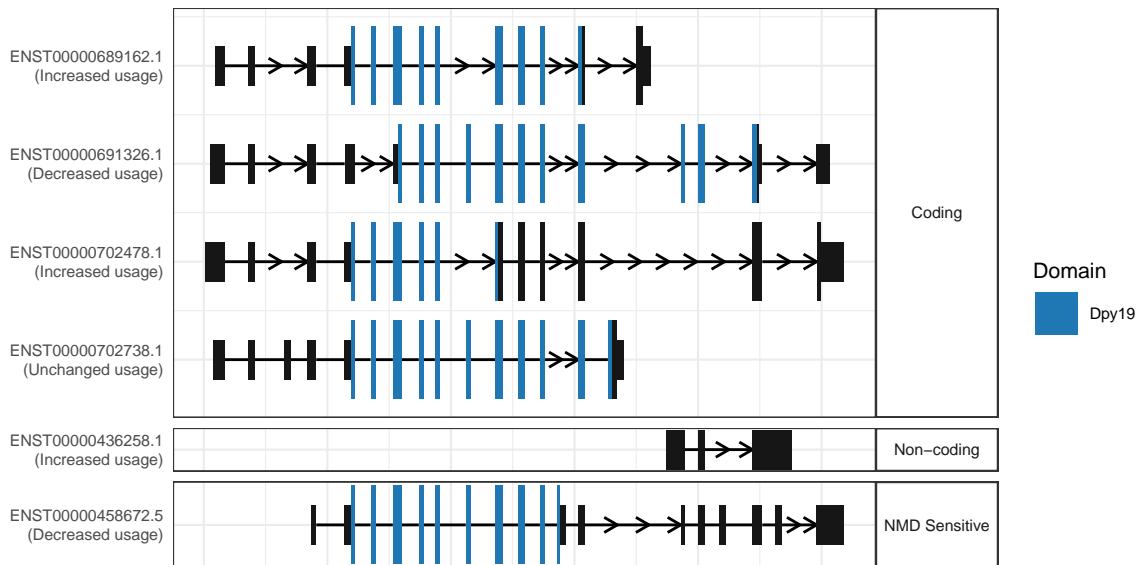


Fanconi_C

The isoform switch in RMST (FT vs RGC)



The isoform switch in (FT vs RGC)

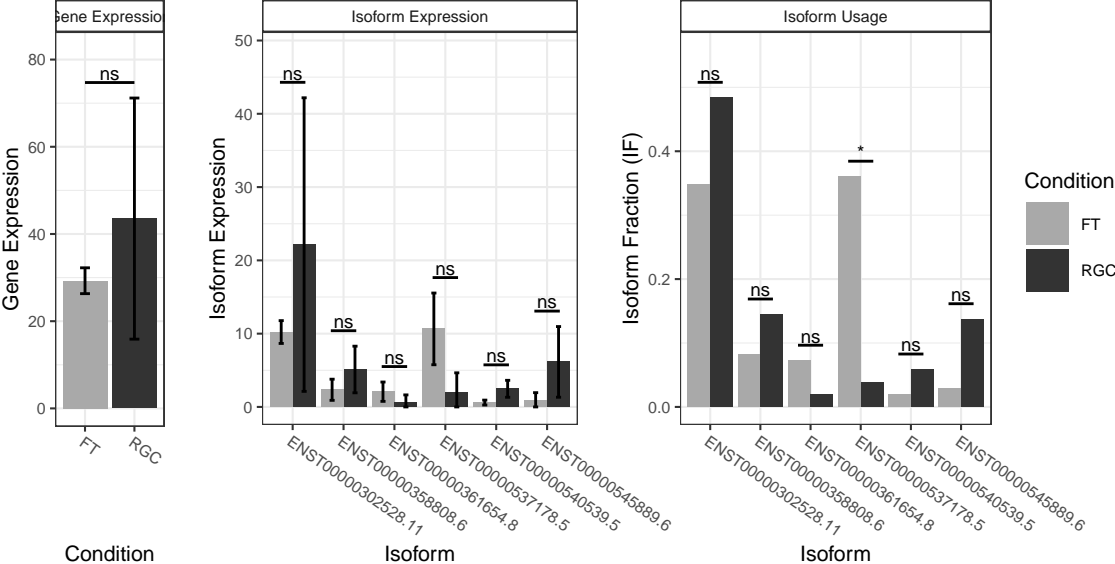
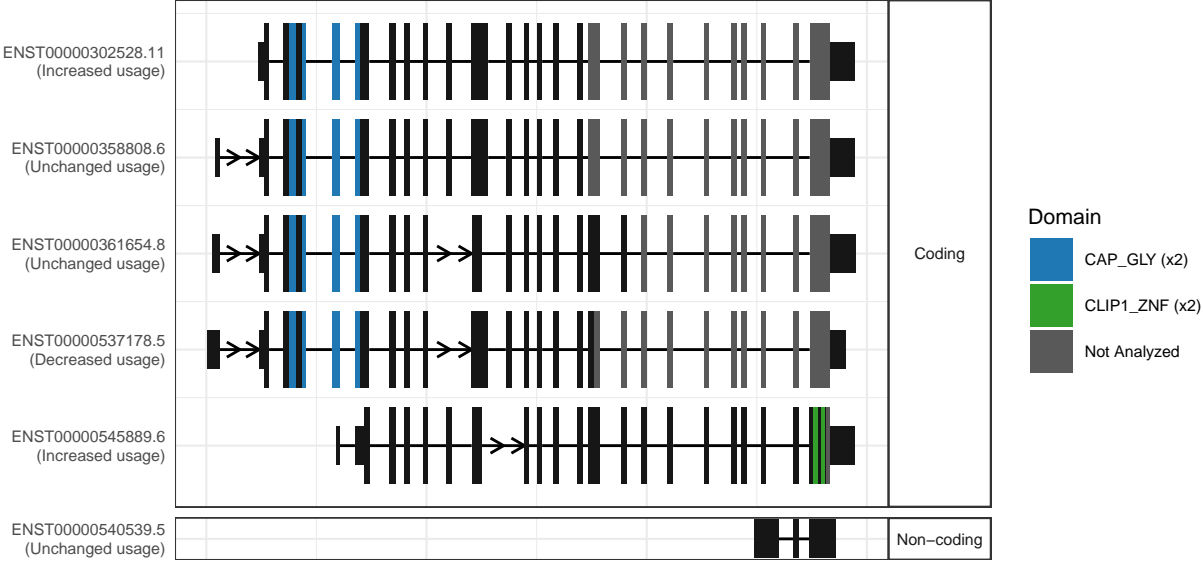


Domain






Dpy19

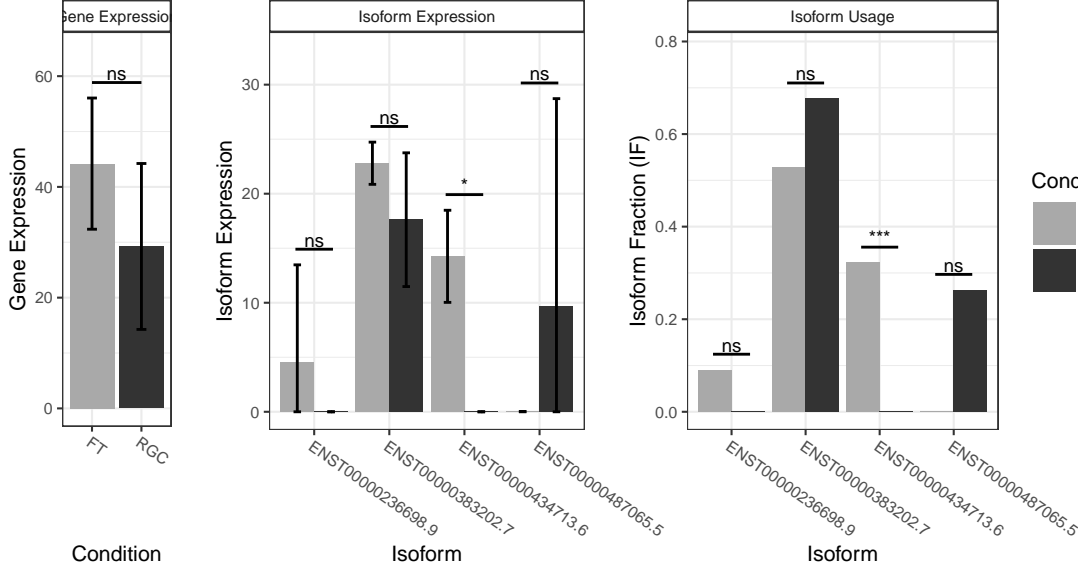
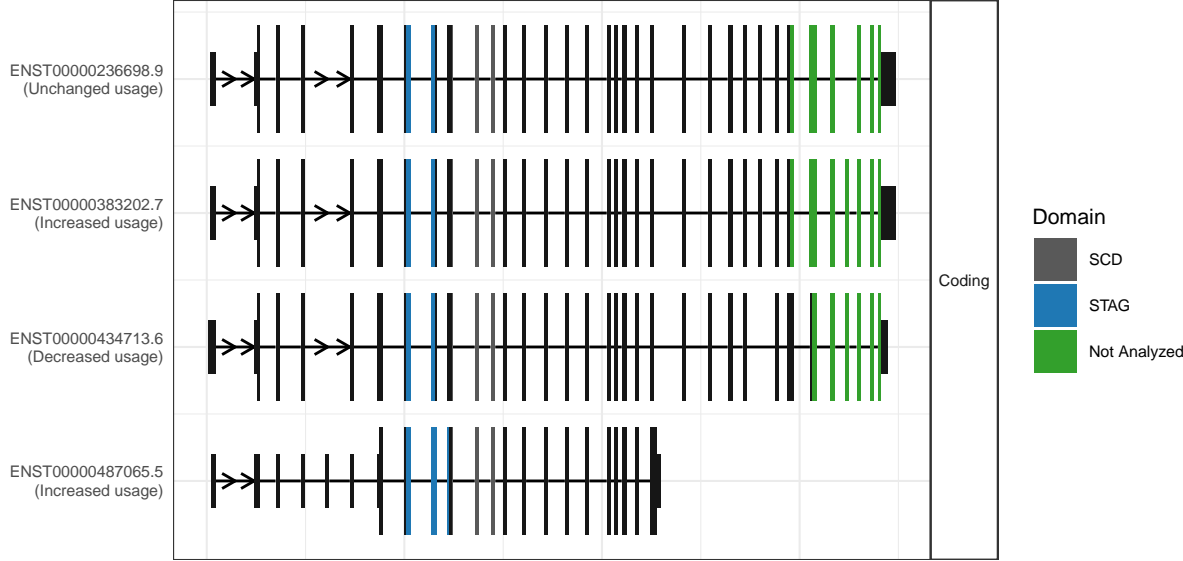
The isoform switch in CLIP1 (FT vs RGC)



Domain

	CAP_GLY (x2)
	CLIP1_ZNF (x2)
	Not Analyzed

The isoform switch in STAG1 (FT vs RGC)



Domain



SCD

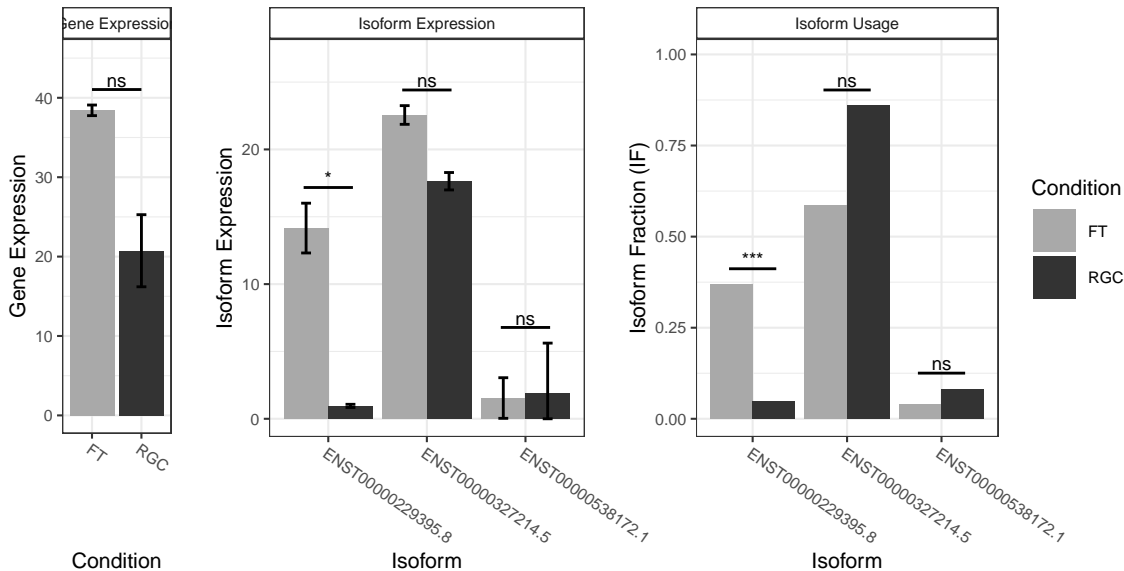
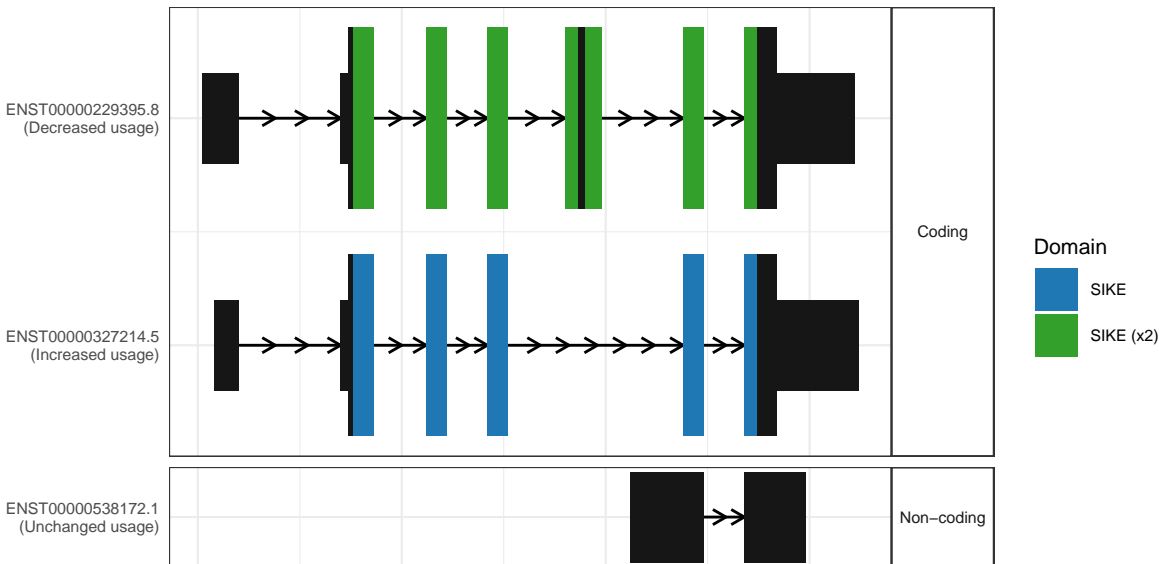


STAG



Not Analyzed

The isoform switch in FGFR1OP2 (FT vs RGC)

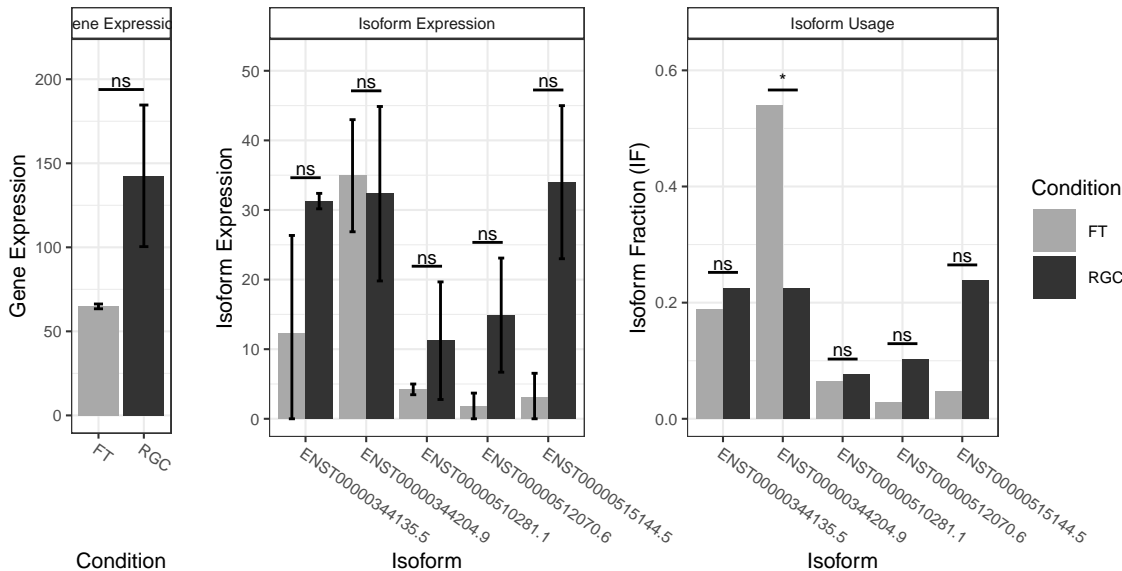
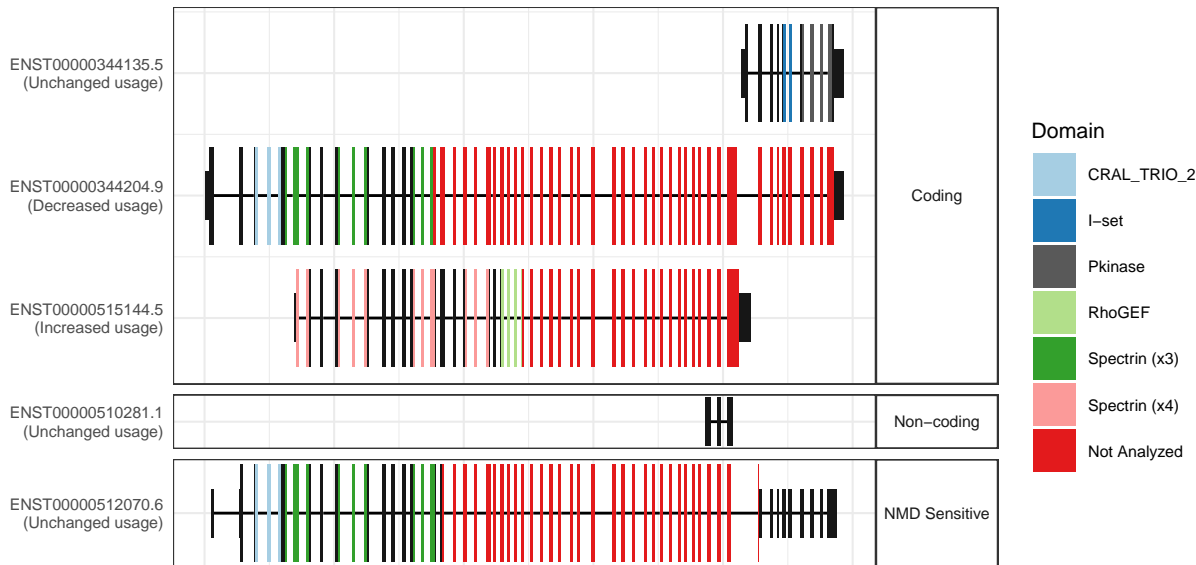


Domain

 SIKE

 SIKE (x2)

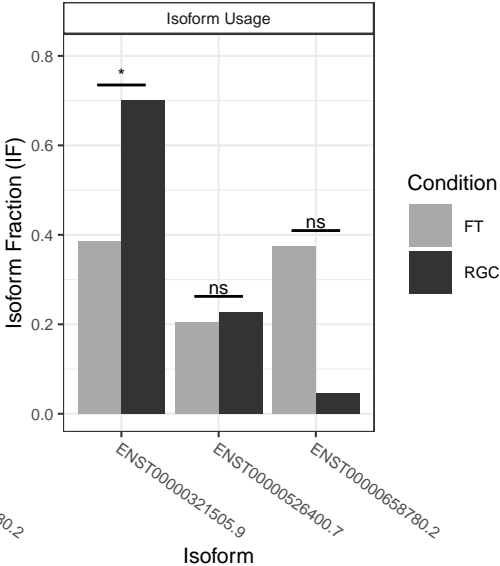
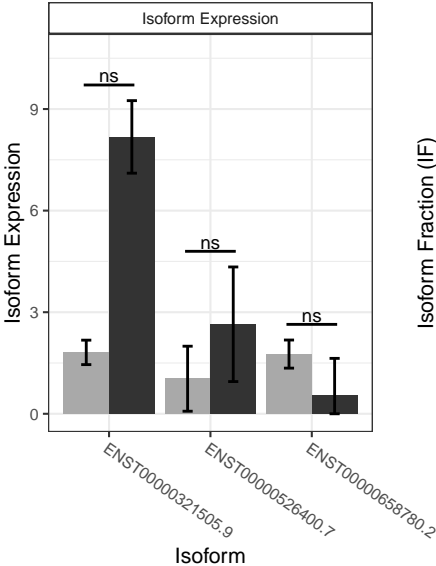
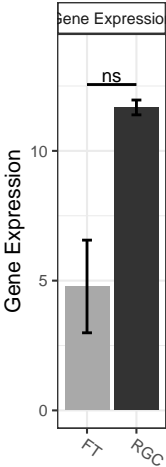
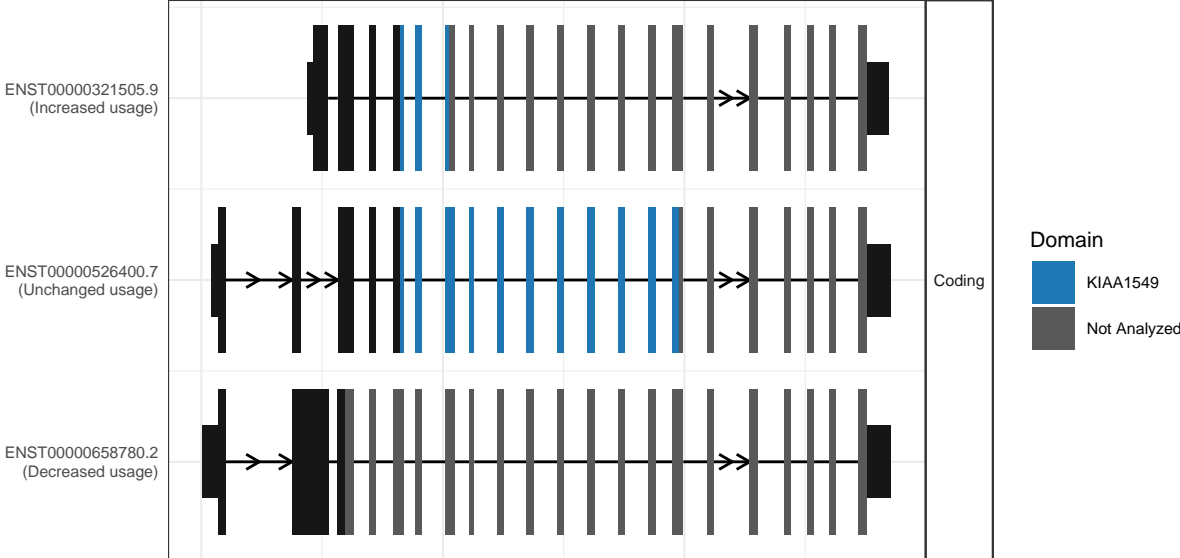
The isoform switch in TRIO (FT vs RGC)



Domain

	CRAL_TRIO_2
	I-set
	Pkinase
	RhoGEF
	Spectrin (x3)
	Spectrin (x4)
	Not Analyzed

The isoform switch in KIAA1549L (FT vs RGC)



Domain

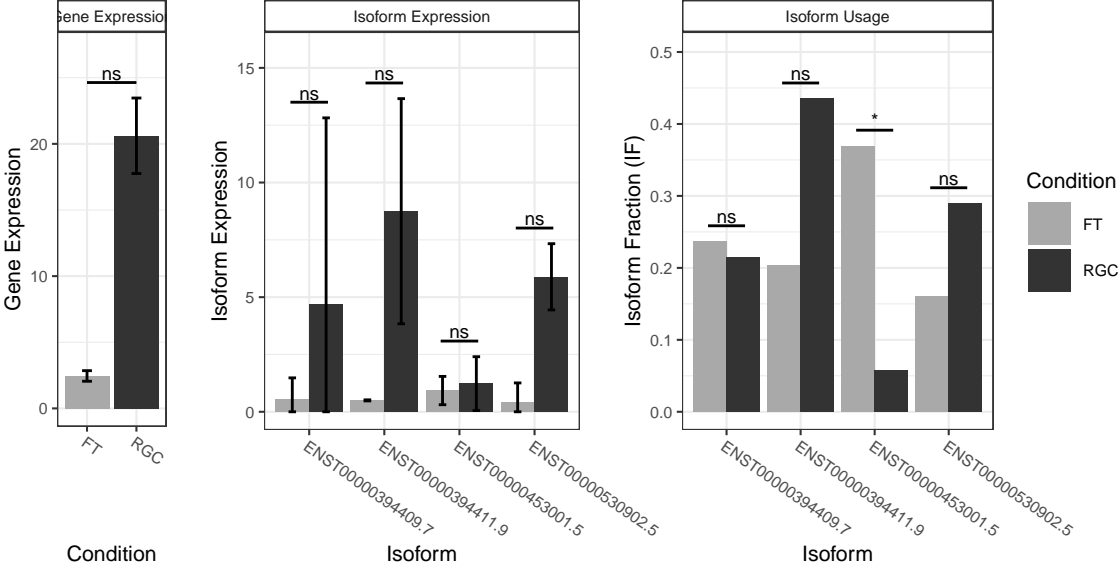
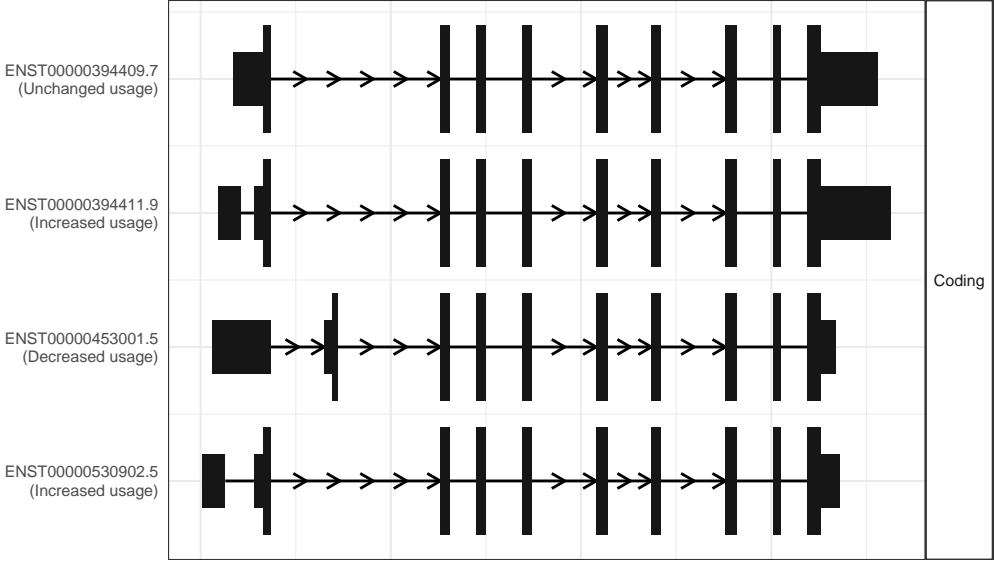


KIAA1549

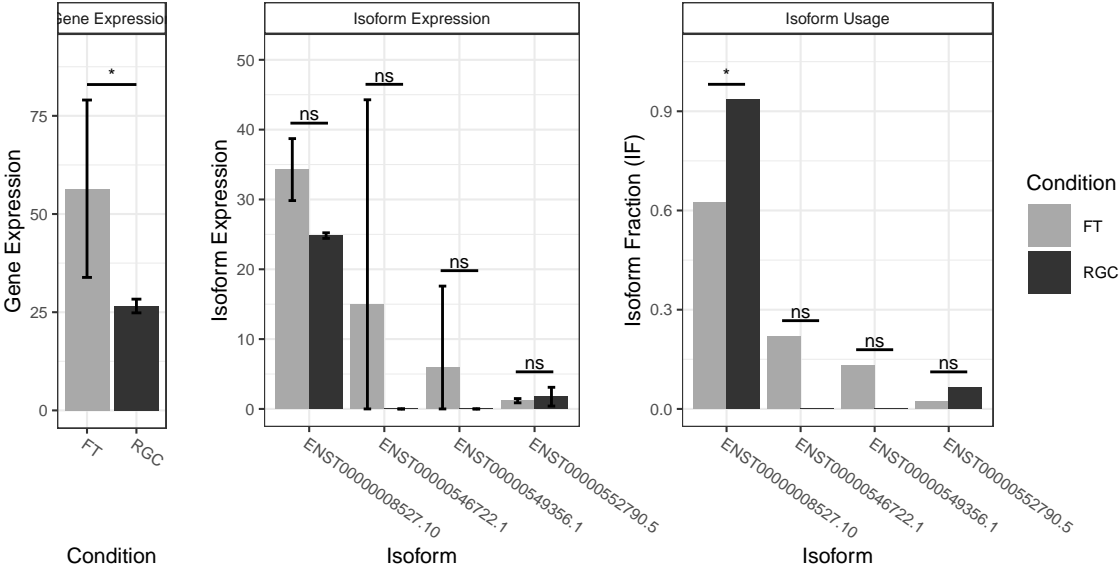
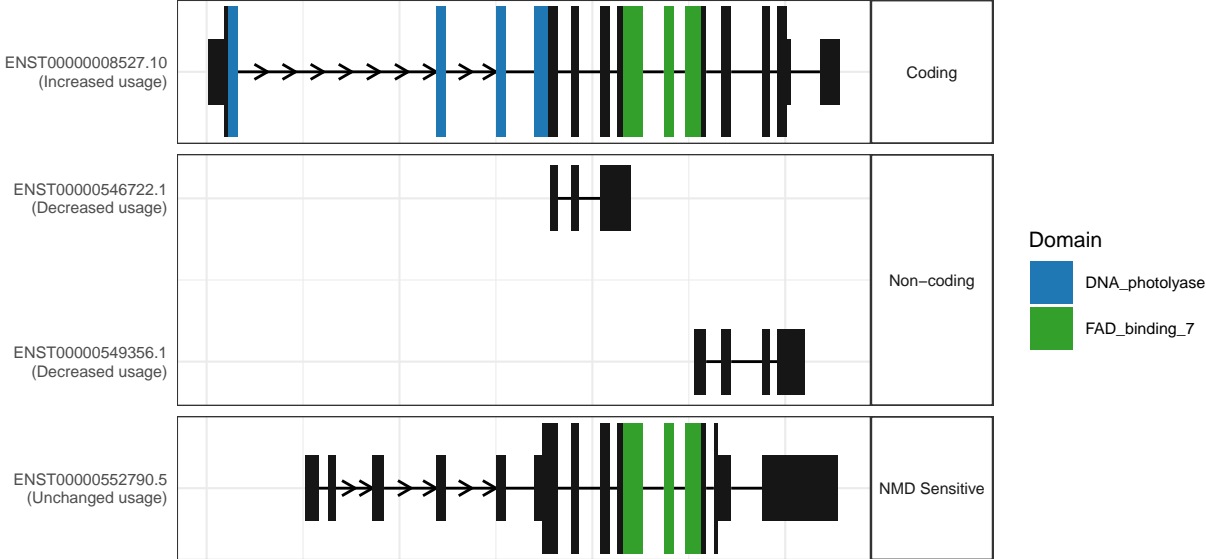


Not Analyzed

The isoform switch in PPP2R2B (FT vs RGC)



The isoform switch in CRY1 (FT vs RGC)



Domain

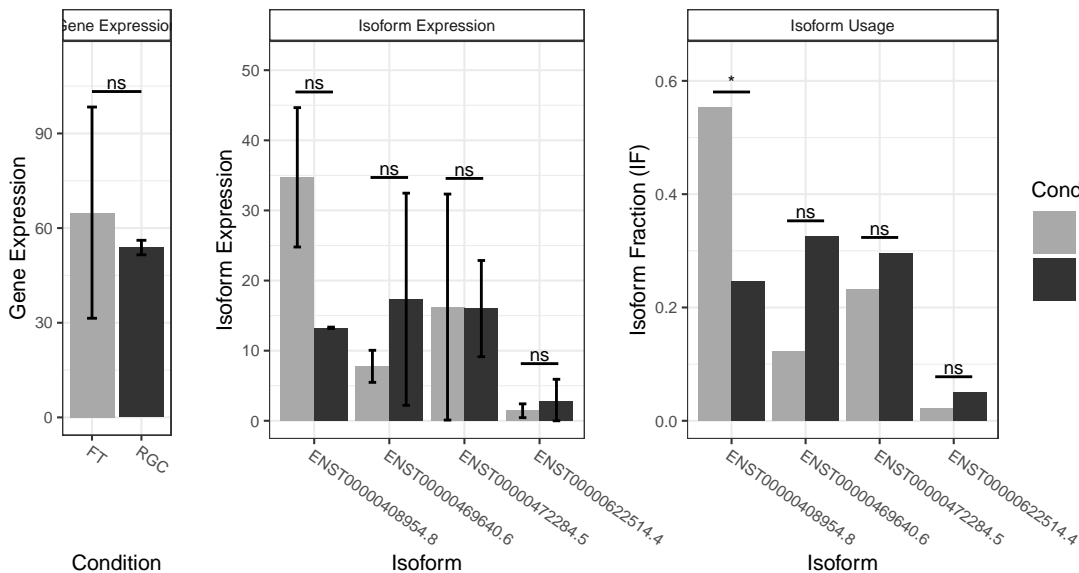
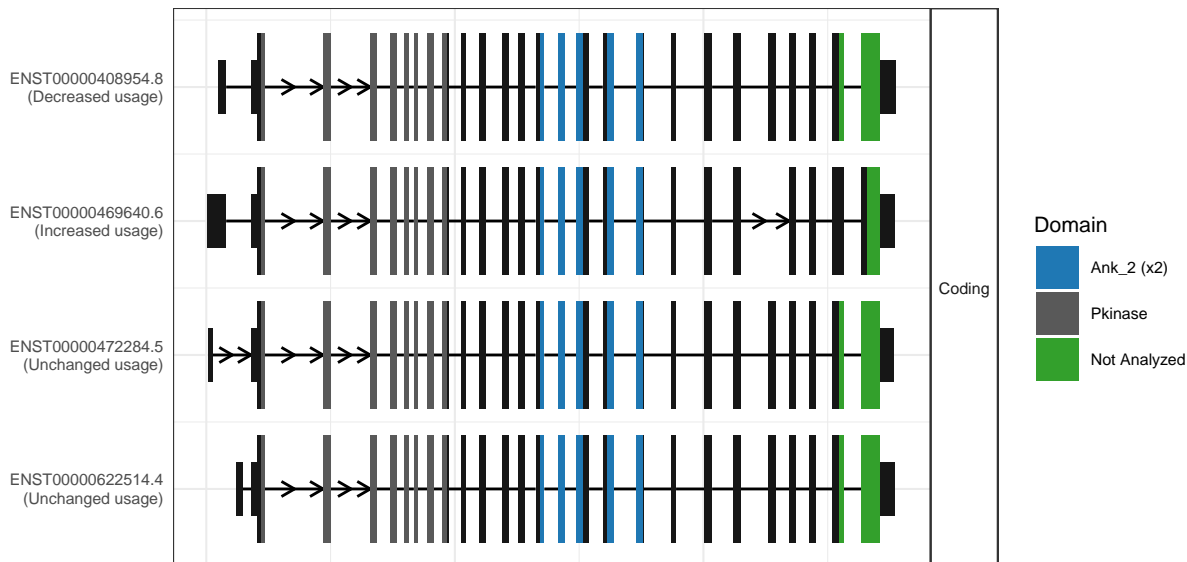


DNA_photolyase

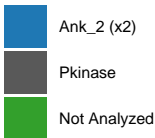


FAD_binding_7

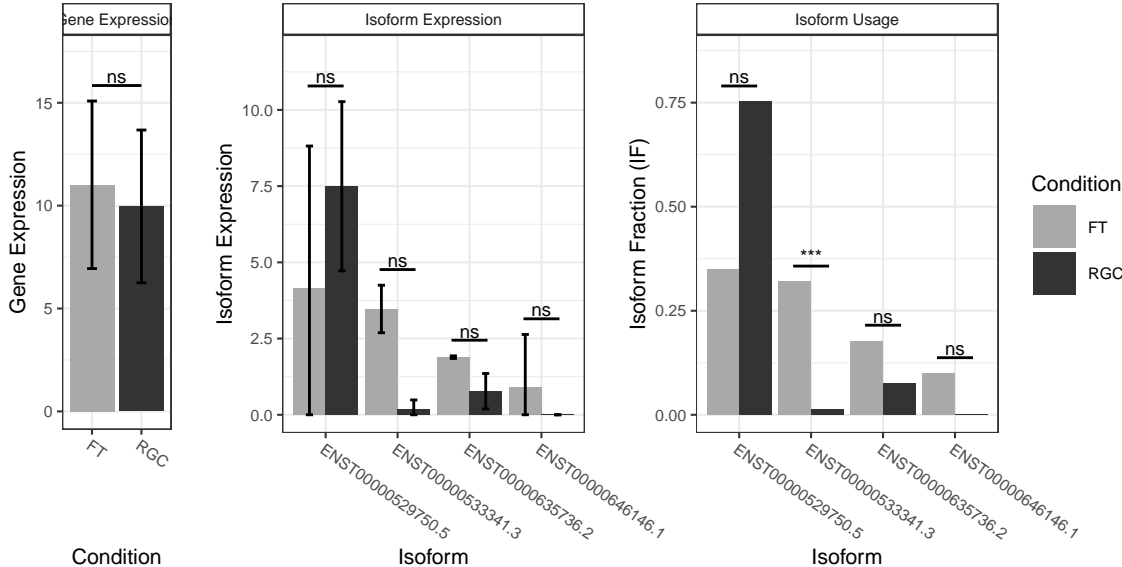
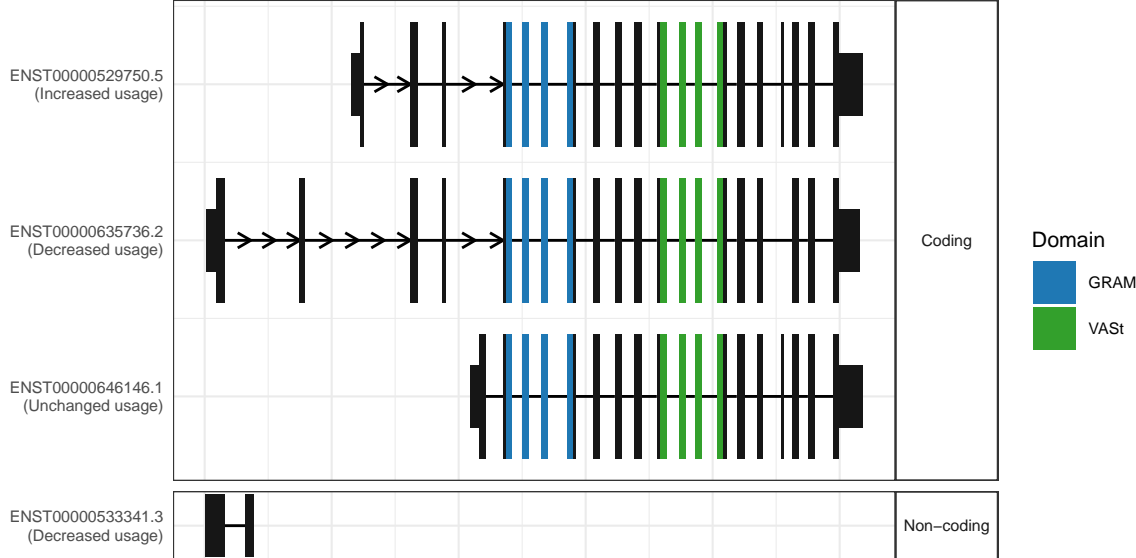
The isoform switch in DAPK1 (FT vs RGC)

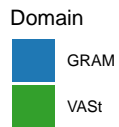


Domain

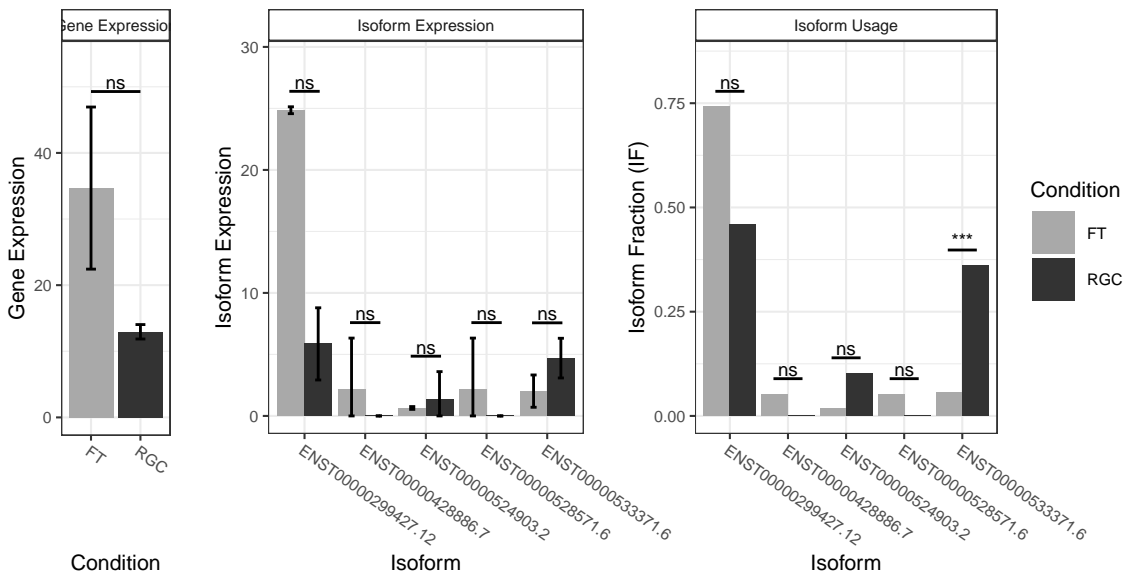
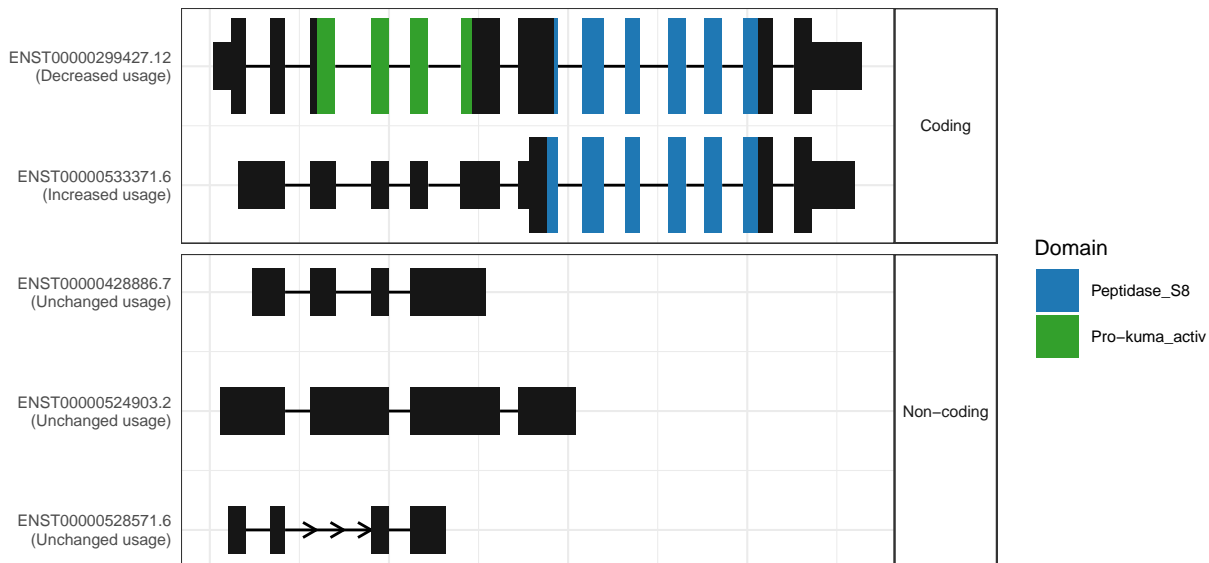


The isoform switch in GRAMD1B (FT vs RGC)





The isoform switch in TPP1 (FT vs RGC)



Domain

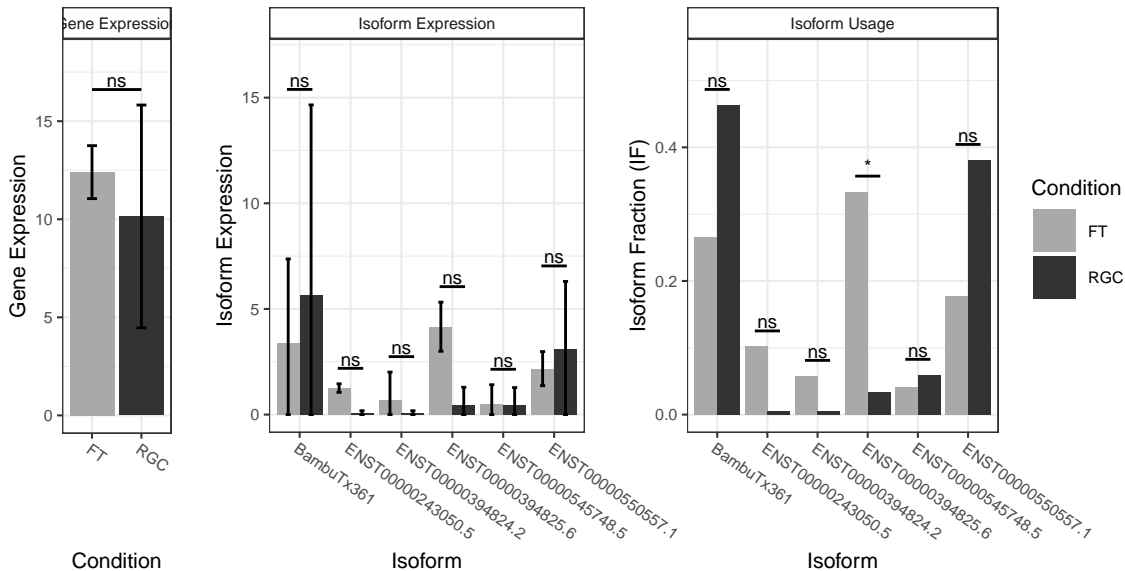
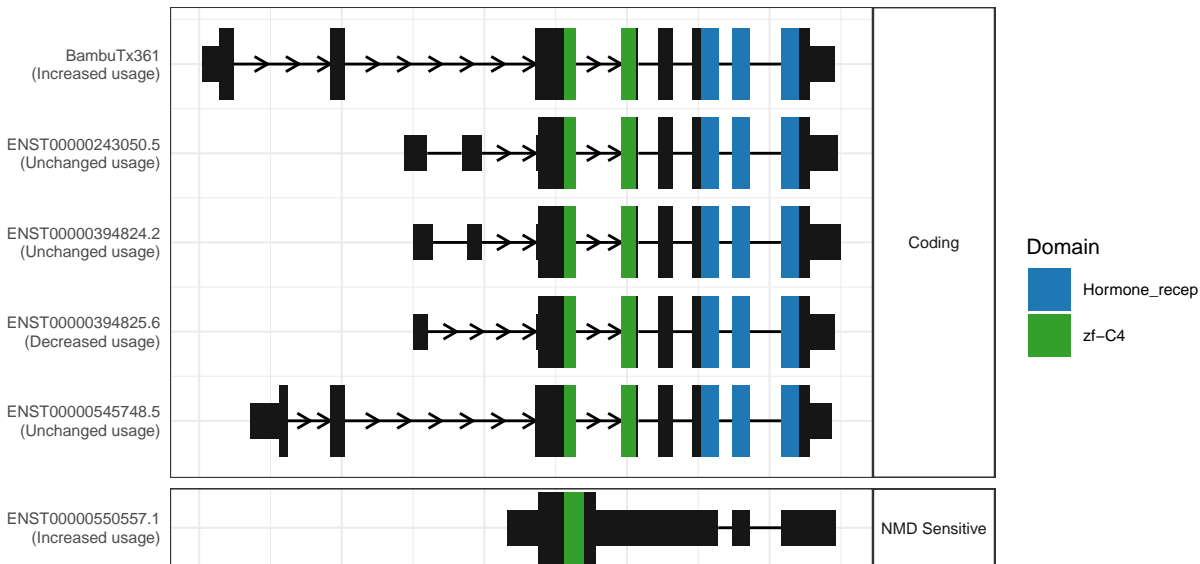


Peptidase_S8



Pro-kuma_activ

The isoform switch in NR4A1 (FT vs RGC)



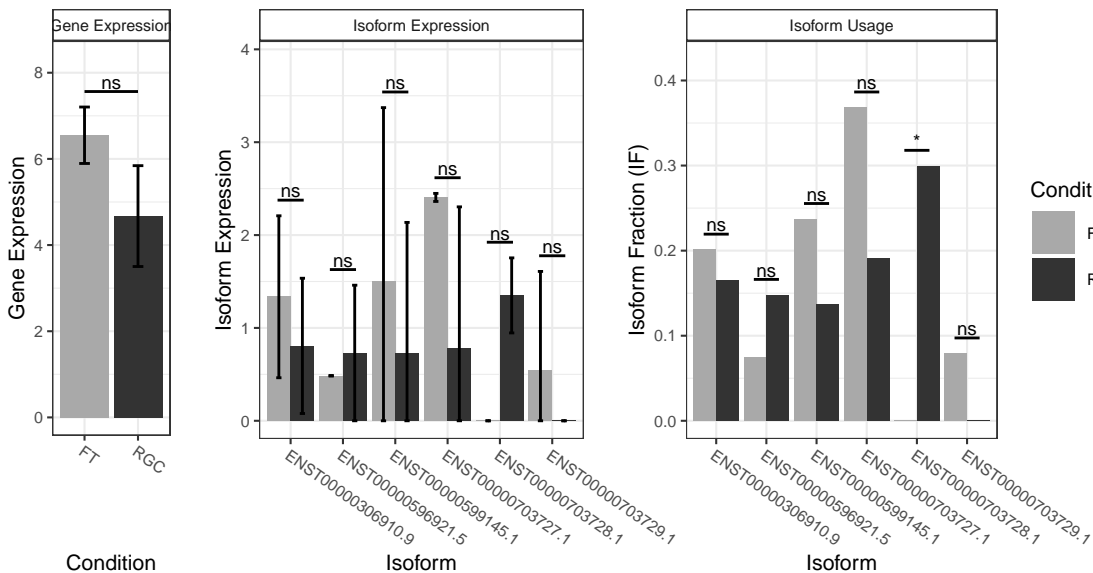
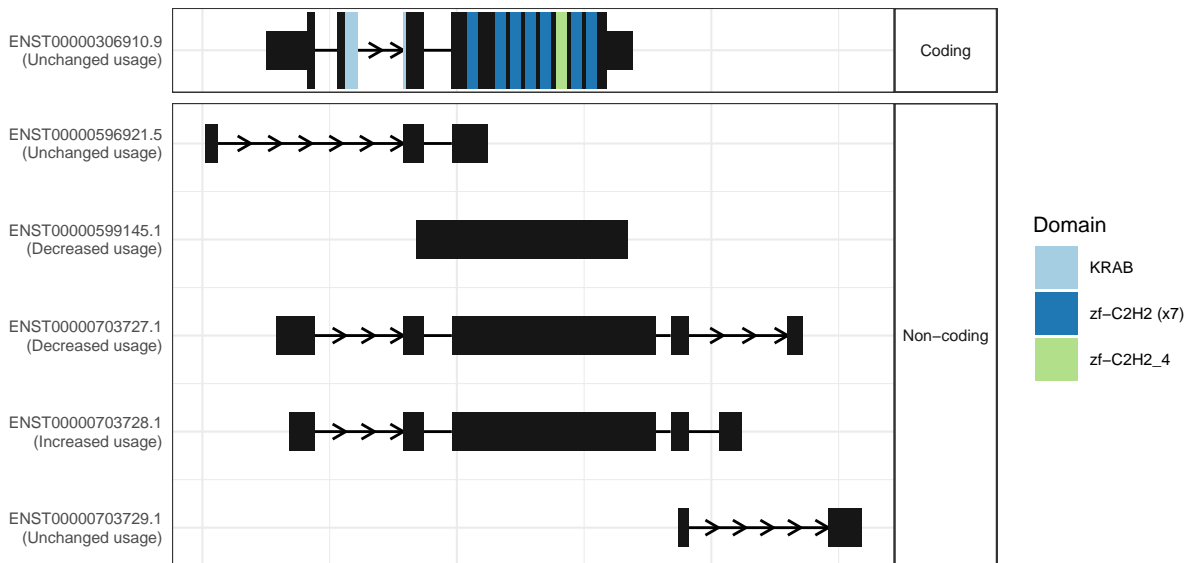
Domain



Hormone_recep

zf-C4

The isoform switch in ZNF584 (FT vs RGC)



Domain



KRAB

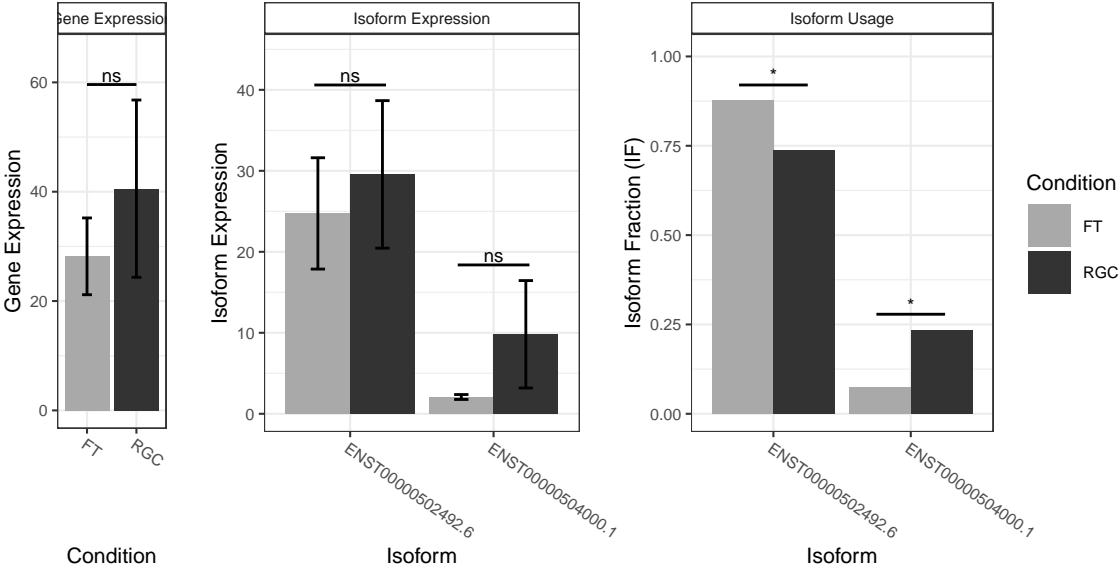
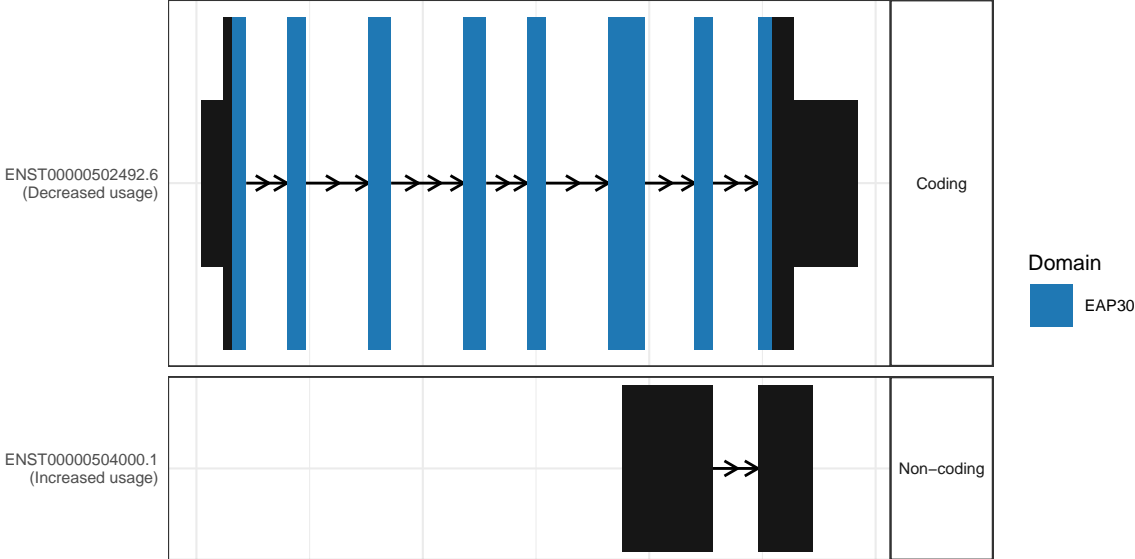


zf-C2H2 (x7)



zf-C2H2_4

The isoform switch in SNF8 (FT vs RGC)

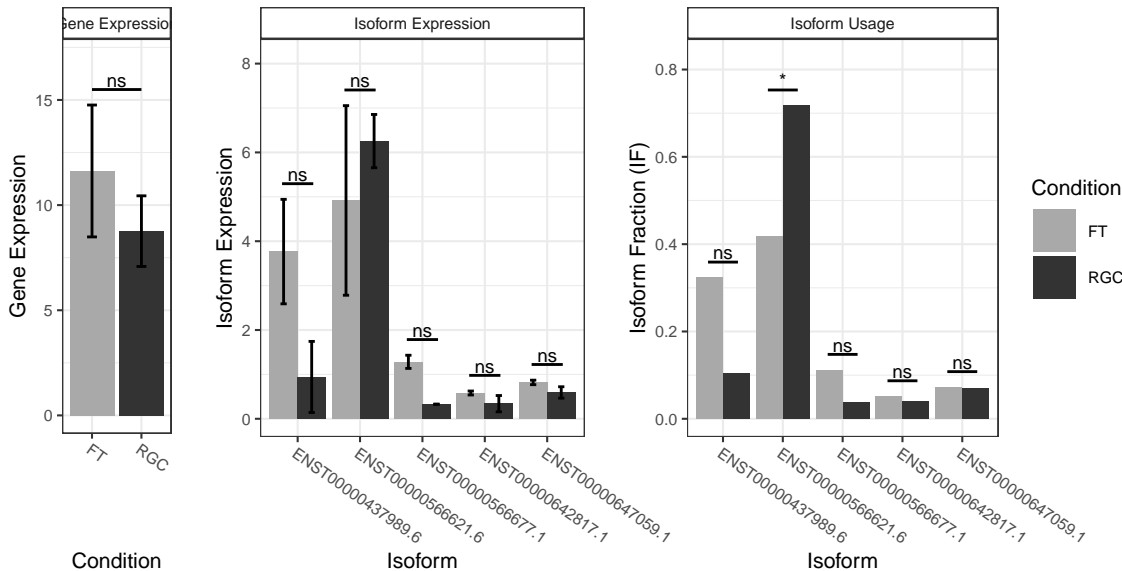
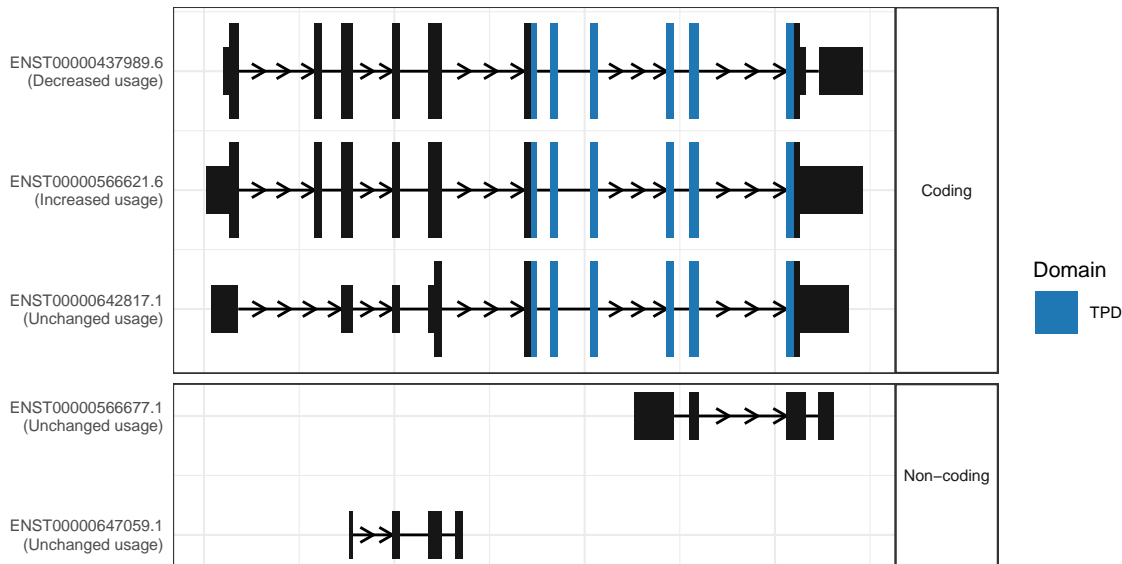


Domain



EAP30

The isoform switch in CDIN1 (FT vs RGC)

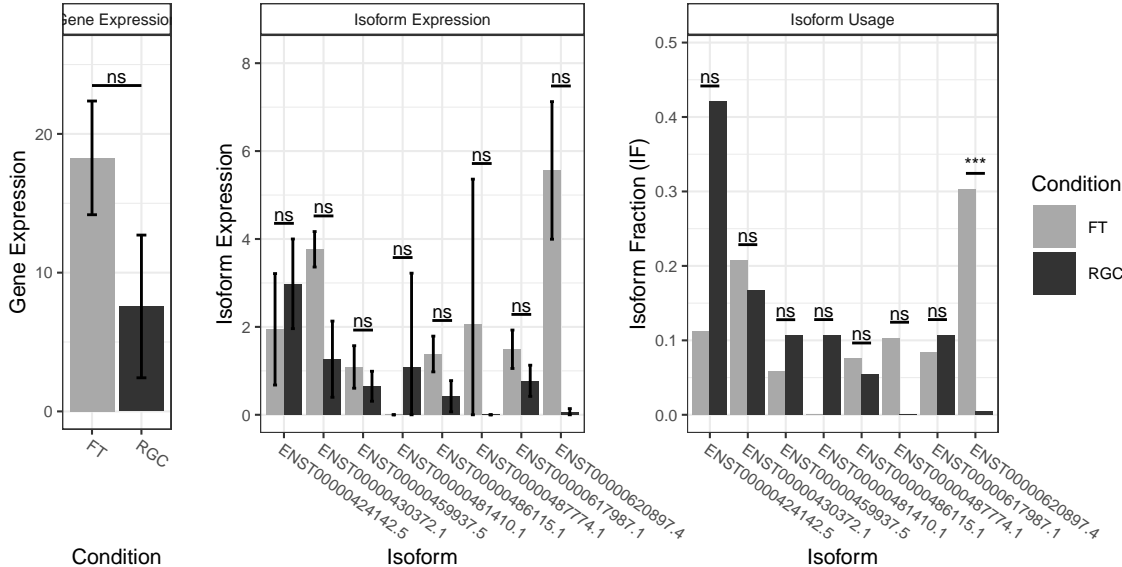
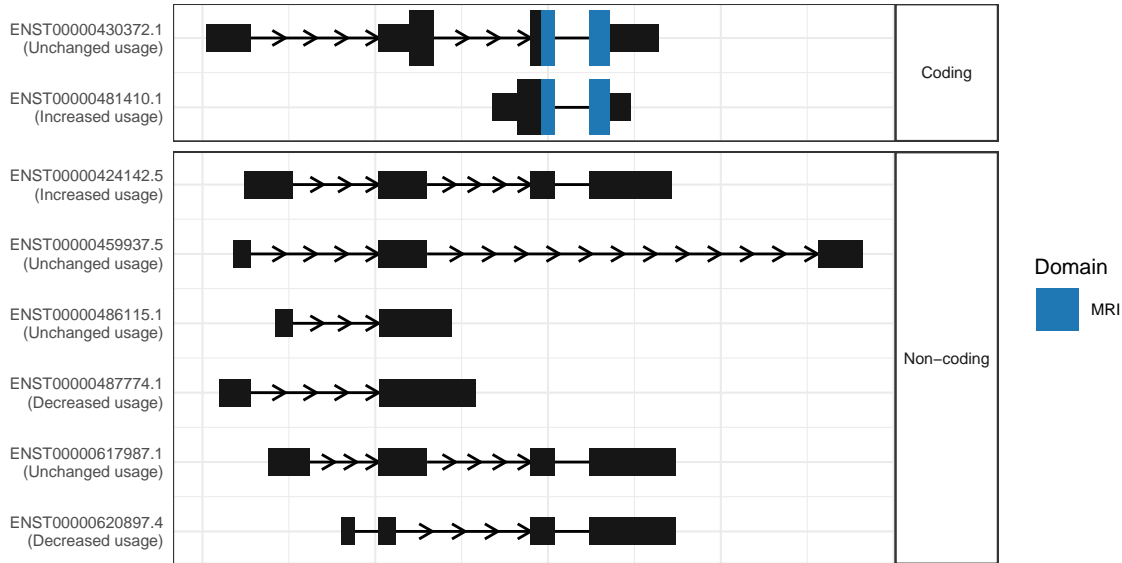


Domain



TPD

The isoform switch in CYREN (FT vs RGC)

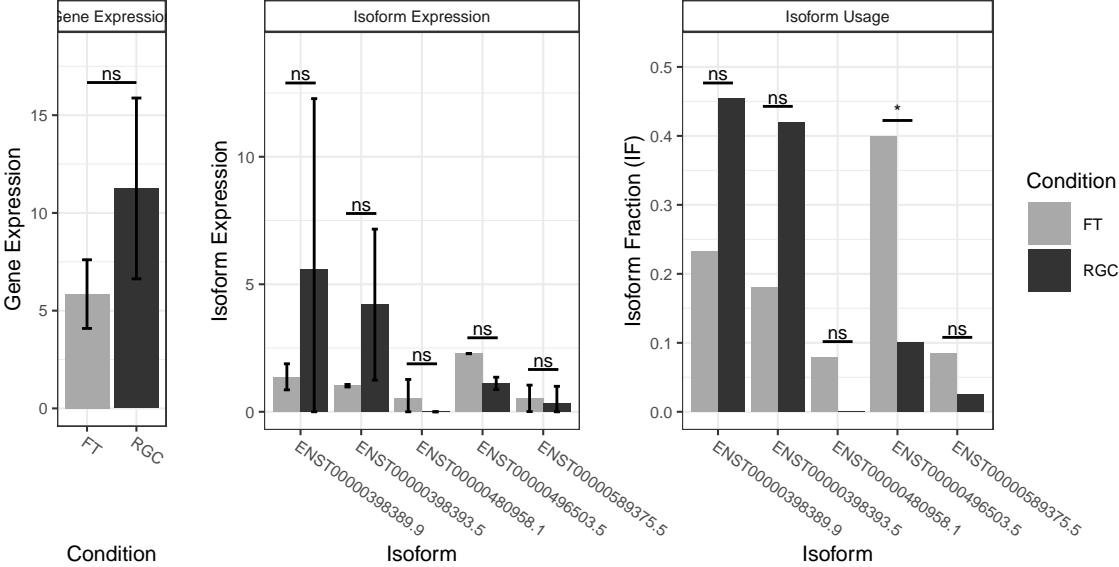
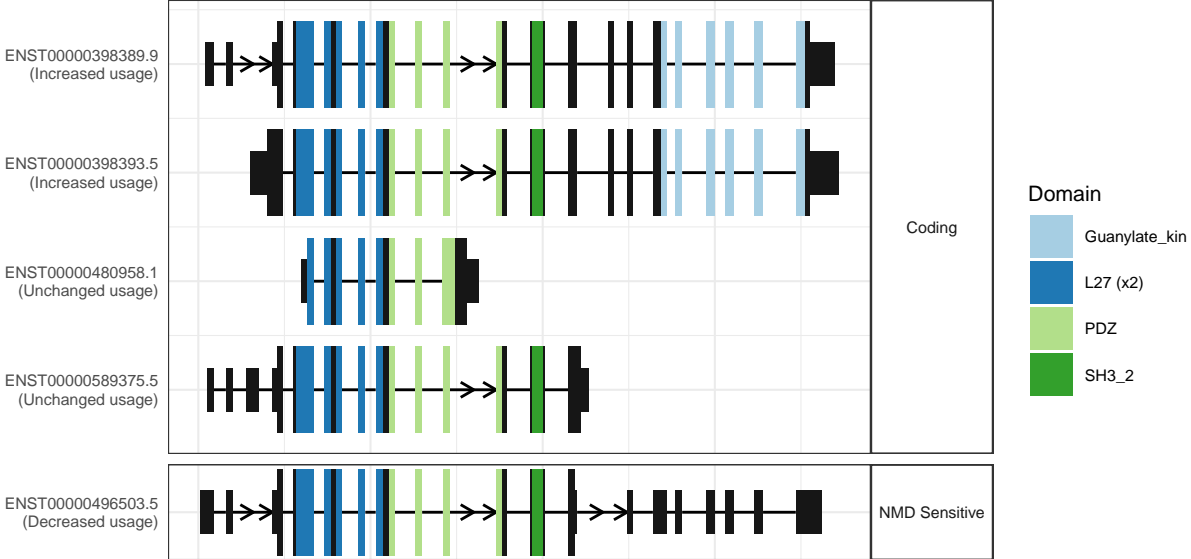


Domain







MRI

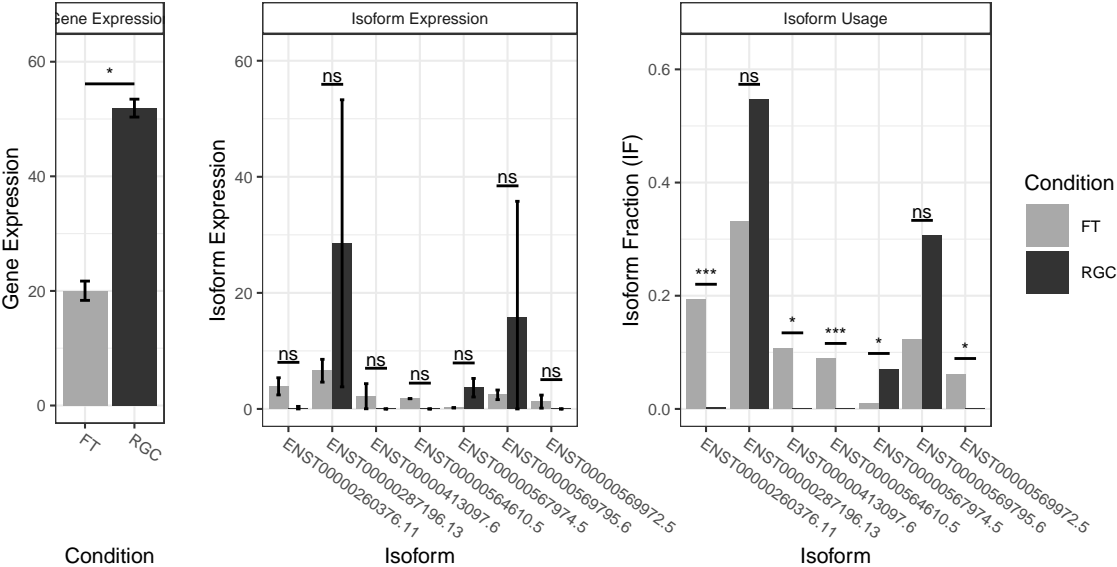
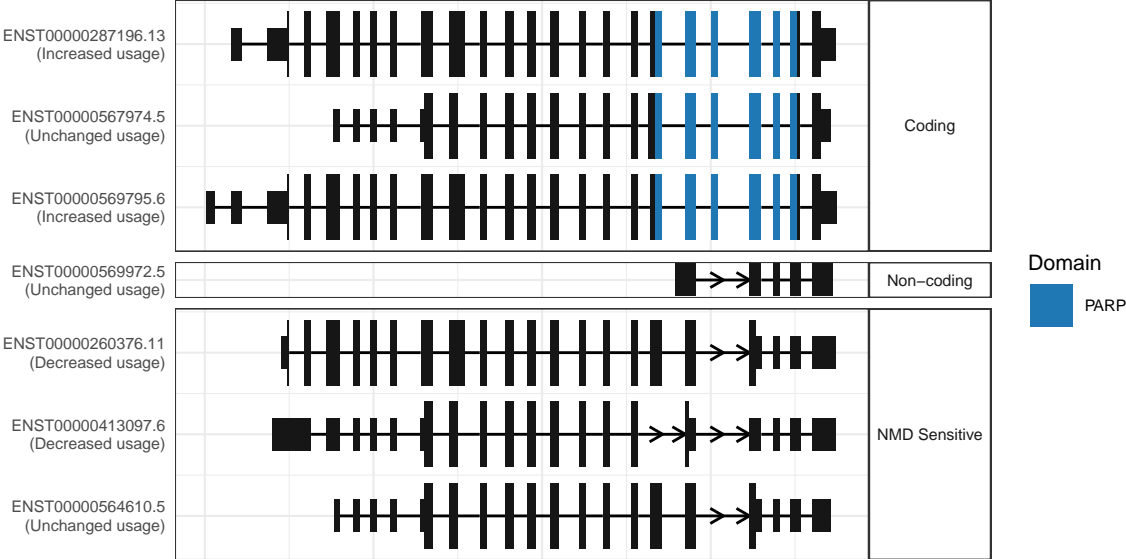
The isoform switch in MPP3 (FT vs RGC)



Domain

	Guanylate_kin
	L27 (x2)
	PDZ
	SH3_2

The isoform switch in PARP6 (FT vs RGC)

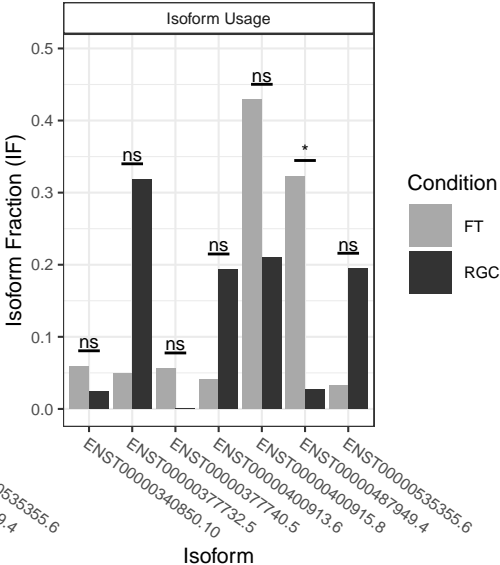
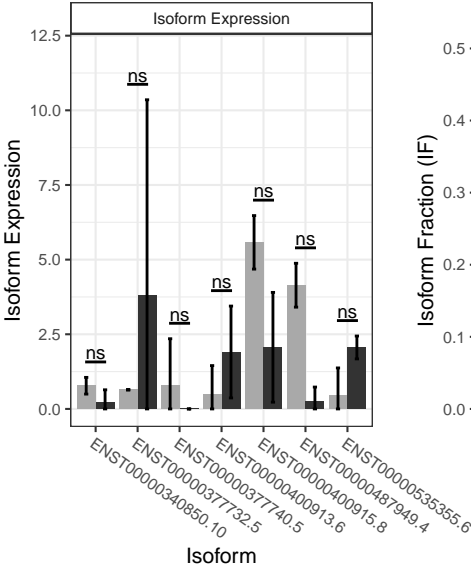
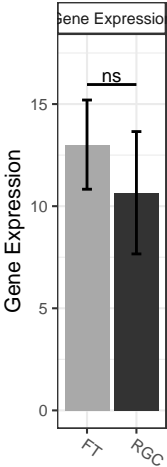
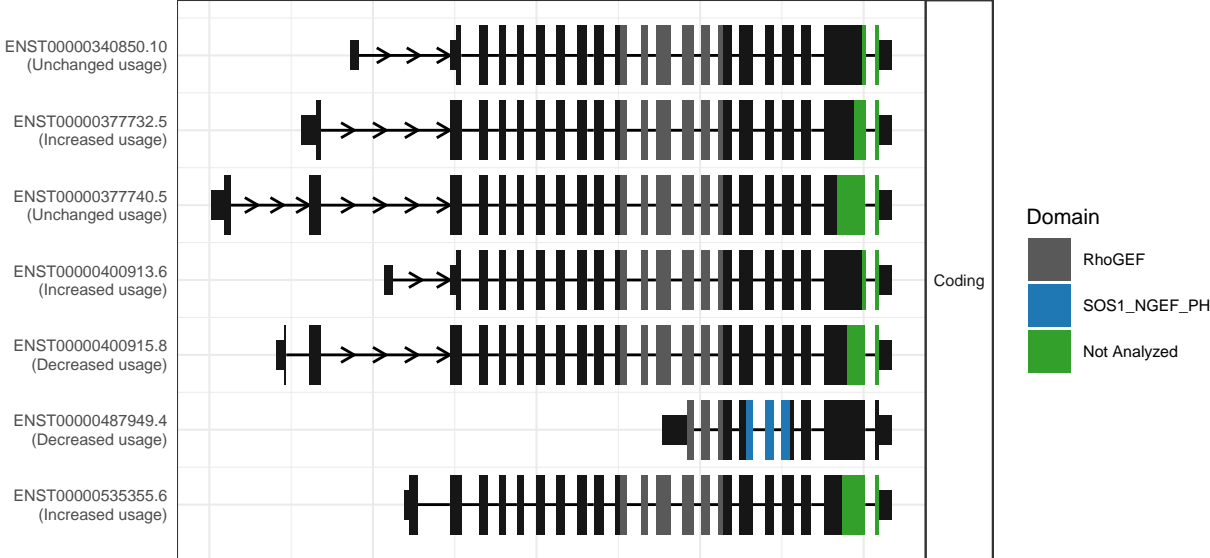


Domain



PARP

The isoform switch in PLEKHG5 (FT vs RGC)



Condition

Isoform

Isoform

Domain



RhoGEF

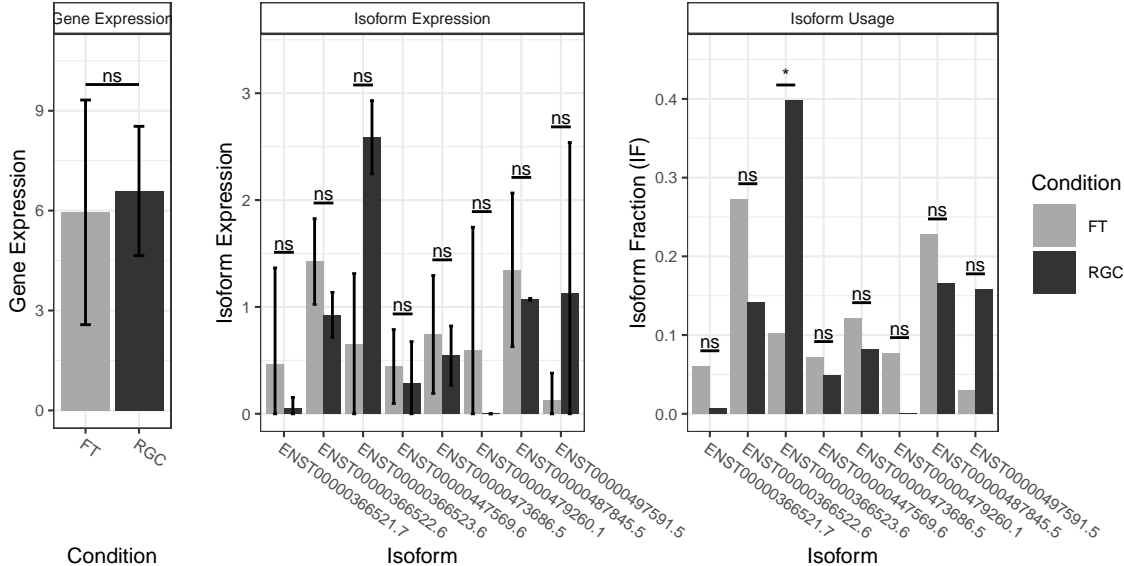
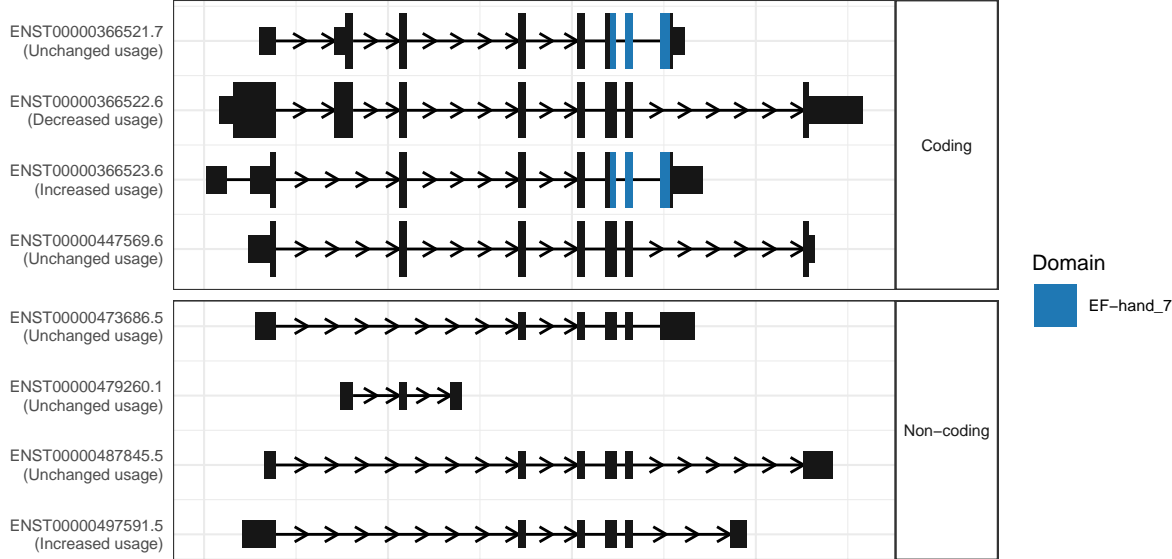


SOS1_NGEF_PH



Not Analyzed

The isoform switch in EFCAB2 (FT vs RGC)

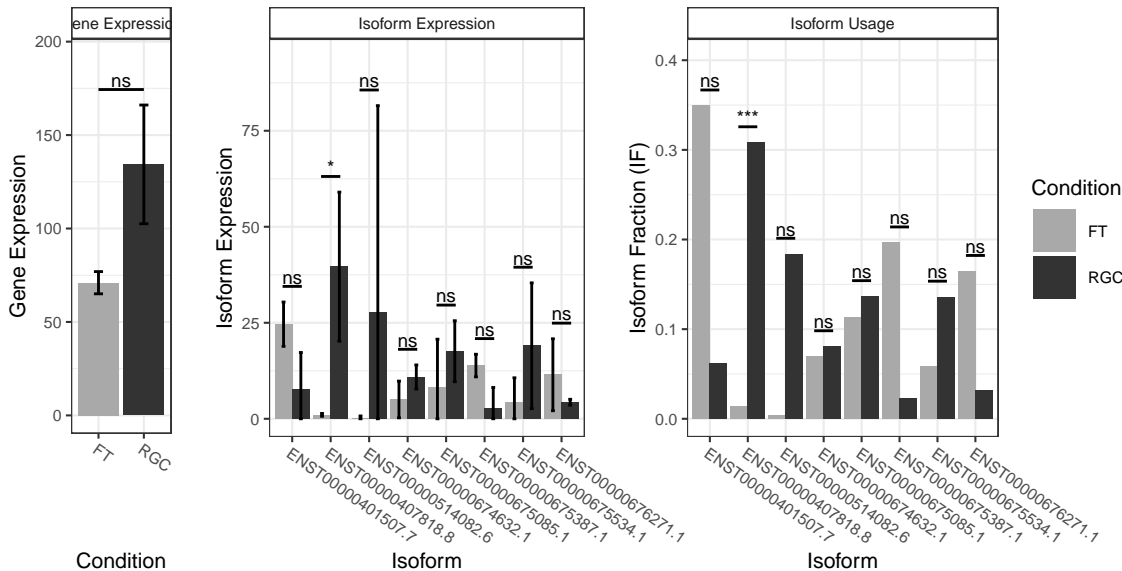
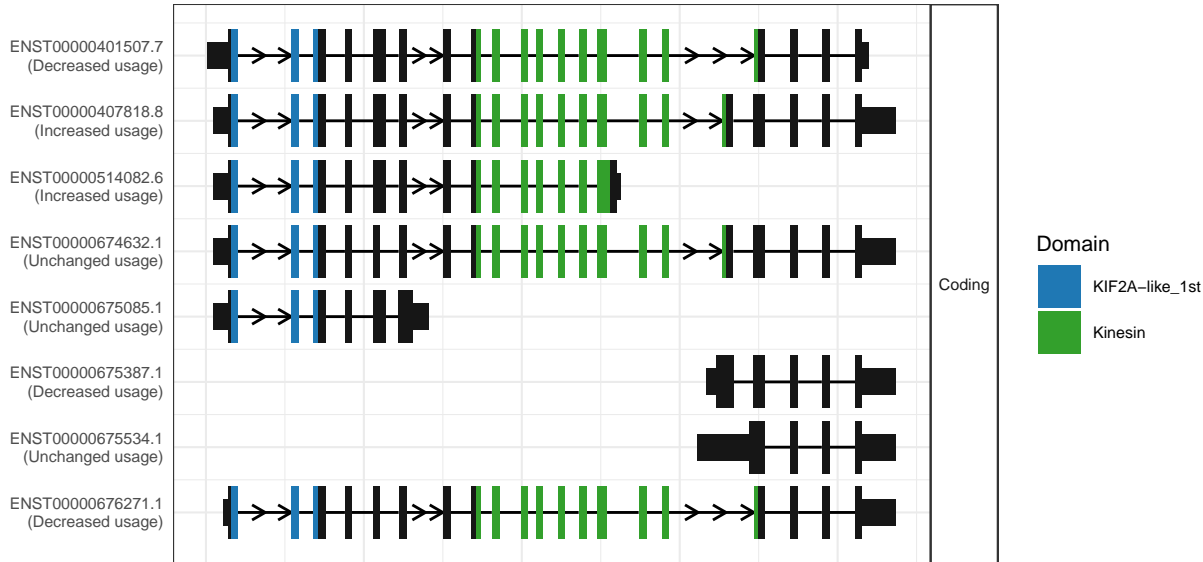


Domain



EF-hand_7

The isoform switch in KIF2A (FT vs RGC)



Domain

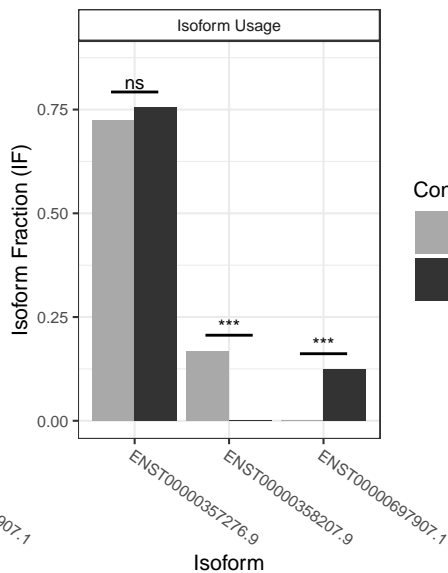
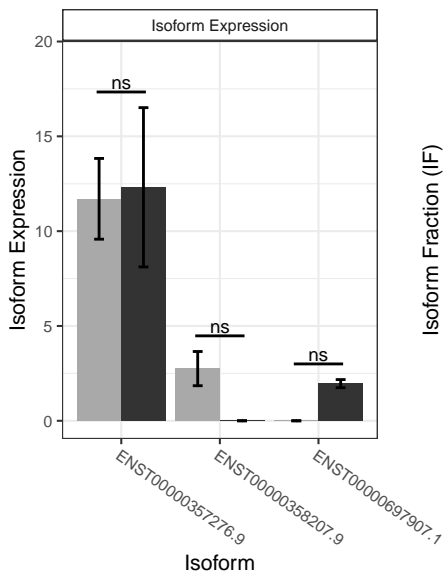
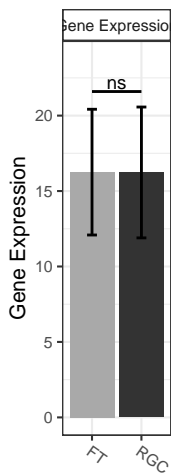
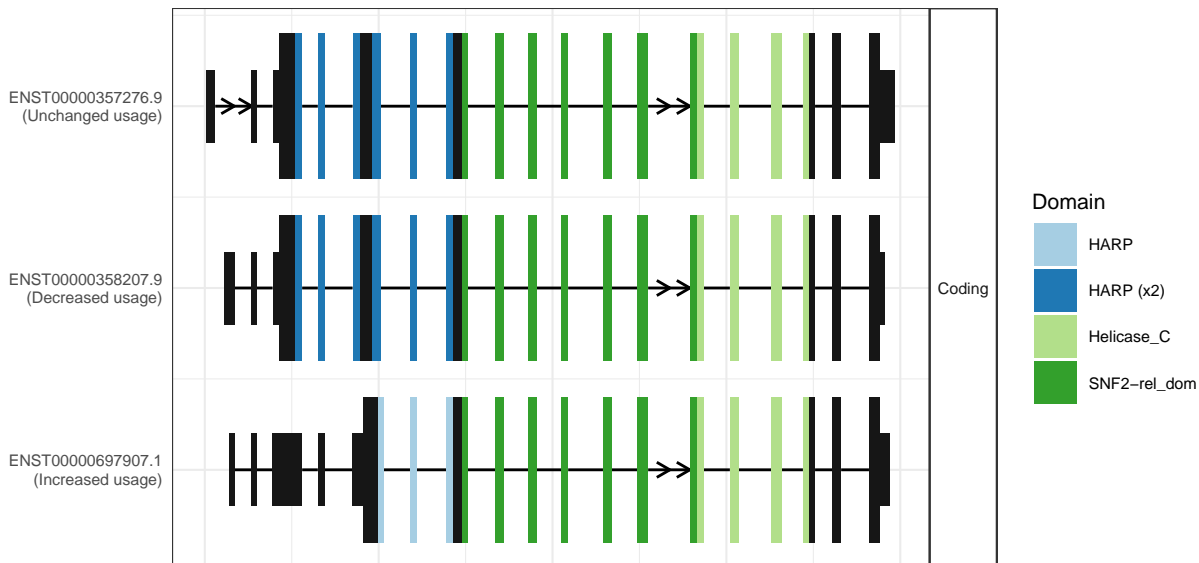


KIF2A-like_1st



Kinesin

The isoform switch in SMARCAL1 (FT vs RGC)



Domain



HARP



HARP (x2)

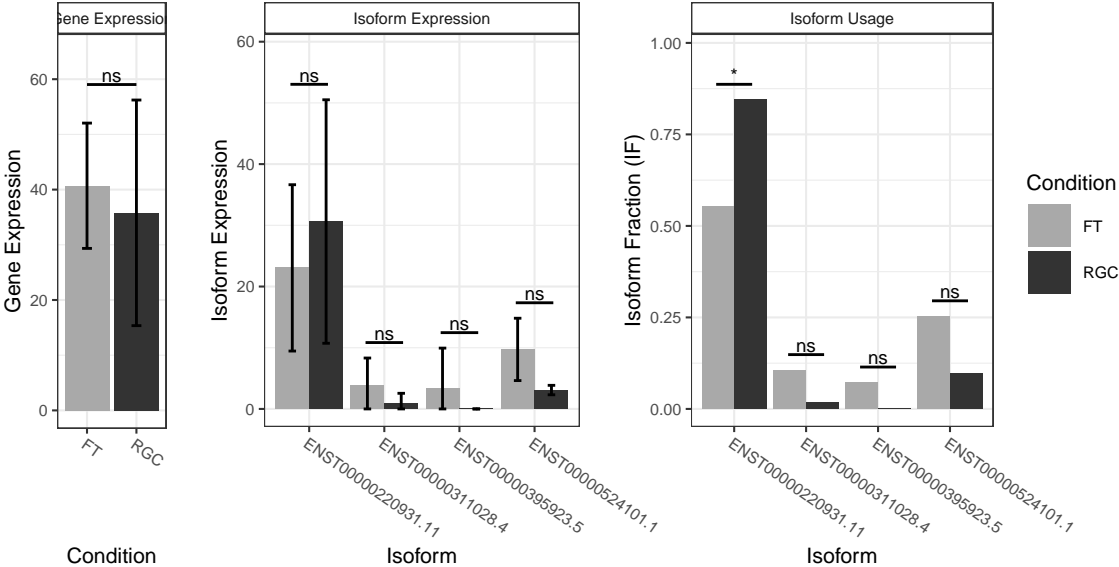
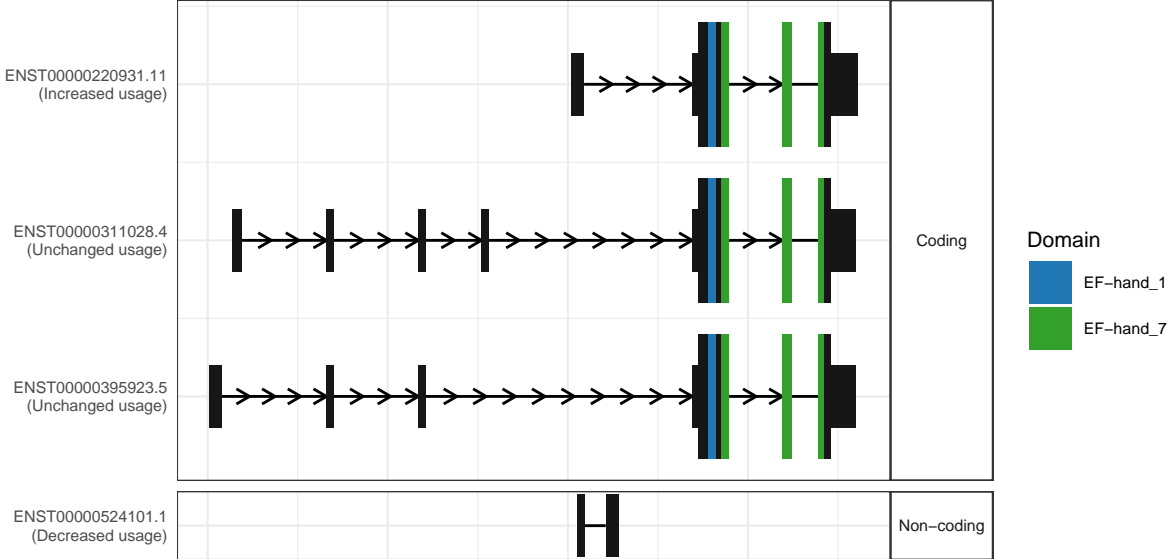


Helicase_C



SNF2-rel_dom

The isoform switch in NCALD (FT vs RGC)



Domain

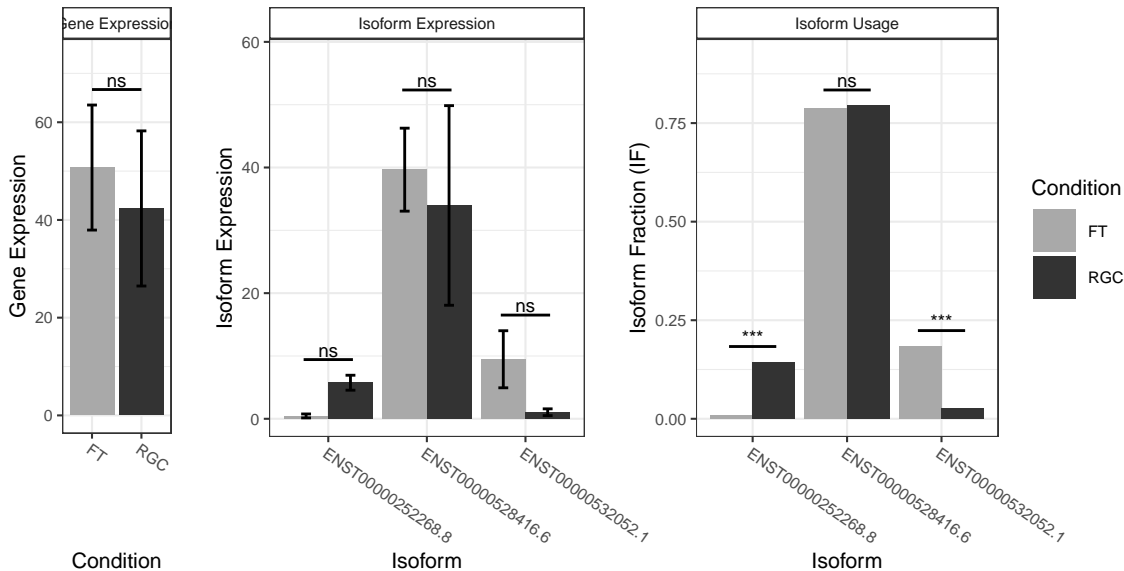
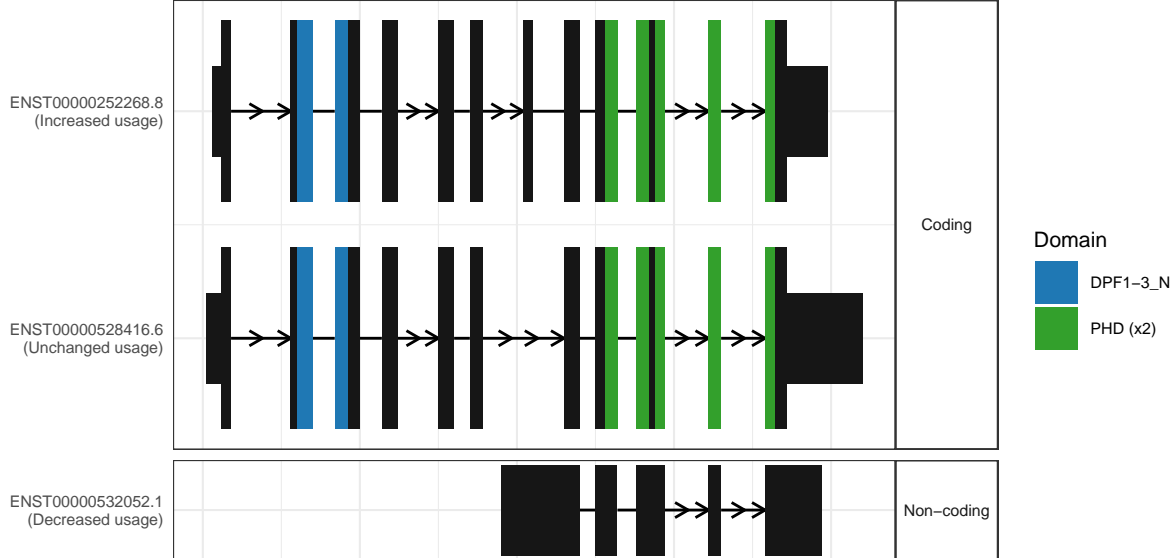


EF-hand_1



EF-hand_7

The isoform switch in DPF2 (FT vs RGC)



Domain

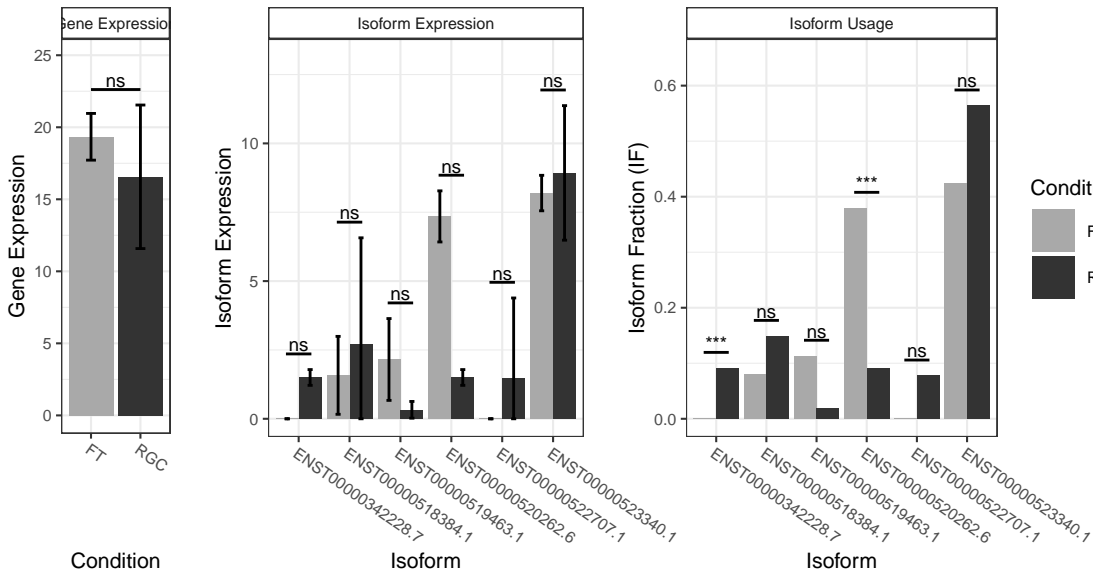
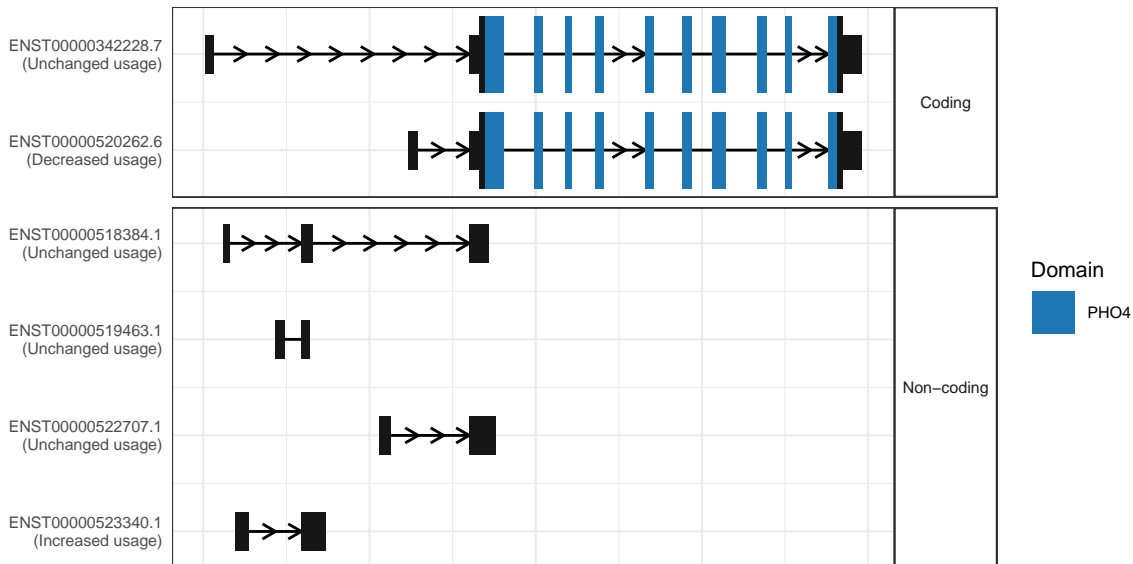


DPF1-3_N



PHD (x2)

The isoform switch in SLC20A2 (FT vs RGC)

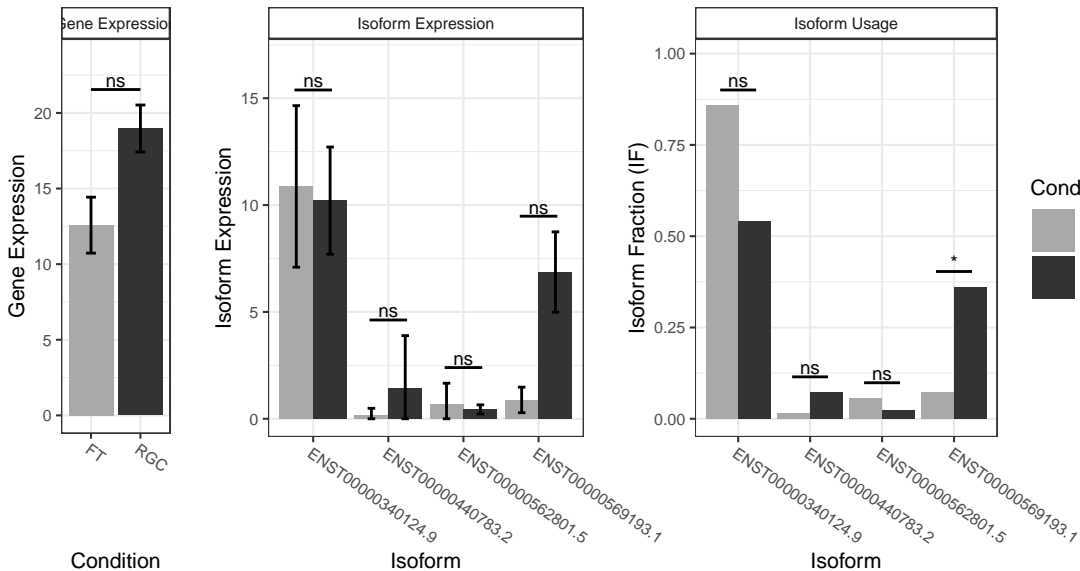
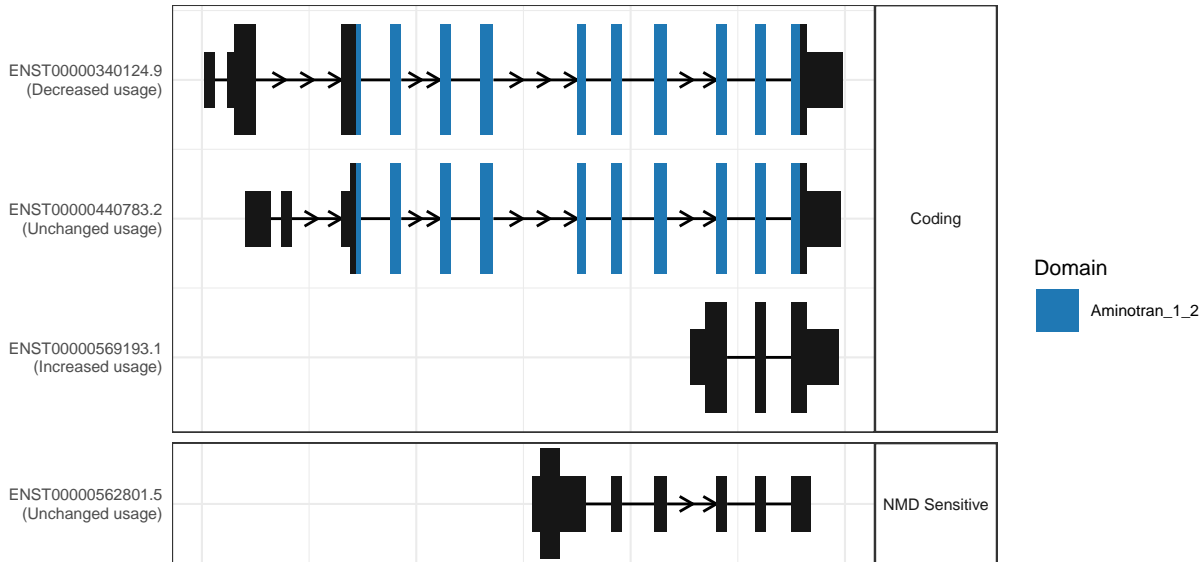


Domain



PHO4

The isoform switch in GPT2 (FT vs RGC)

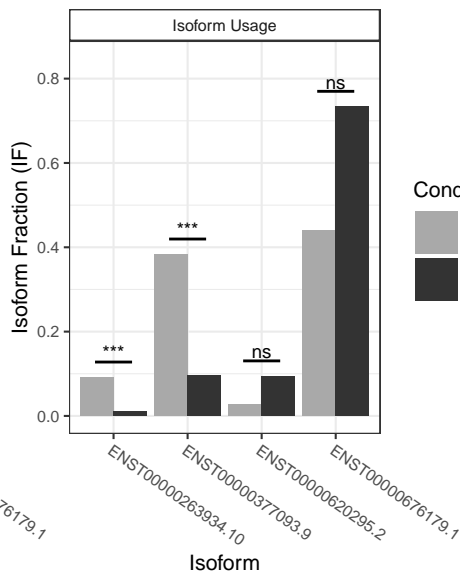
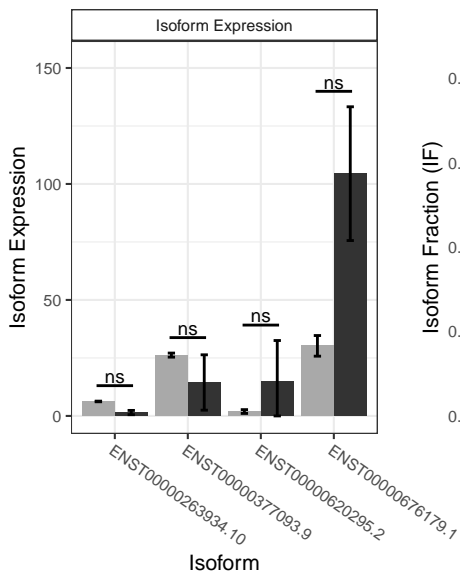
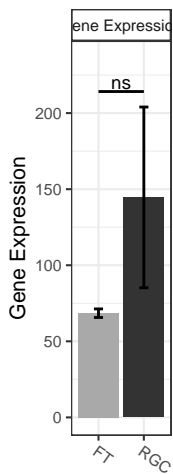
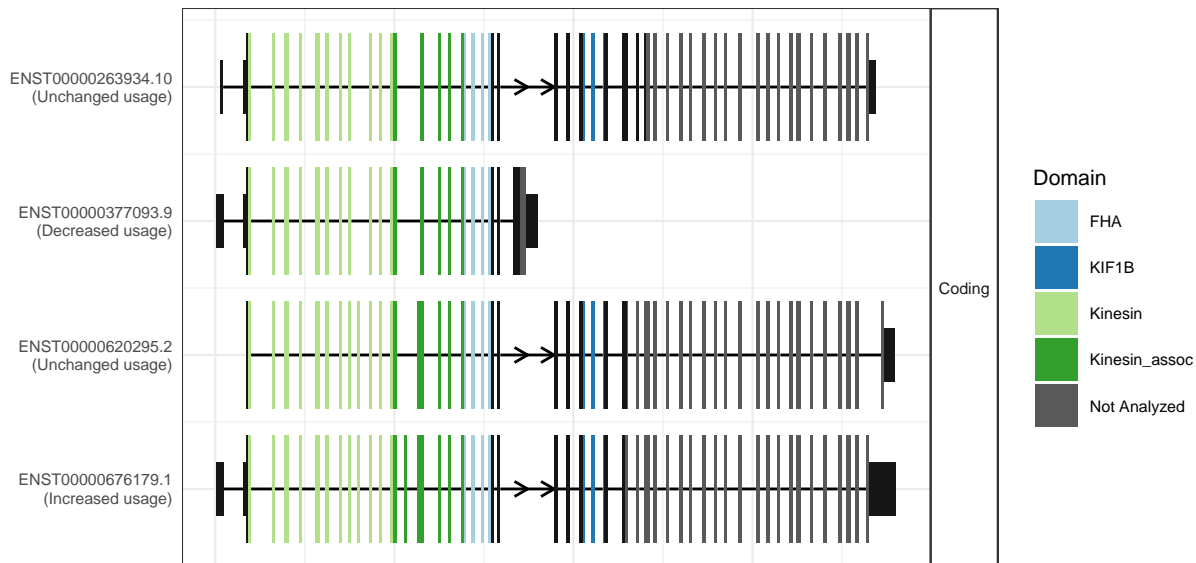


Domain





Aminotran_1_2

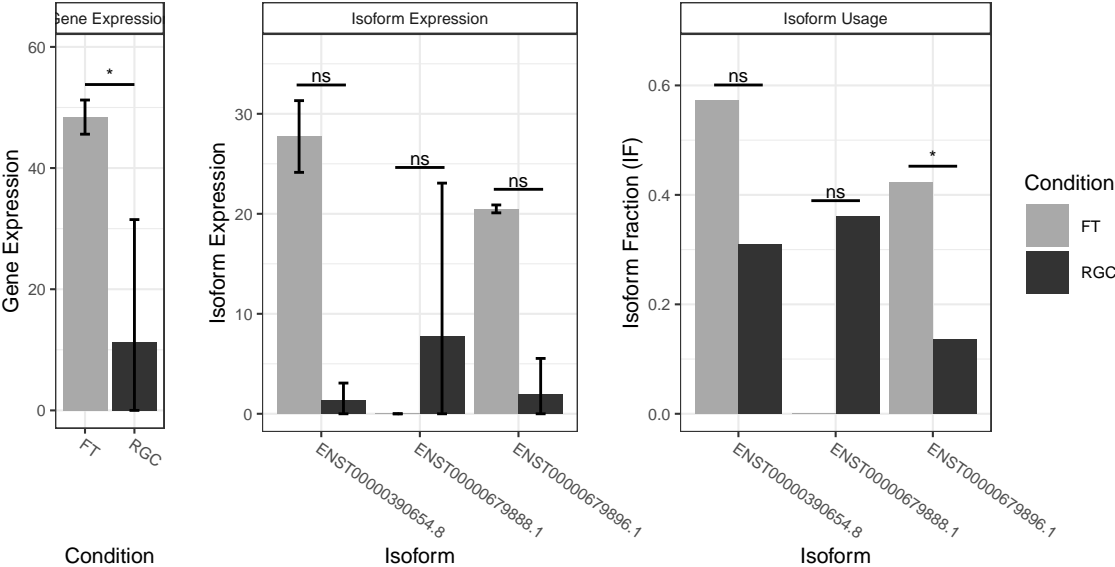
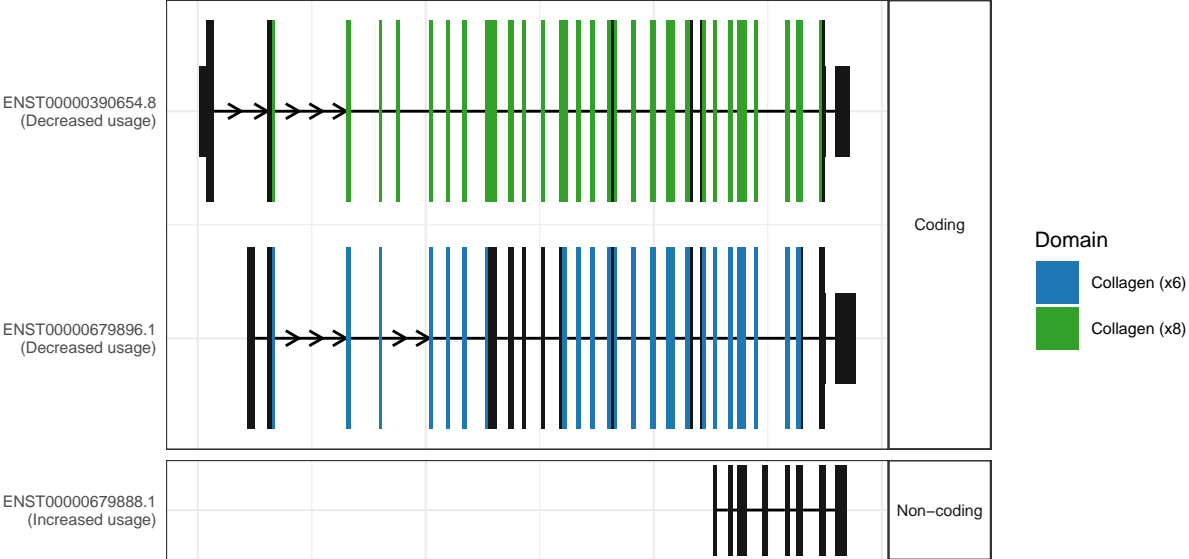
The isoform switch in KIF1B (FT vs RGC)



Domain

	FHA
	KIF1B
	Kinesin
	Kinesin_assoc
	Not Analyzed

The isoform switch in COL23A1 (FT vs RGC)



Domain

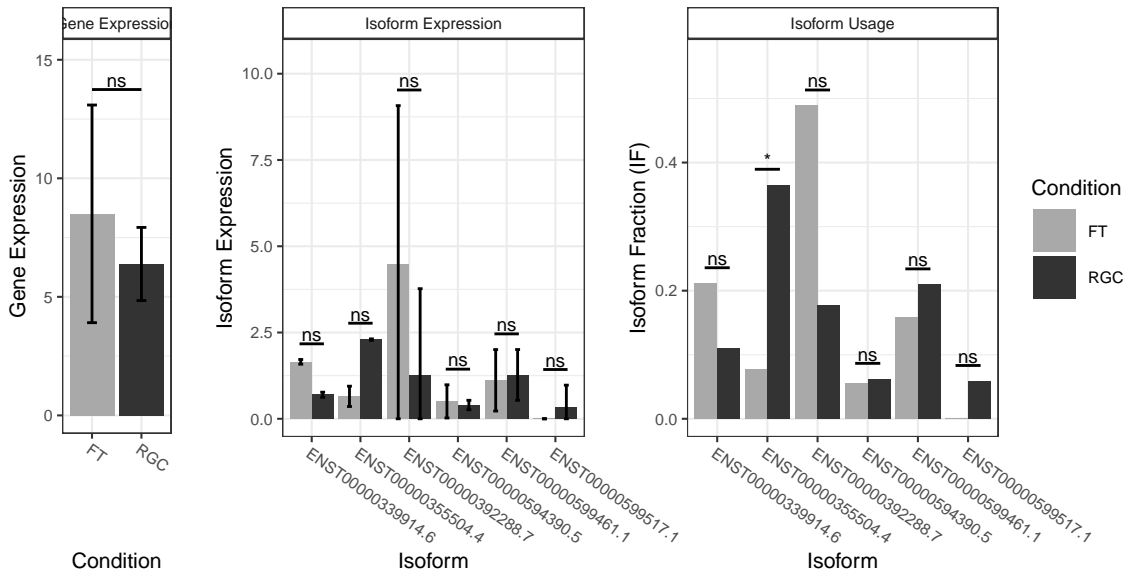
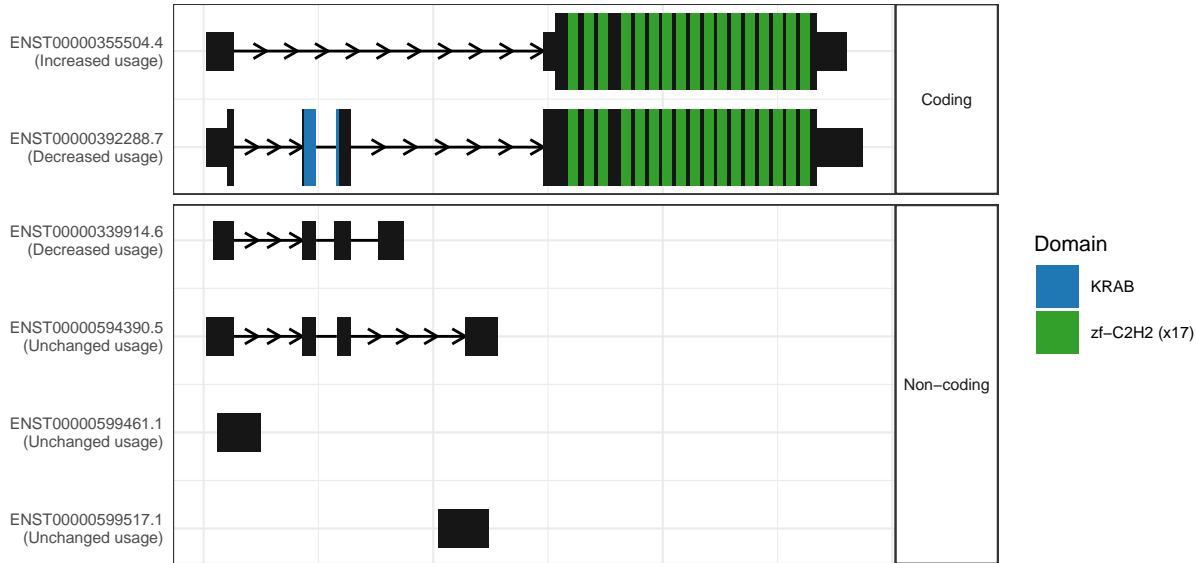


Collagen (x6)



Collagen (x8)

The isoform switch in ZNF493 (FT vs RGC)



Domain

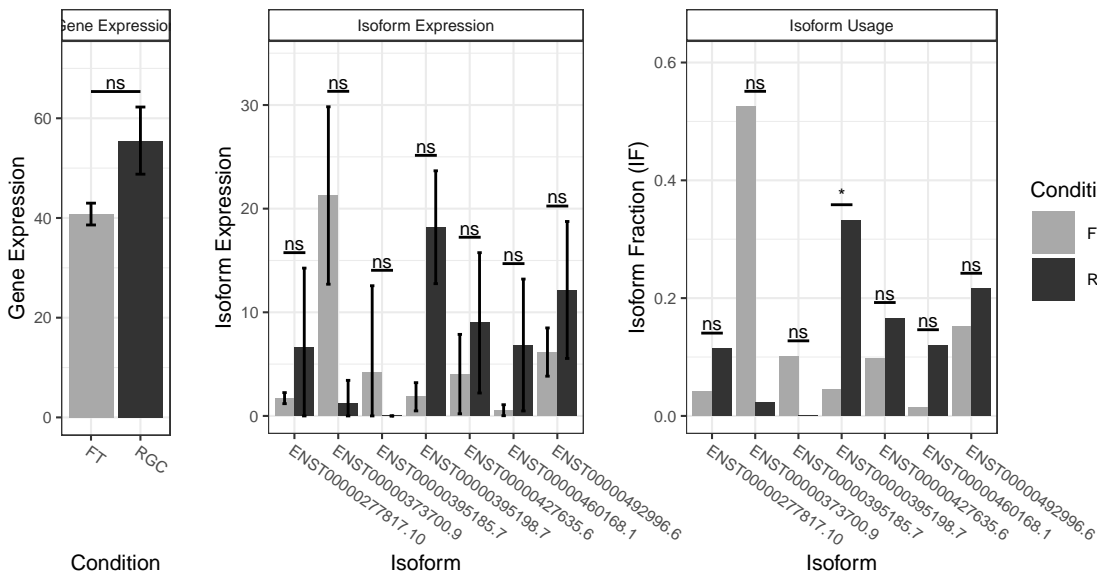
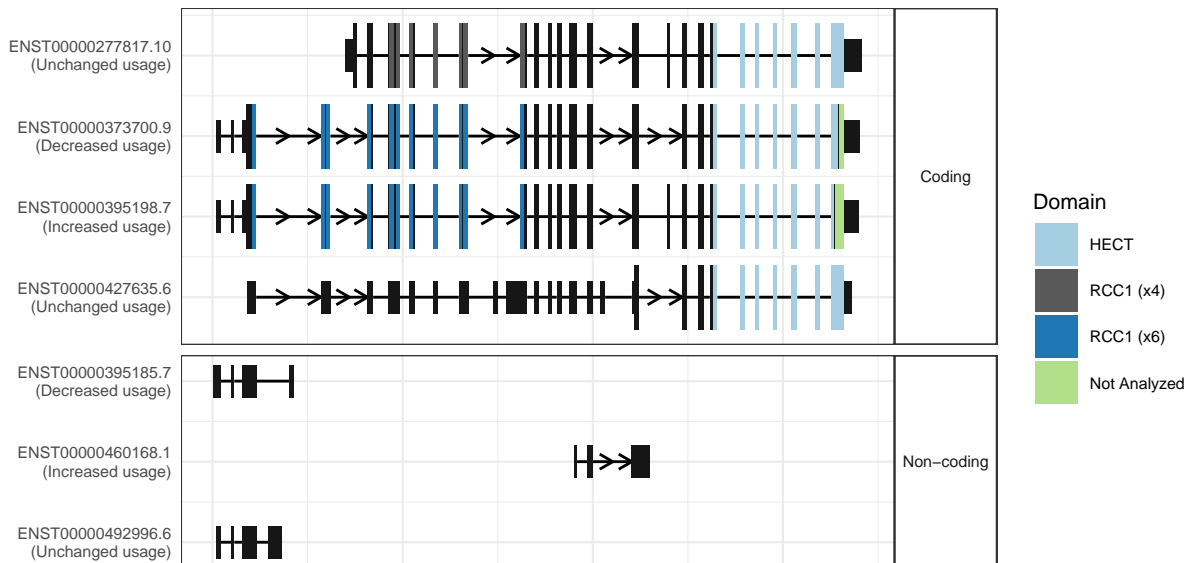


KRAB

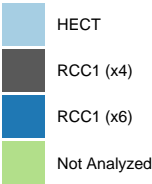


zf-C2H2 (x17)

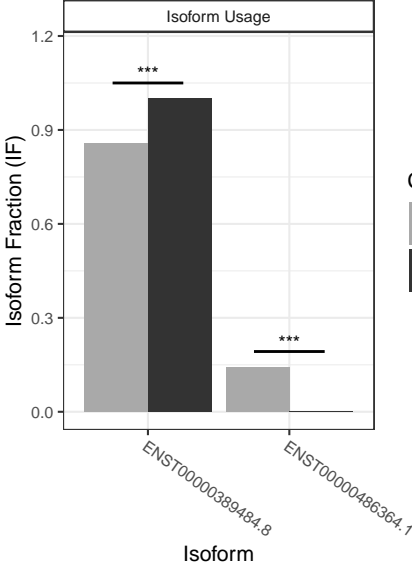
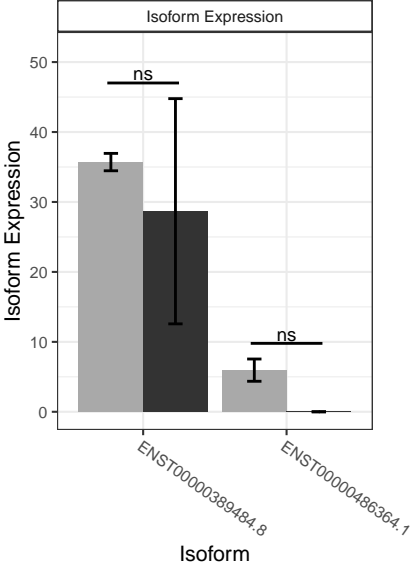
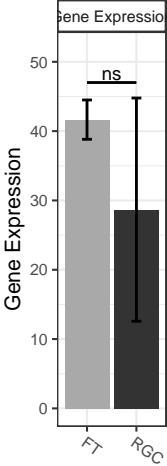
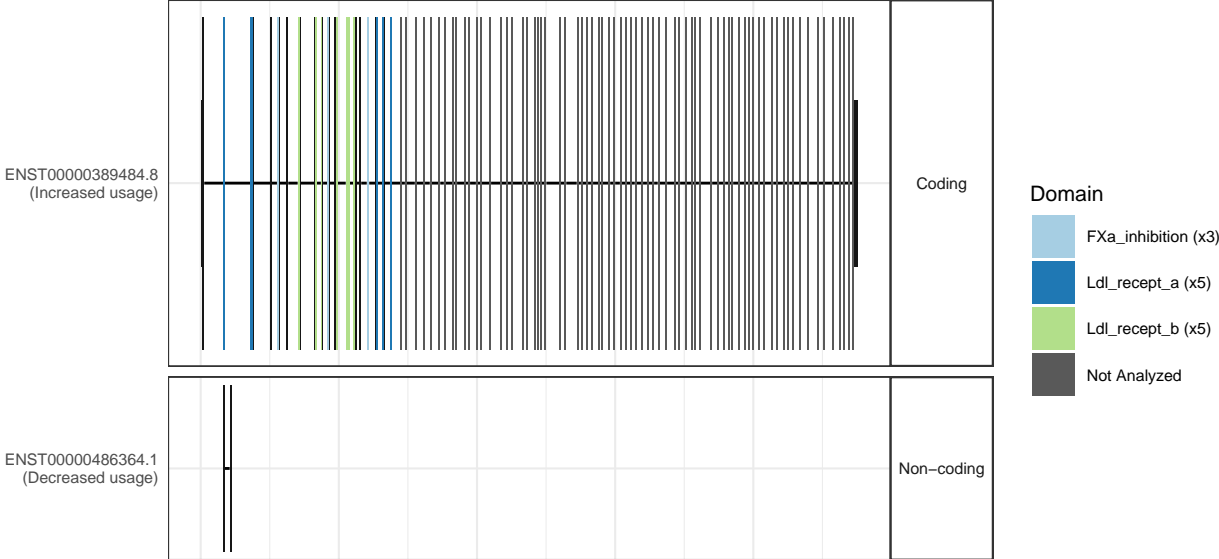
The isoform switch in HERC4 (FT vs RGC)



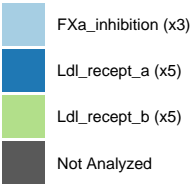
Domain



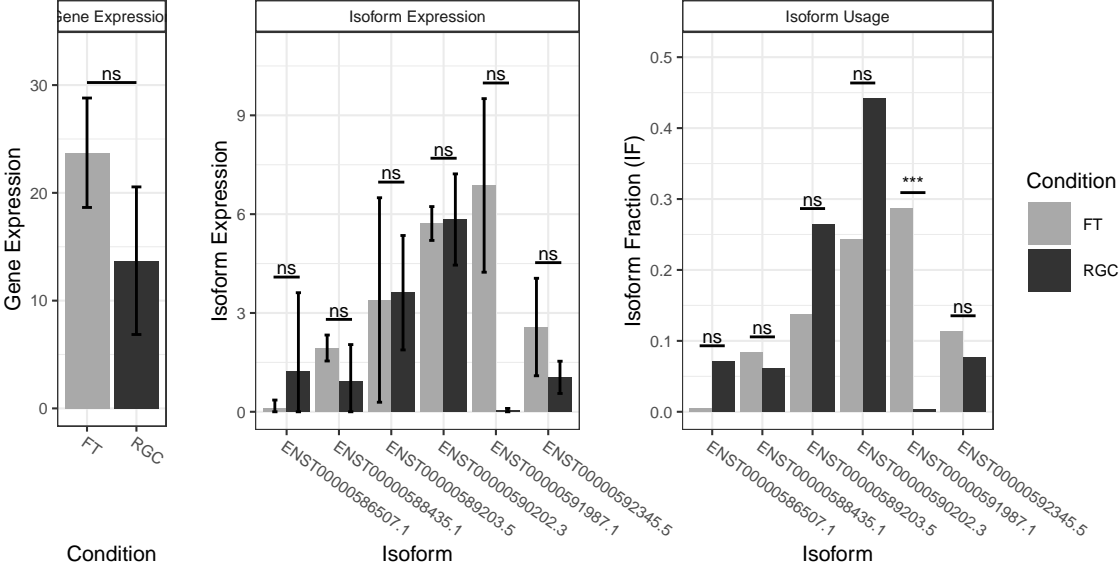
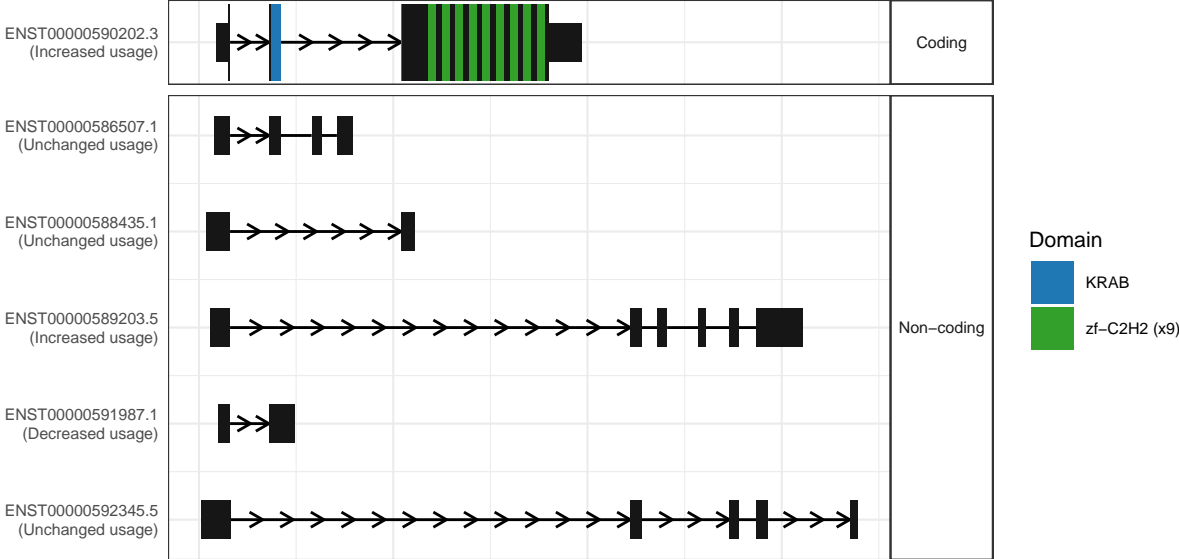
The isoform switch in LRP1B (FT vs RGC)



Domain



The isoform switch in ZNF519 (FT vs RGC)



Domain

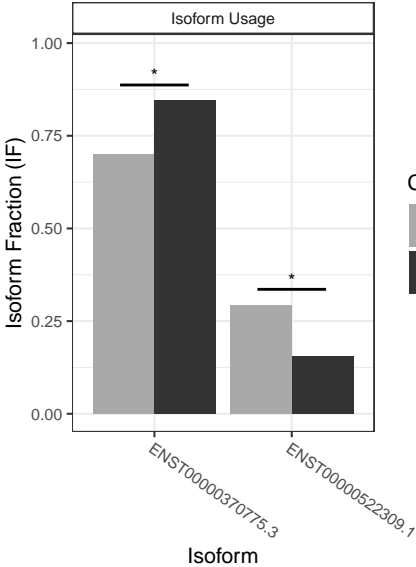
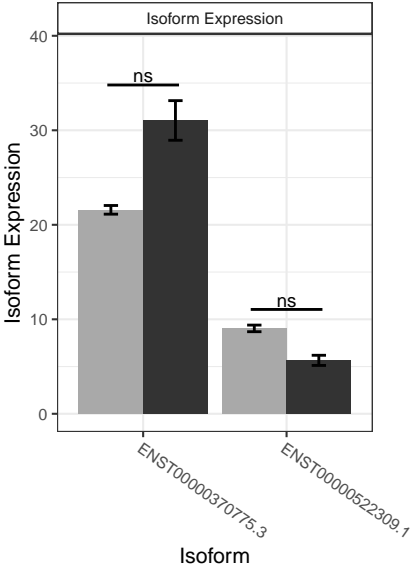
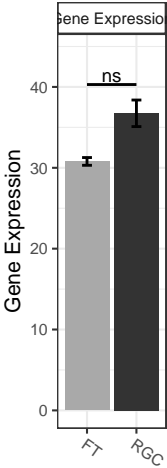
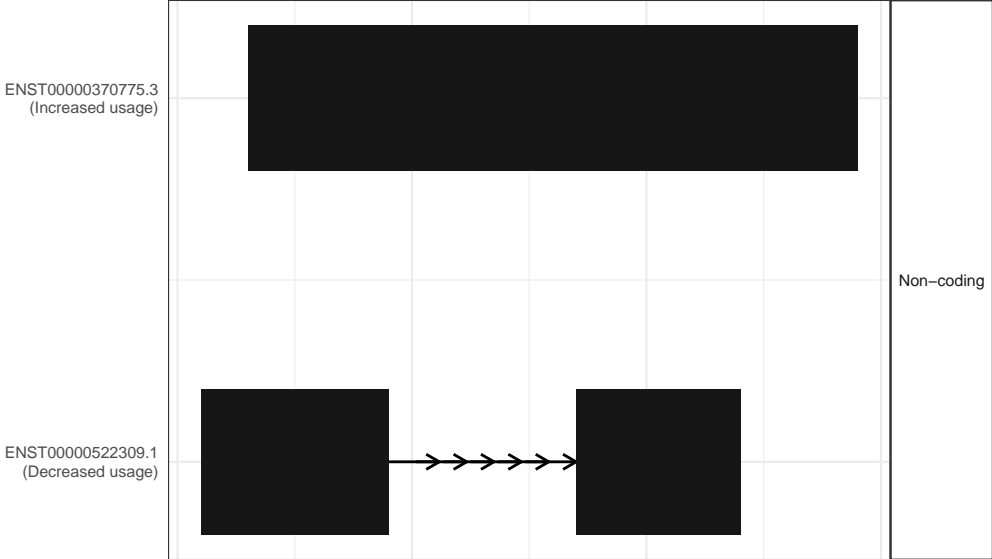


KRAB

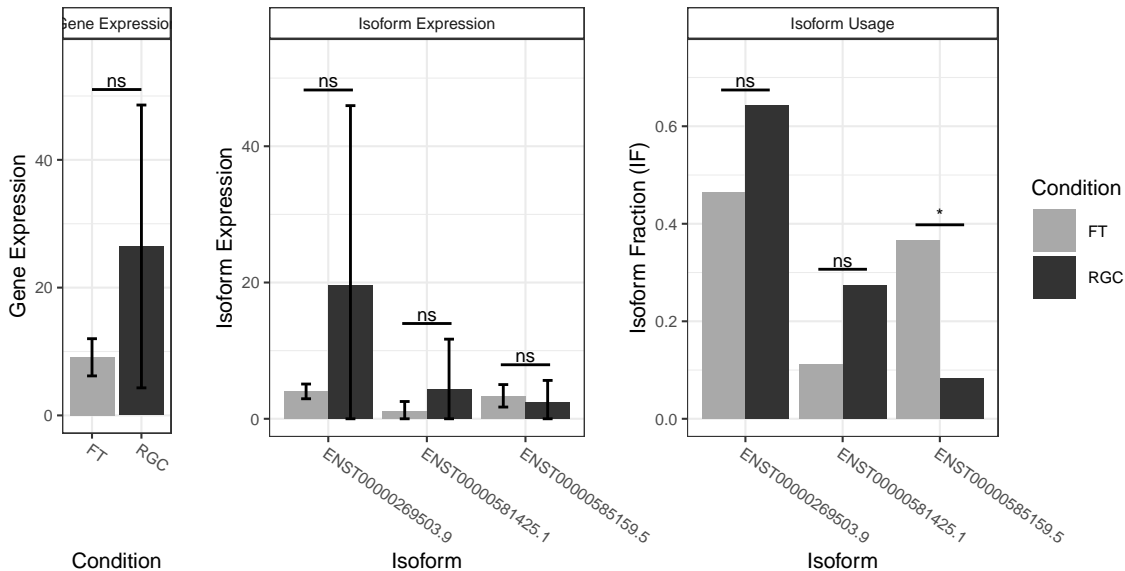
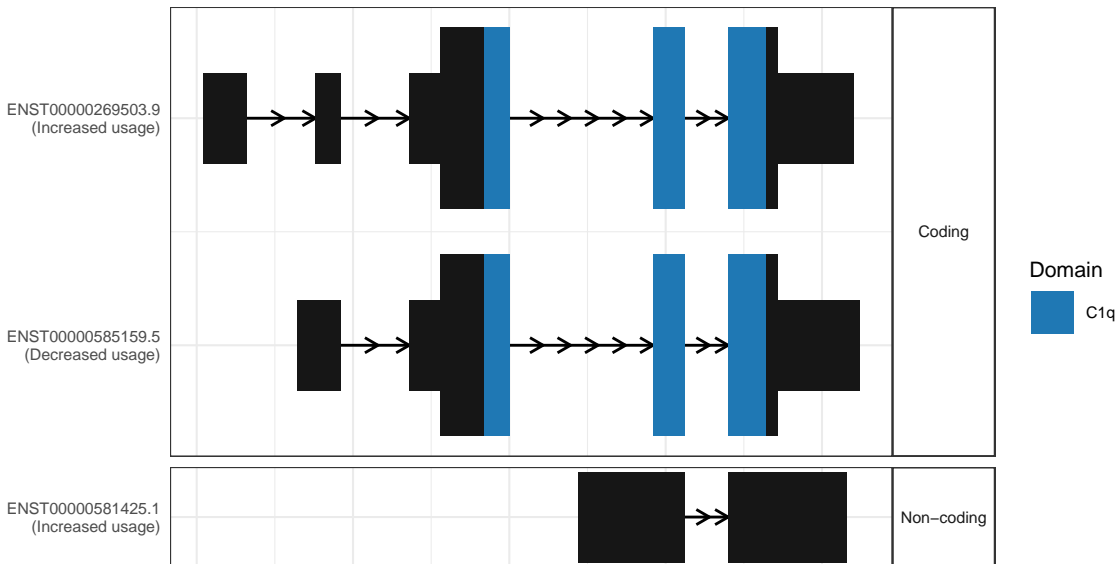


zf-C2H2 (x9)

The isoform switch in RTL8A (FT vs RGC)



The isoform switch in CBLN2 (FT vs RGC)

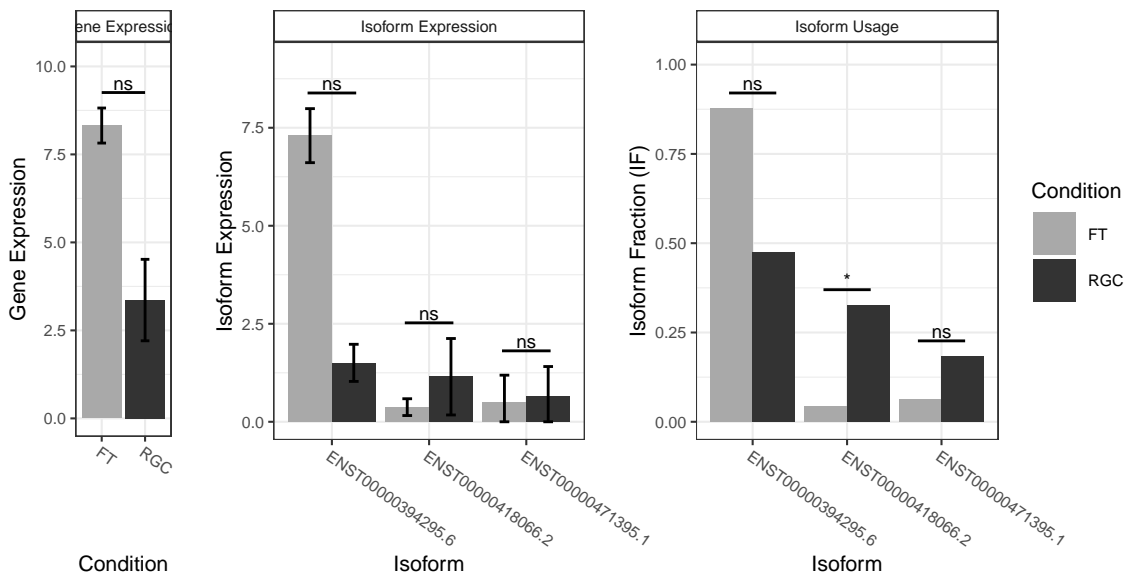
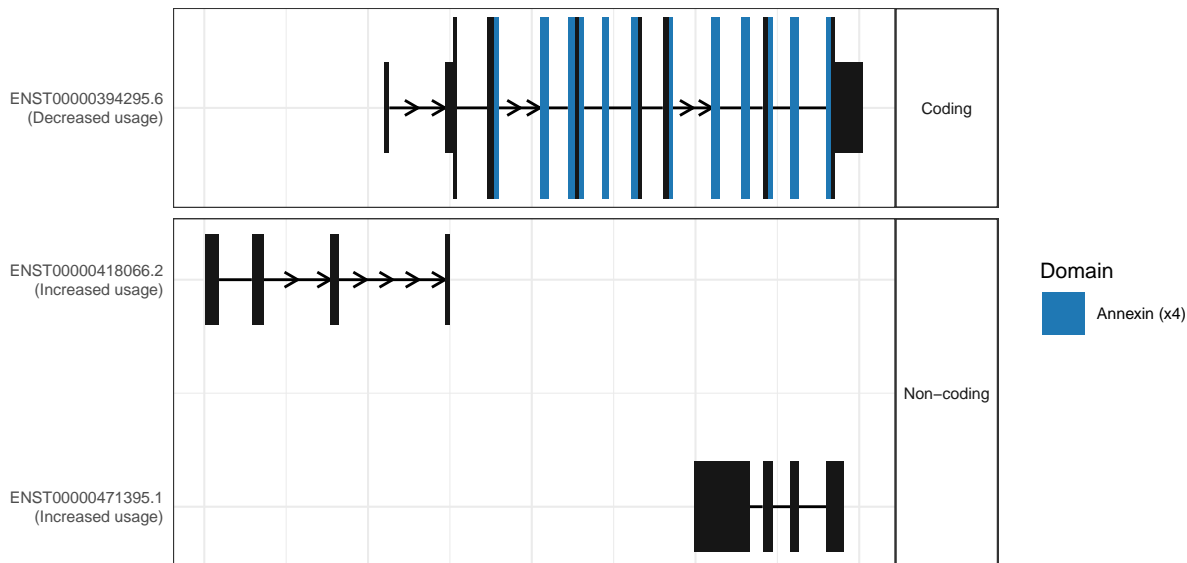


Domain



C1q

The isoform switch in ANXA4 (FT vs RGC)

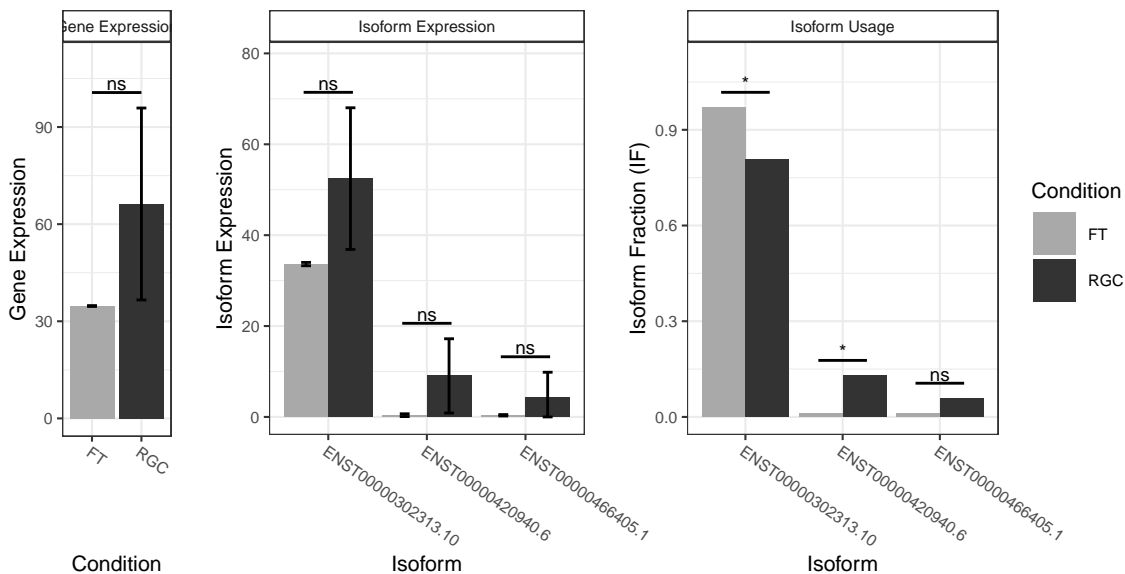
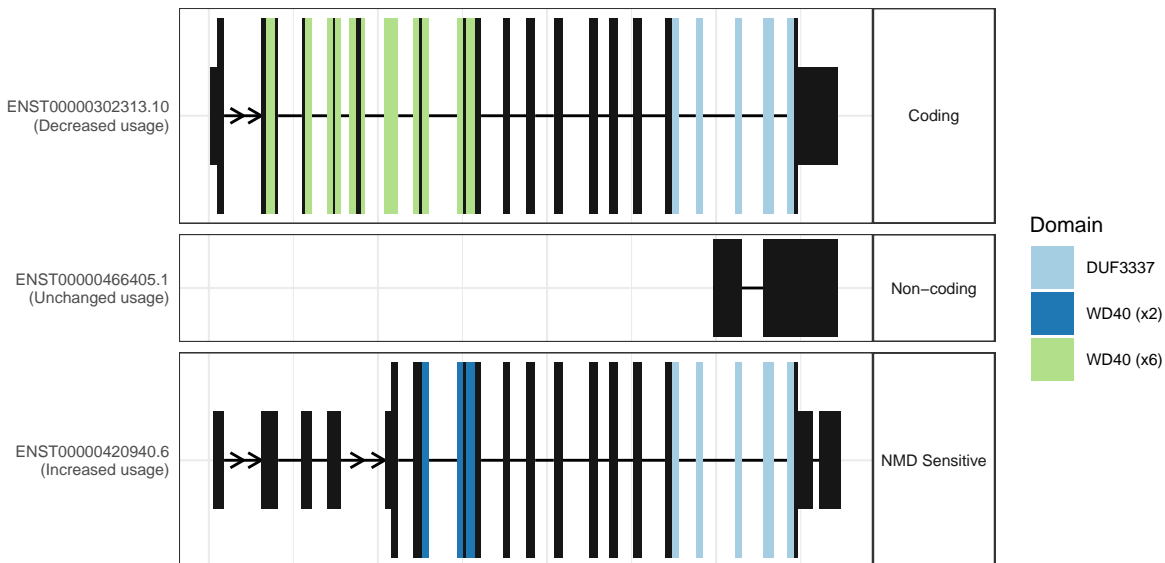


Domain



Annexin (x4)

The isoform switch in WDR48 (FT vs RGC)



Domain



DUF3337

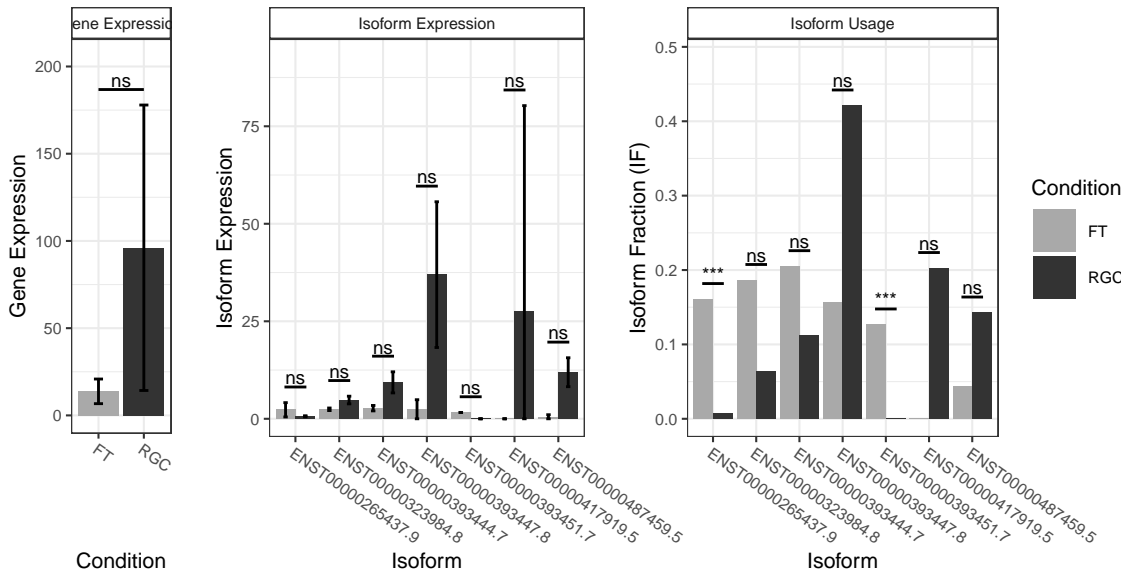
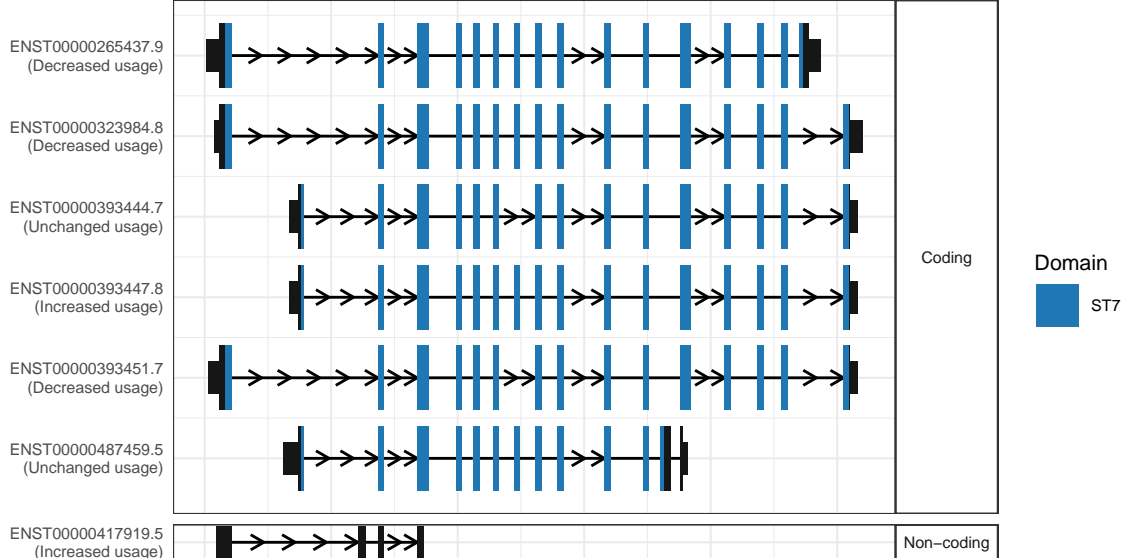


WD40 (x2)



WD40 (x6)

The isoform switch in ST7 (FT vs RGC)

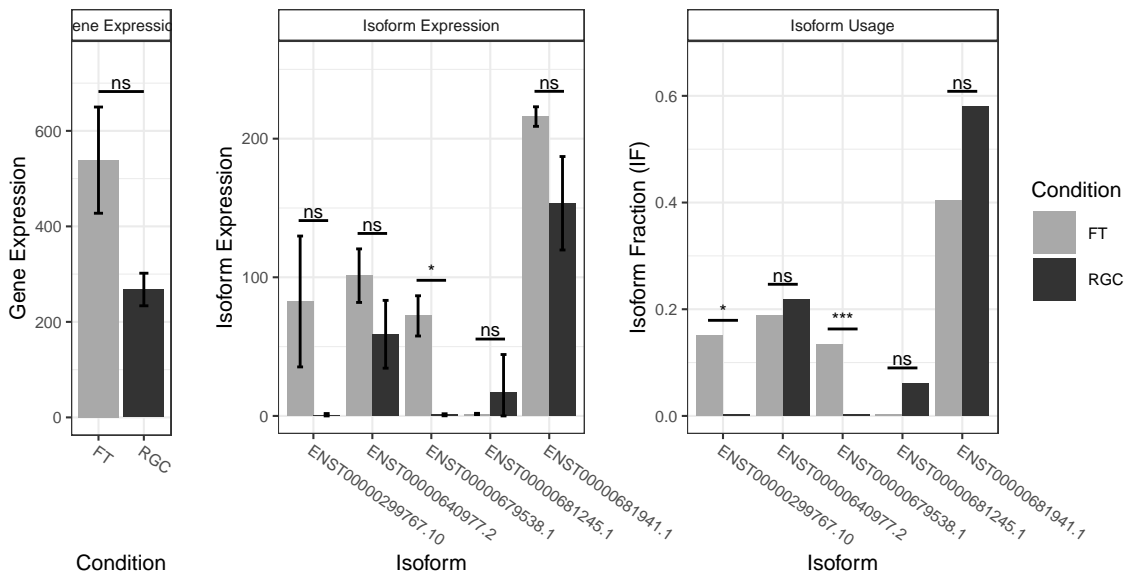
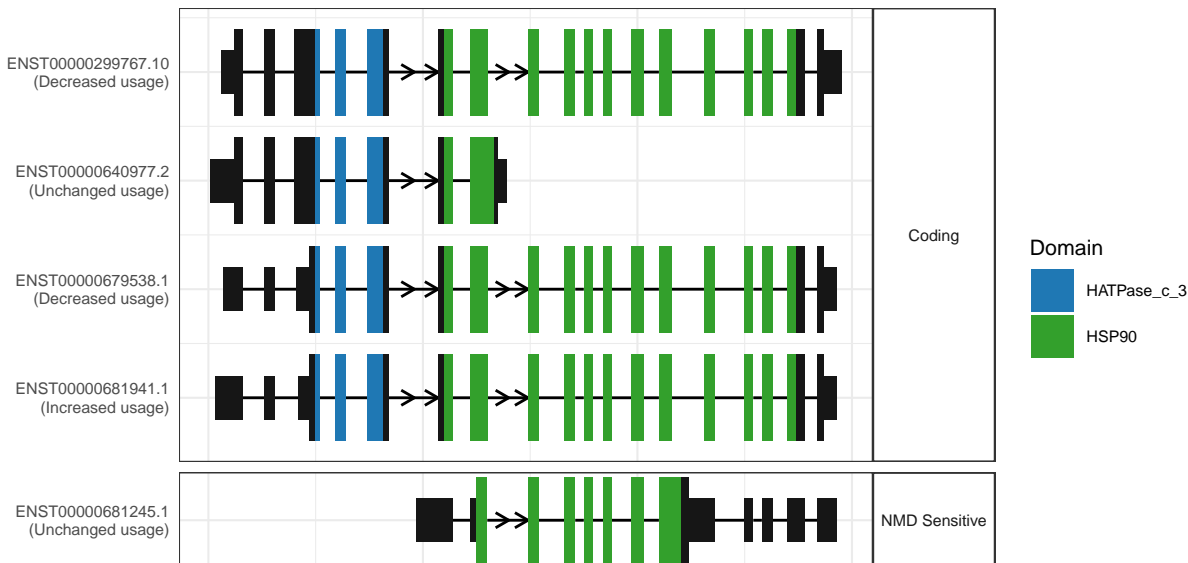


Domain



ST7

The isoform switch in HSP90B1 (FT vs RGC)



Domain

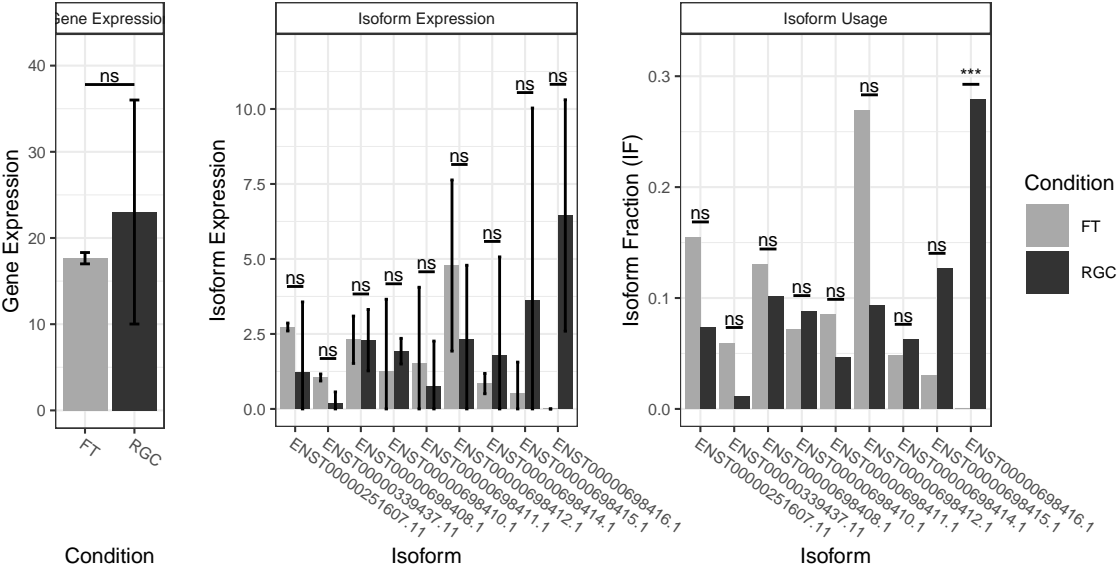
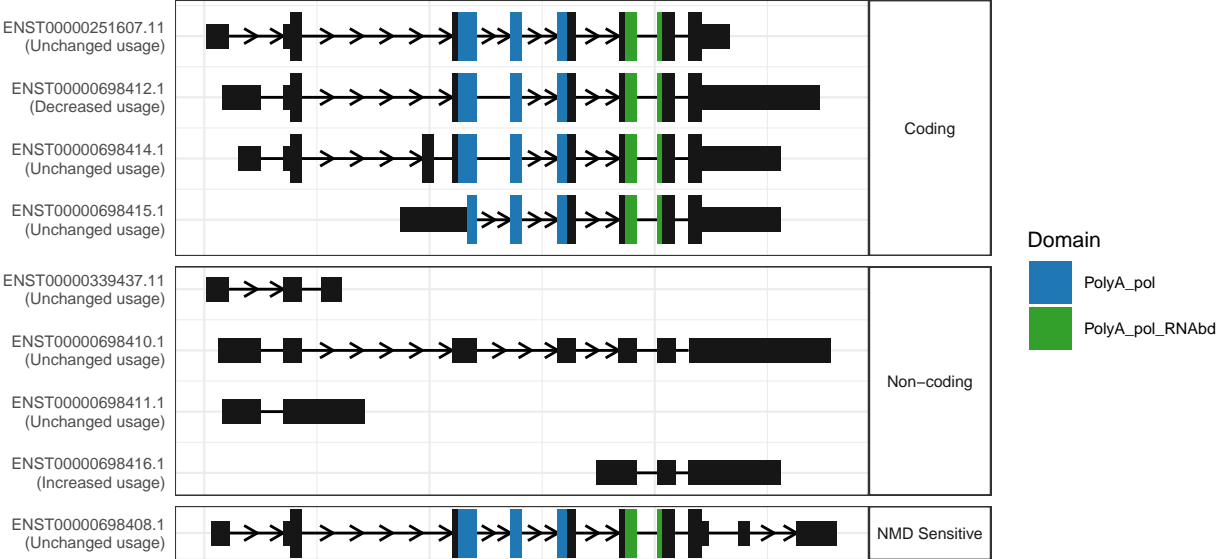


HATPase_c_3



HSP90

The isoform switch in TRNT1 (FT vs RGC)



Domain

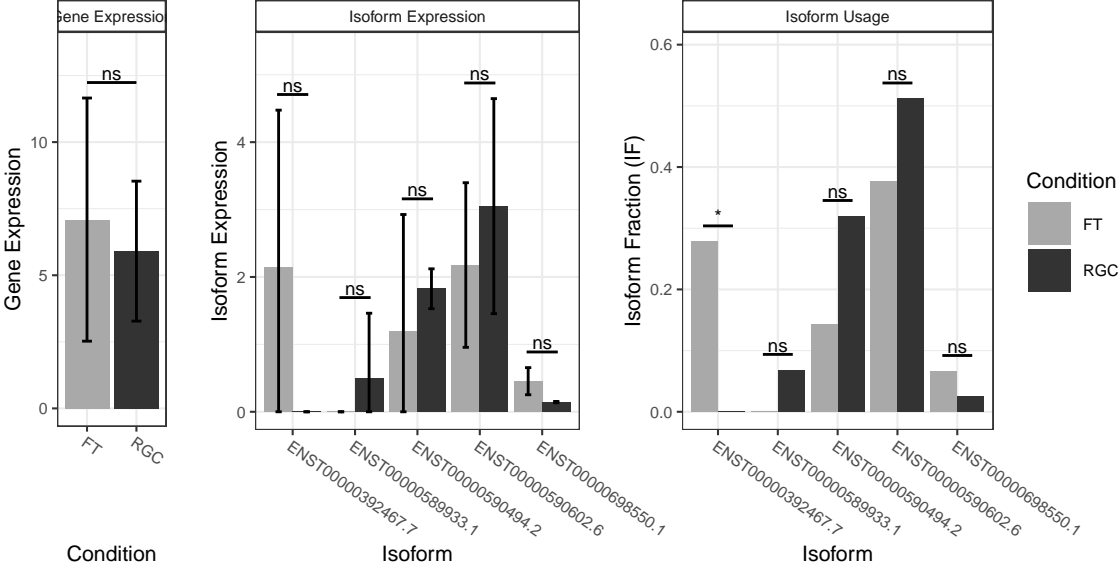
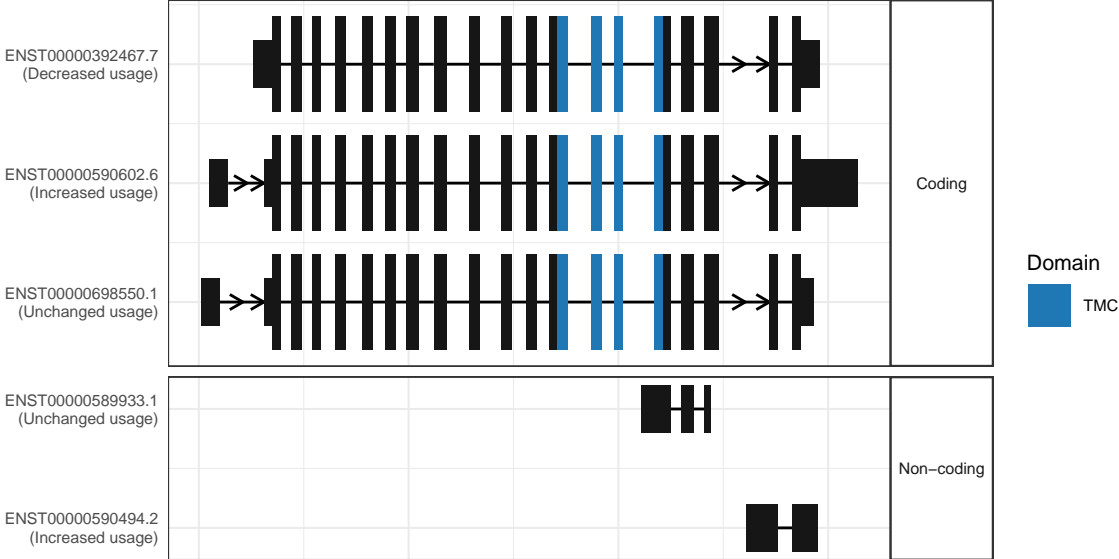


PolyA_pol



PolyA_pol_RNAAbd

The isoform switch in TMC6 (FT vs RGC)

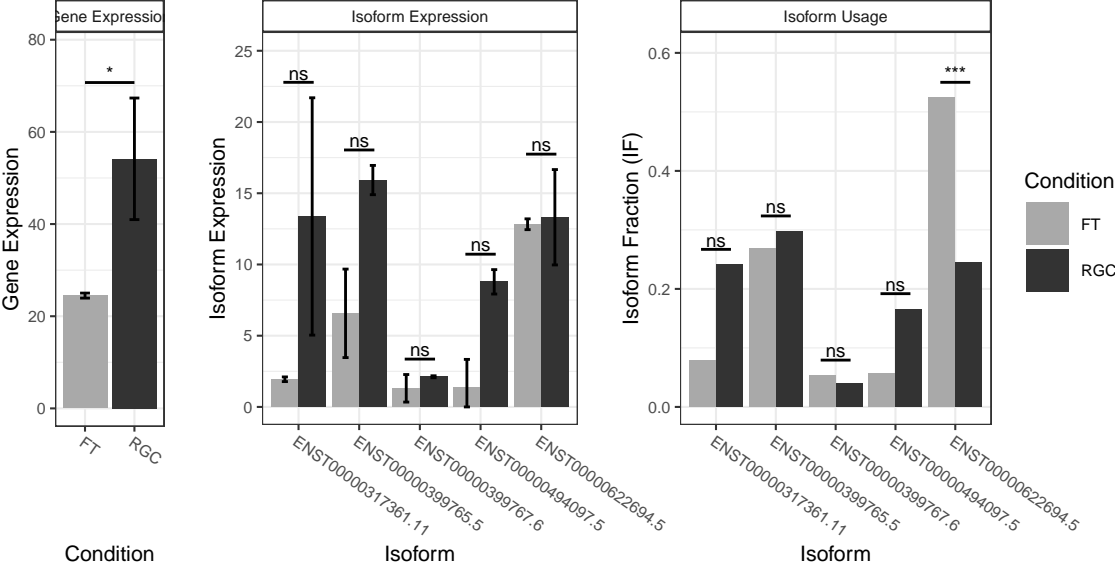
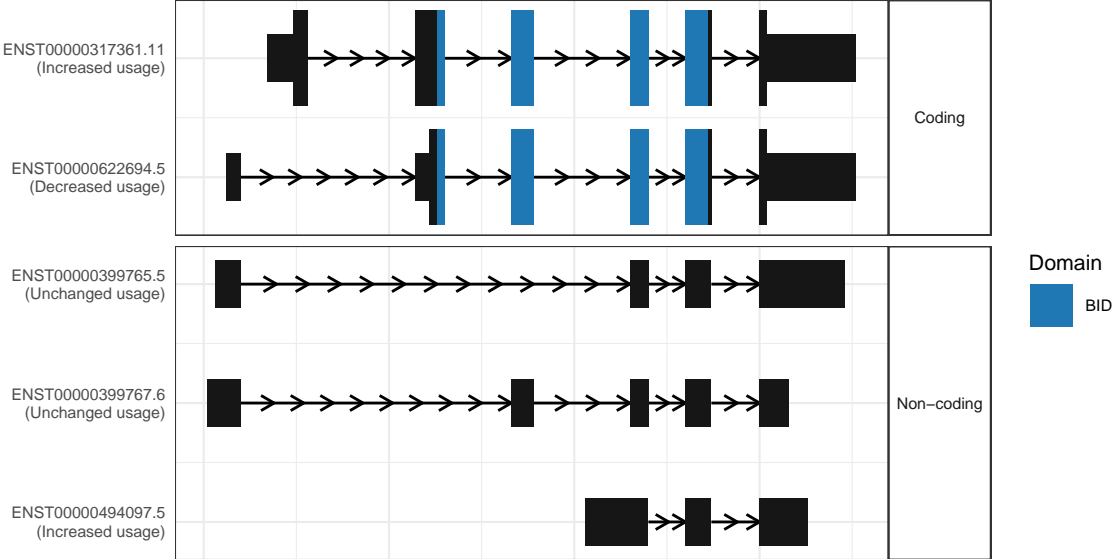


Domain



TMC

The isoform switch in BID (FT vs RGC)

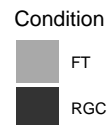
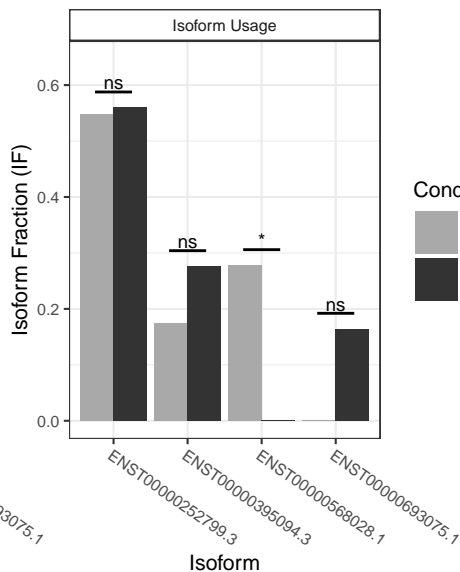
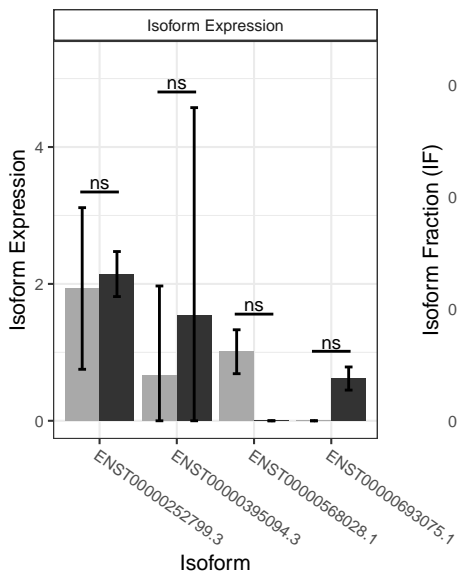
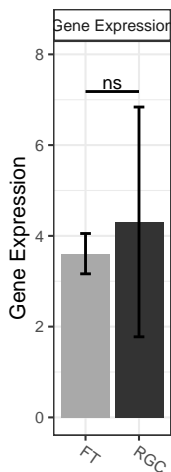
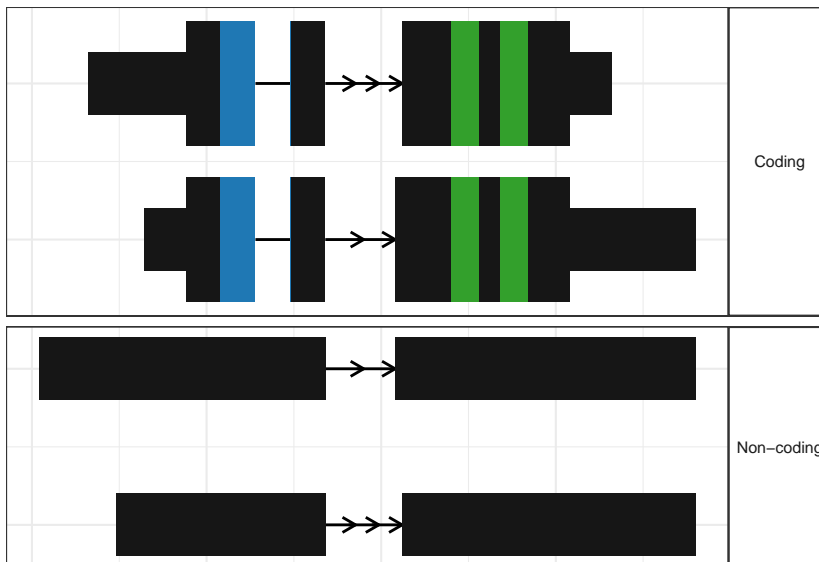


Domain



BID

The isoform switch in ZNF747 (FT vs RGC)



Domain

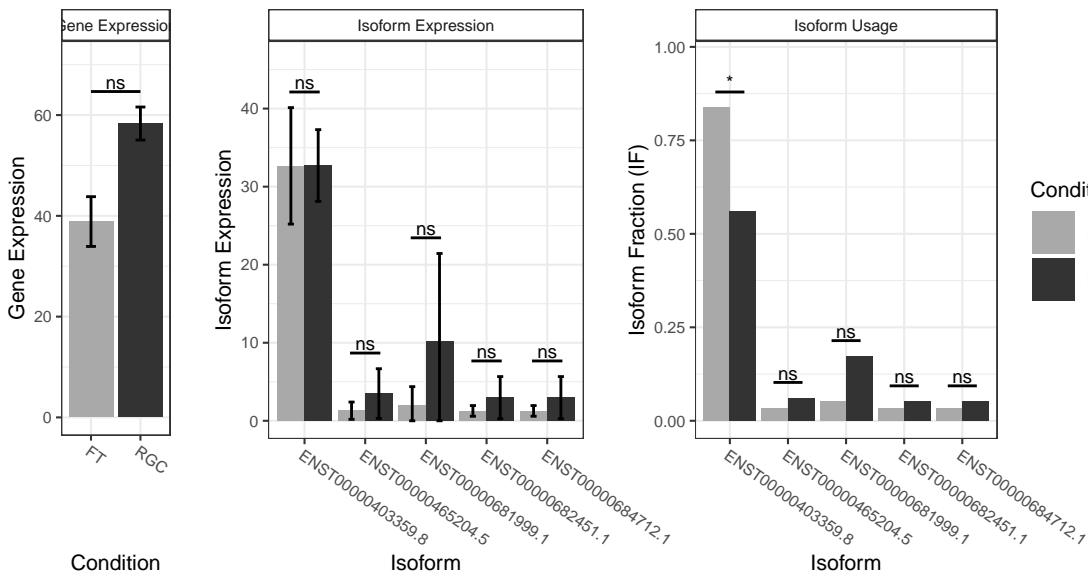
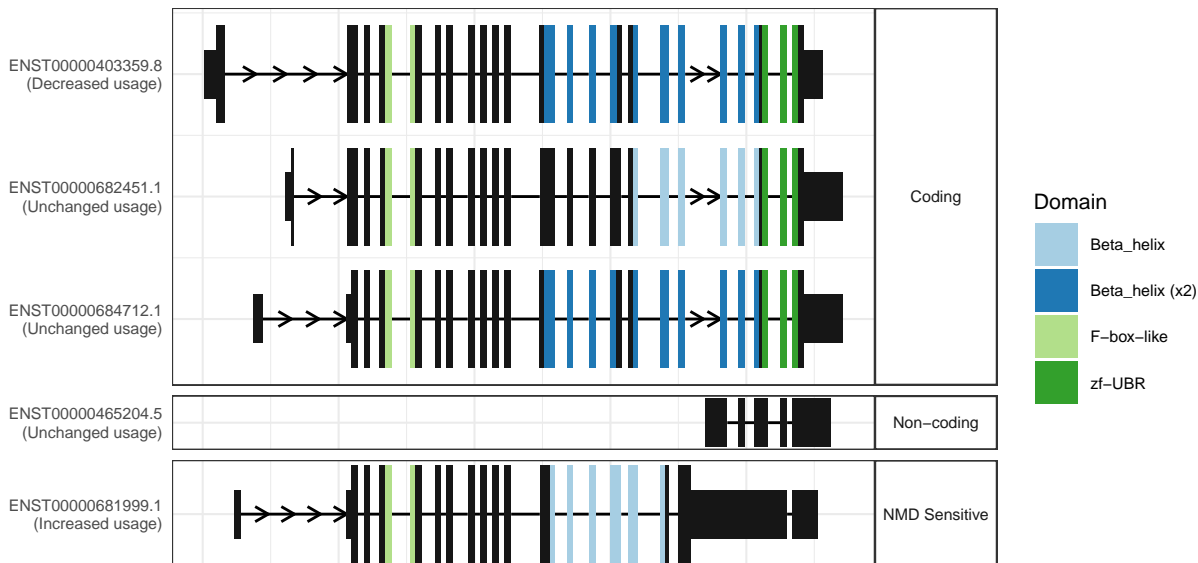


KRAB







zf-C2H2 (x2)

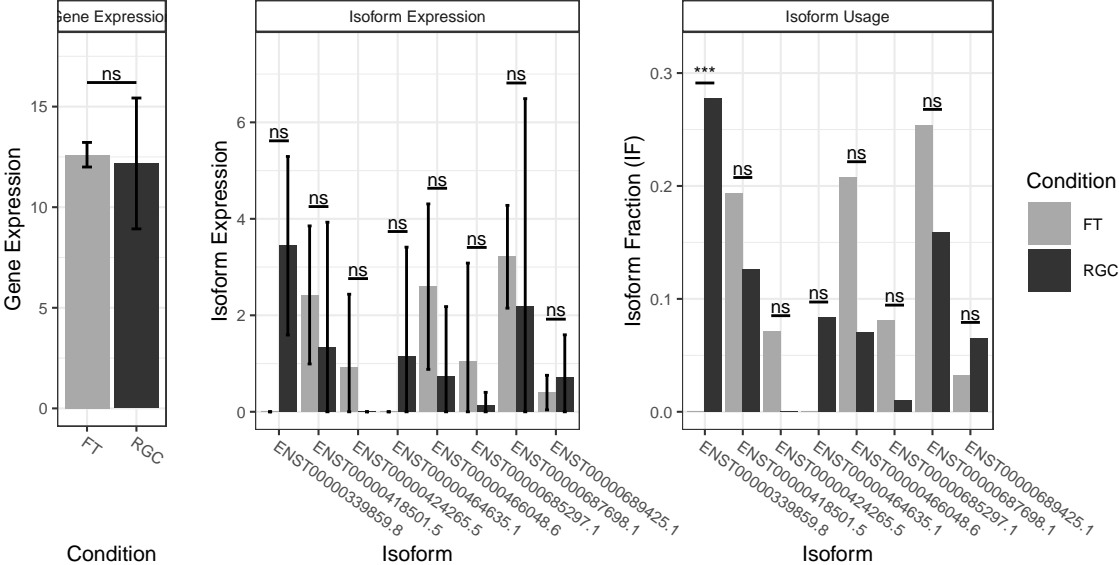
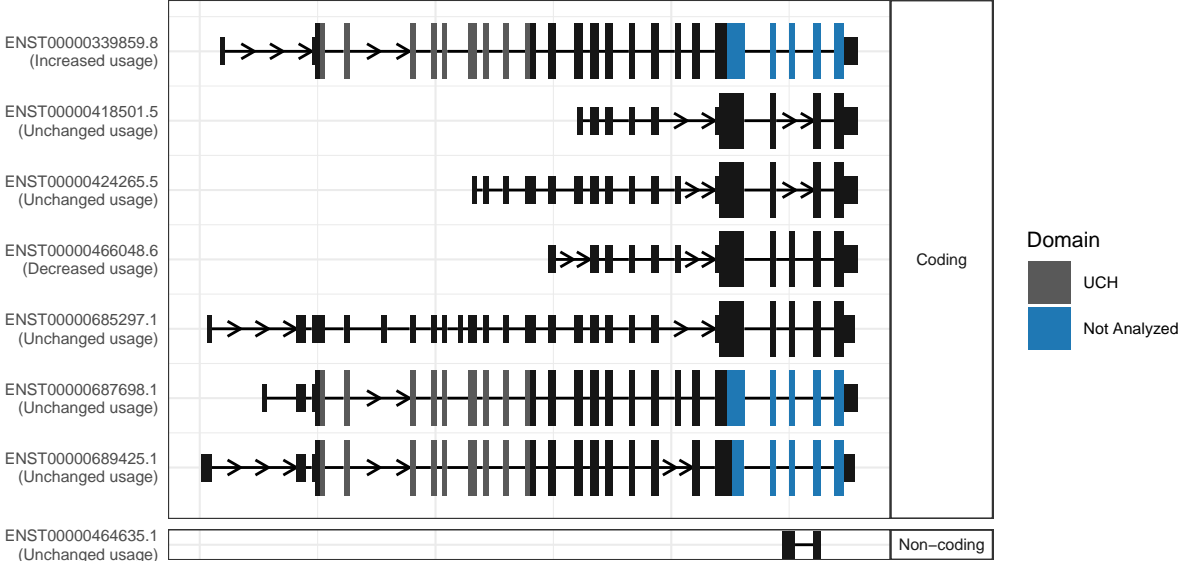
The isoform switch in FBXO11 (FT vs RGC)



Domain

	Beta_helix
	Beta_helix (x2)
	F-box-like
	zf-UBR

The isoform switch in USP54 (FT vs RGC)



Domain

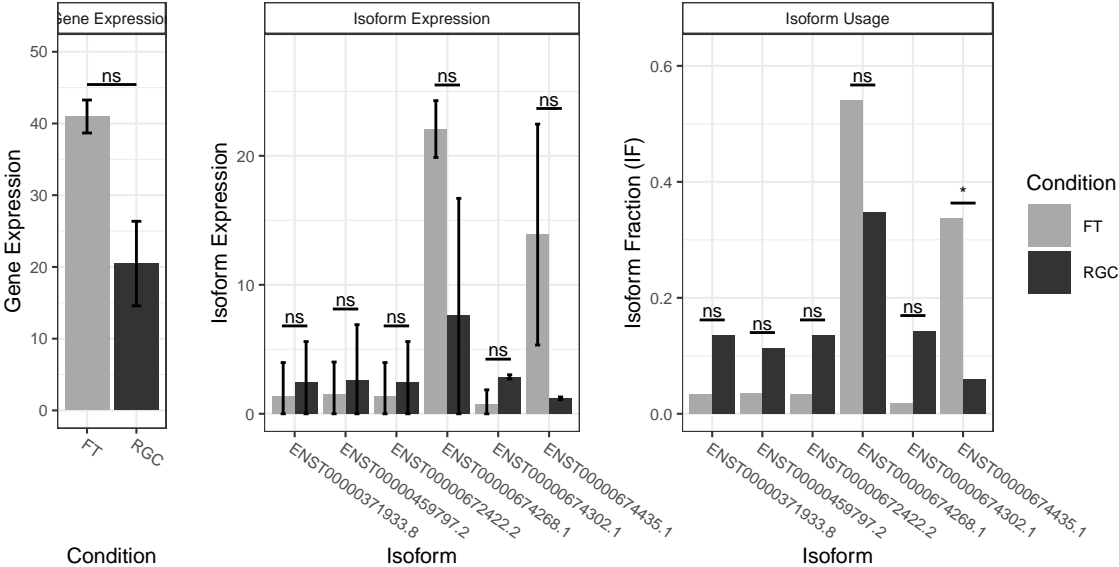
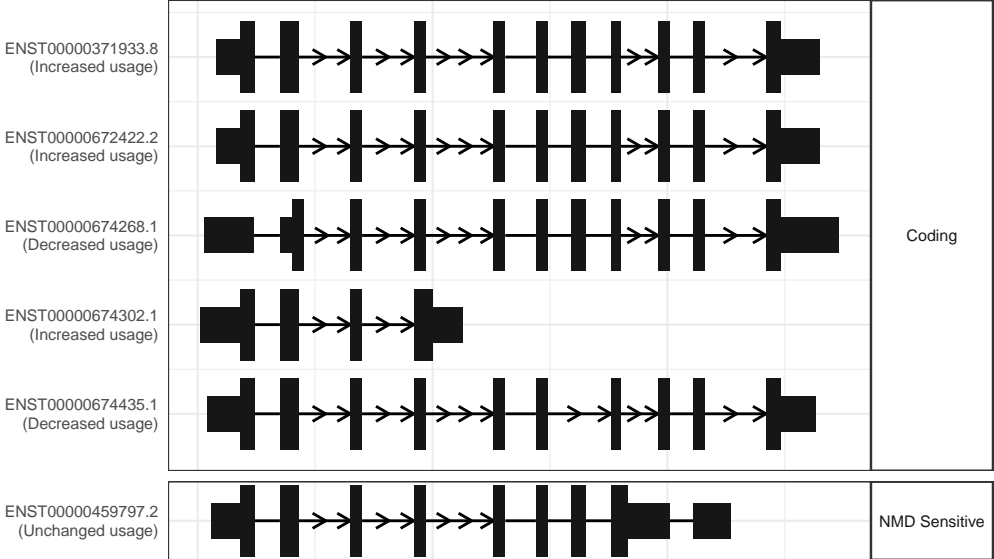


UCH

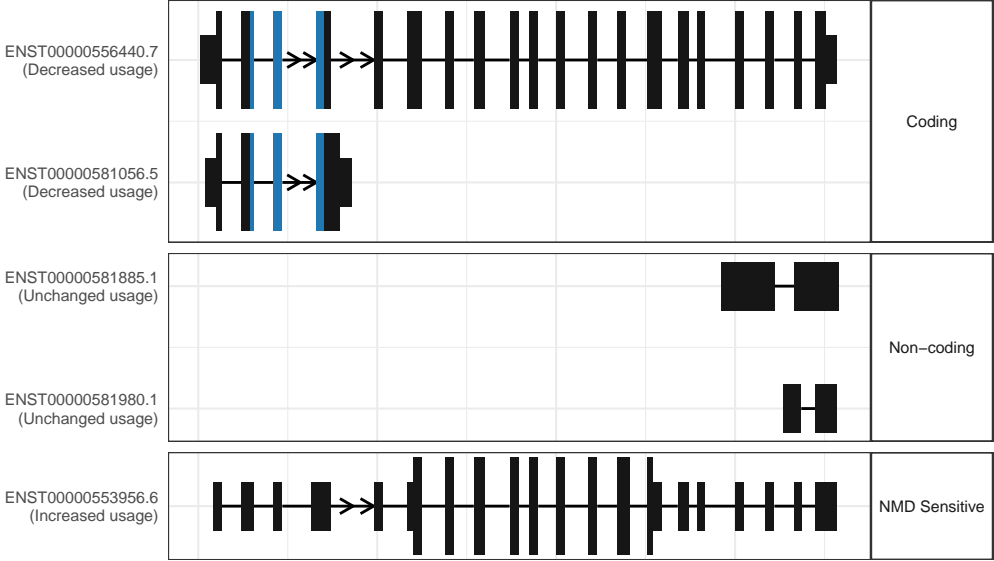


Not Analyzed

The isoform switch in EFCAB14 (FT vs RGC)

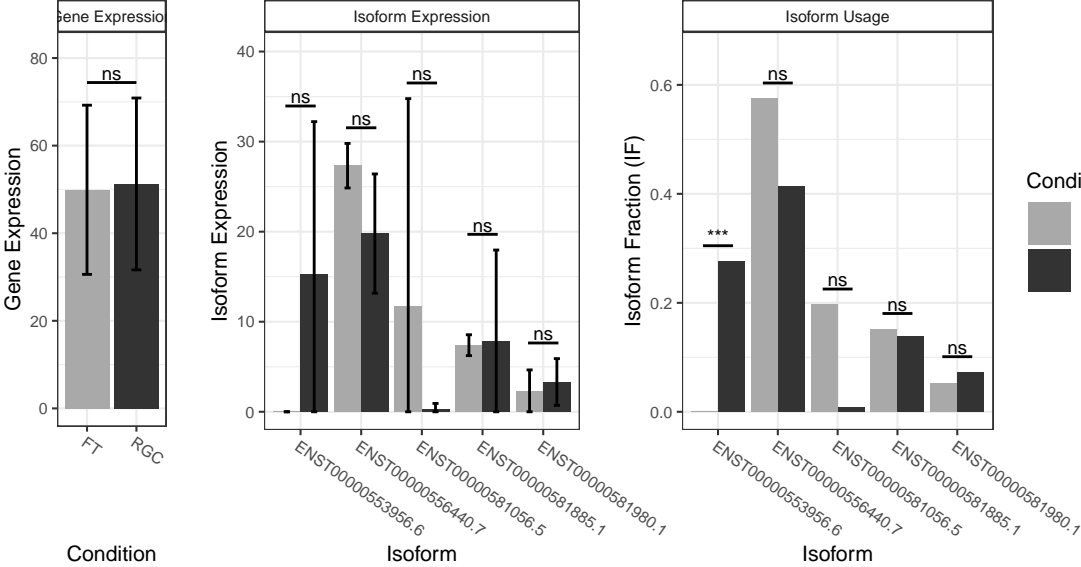


The isoform switch in CEP95 (FT vs RGC)



Domain

DUF5745

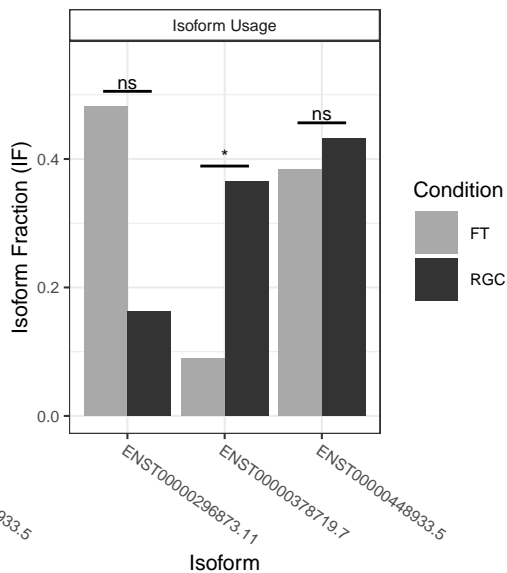
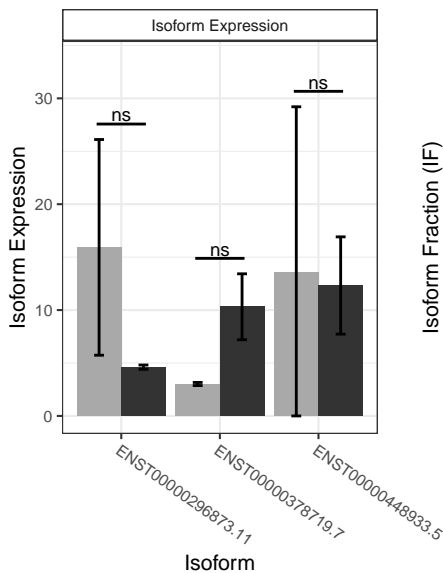
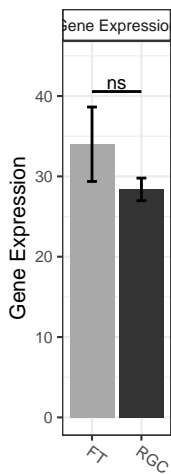
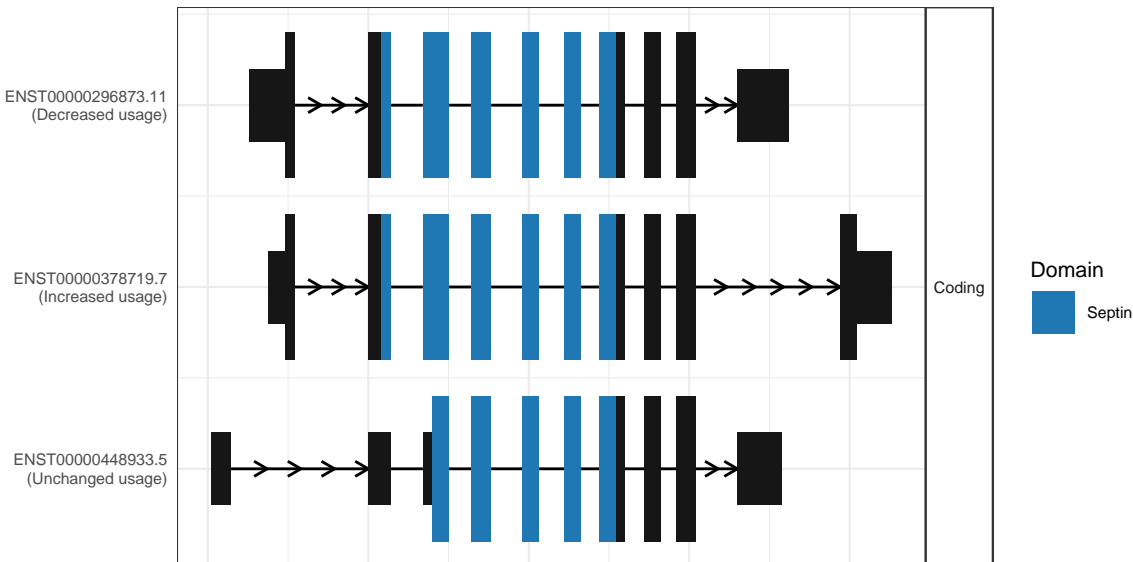


Domain



DUF5745

The isoform switch in SEPTIN8 (FT vs RGC)

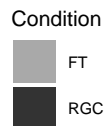
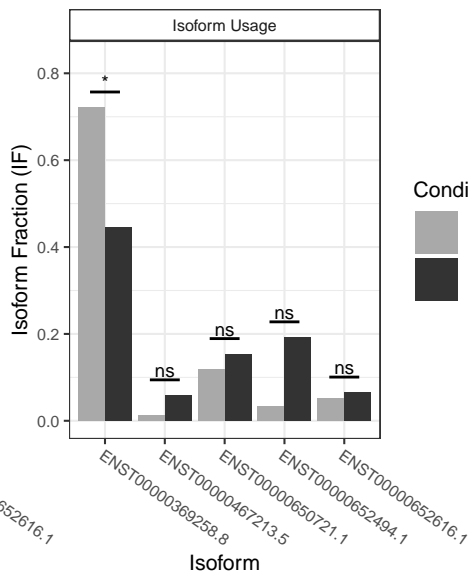
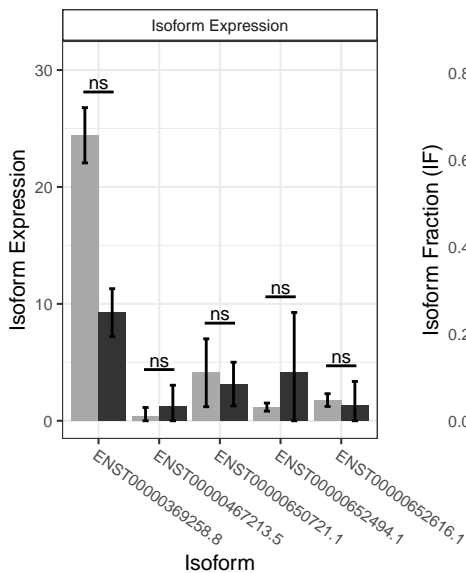
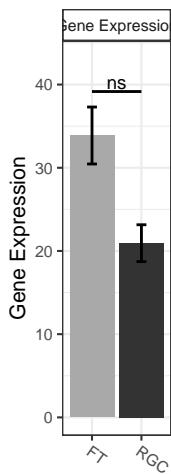
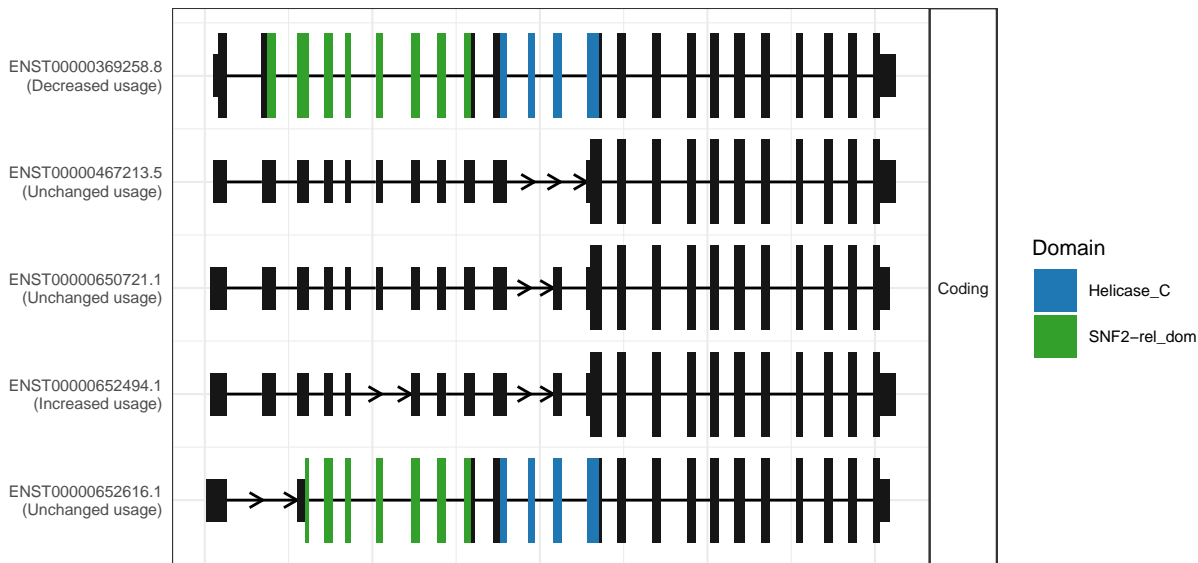


Domain



Septin

The isoform switch in CHD1L (FT vs RGC)



Domain

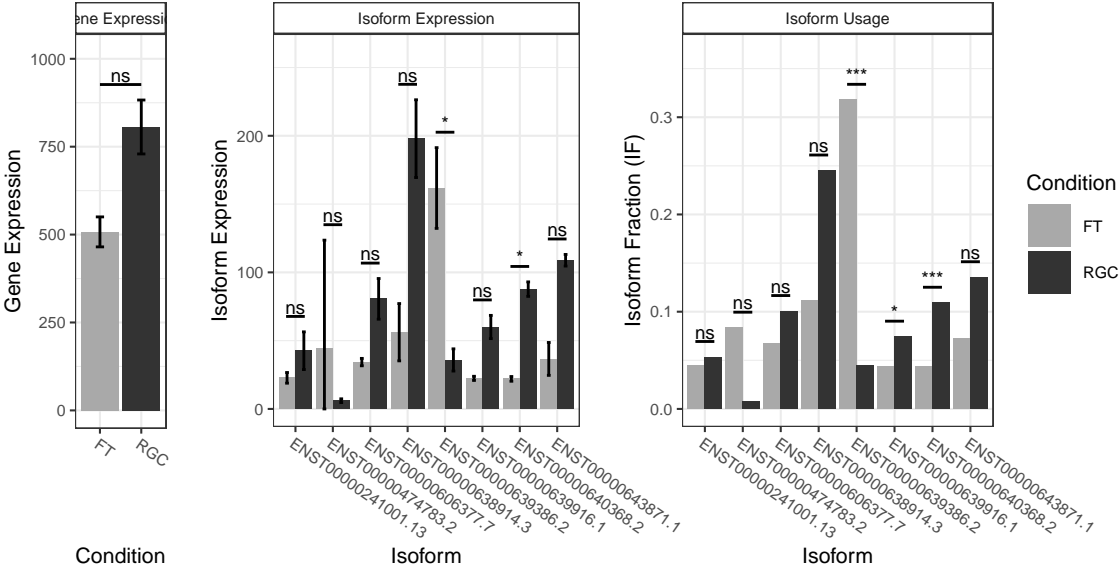
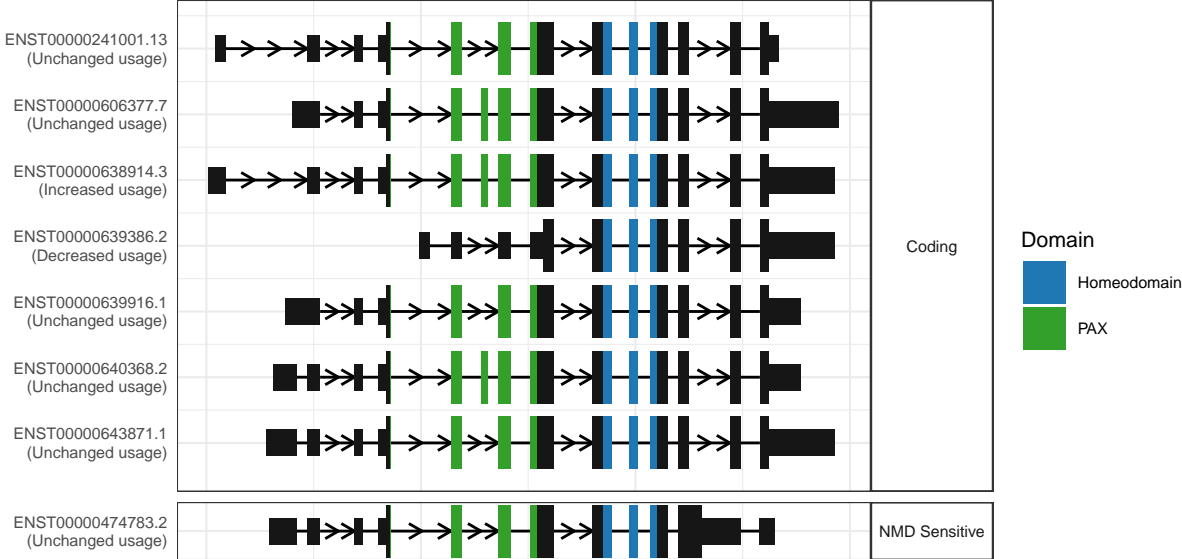


Helicase_C



SNF2-rel_dom

The isoform switch in PAX6 (FT vs RGC)



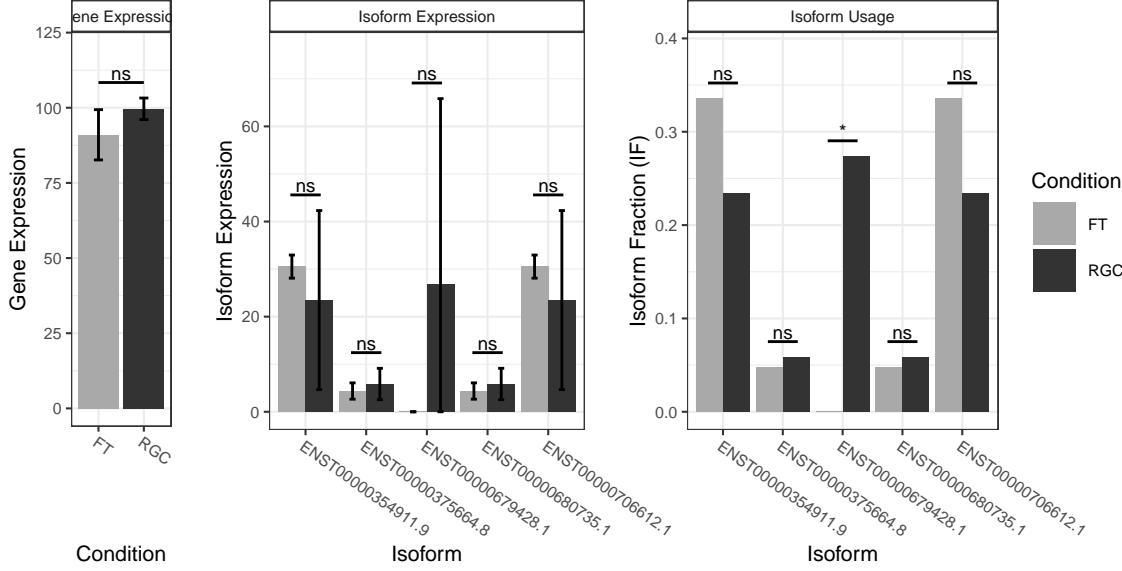
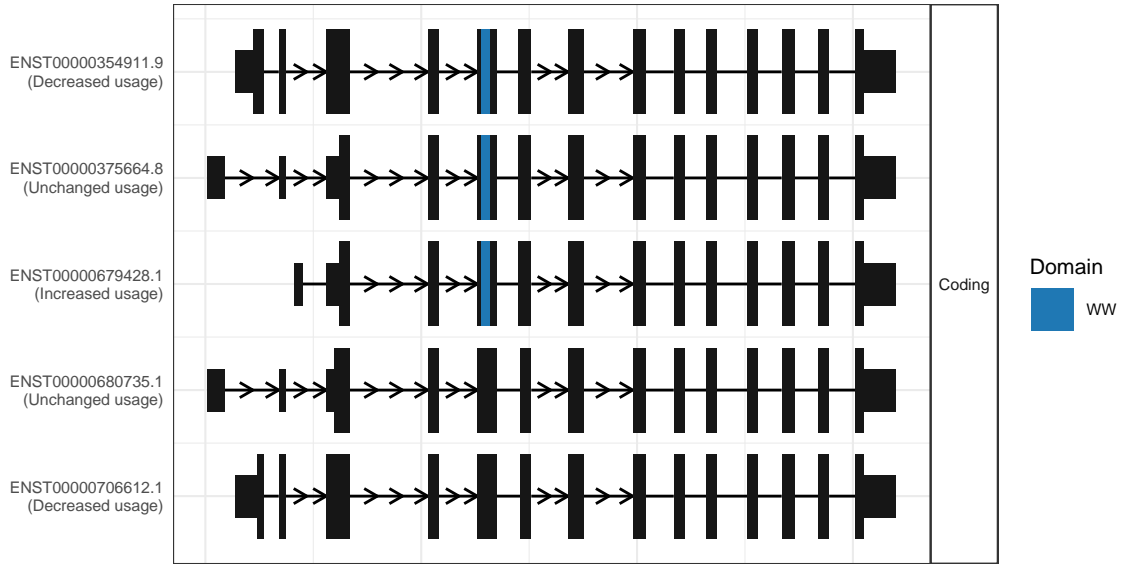
Domain



Homeodomain

PAX

The isoform switch in WAC (FT vs RGC)

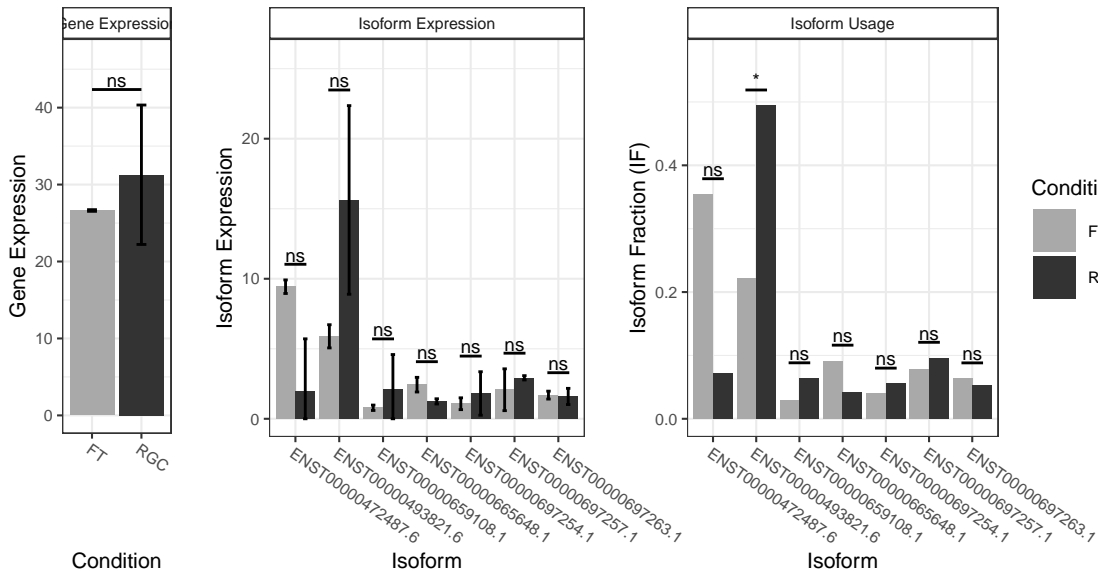
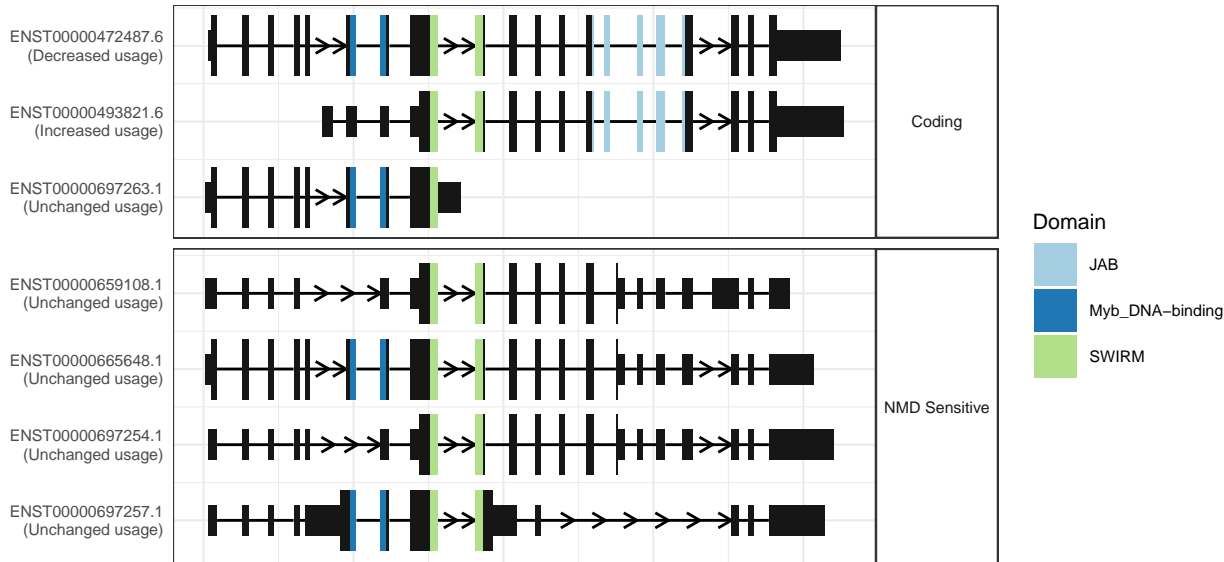


Domain



ww

The isoform switch in MYSM1 (FT vs RGC)



Domain



JAB



Myb_DNA-binding



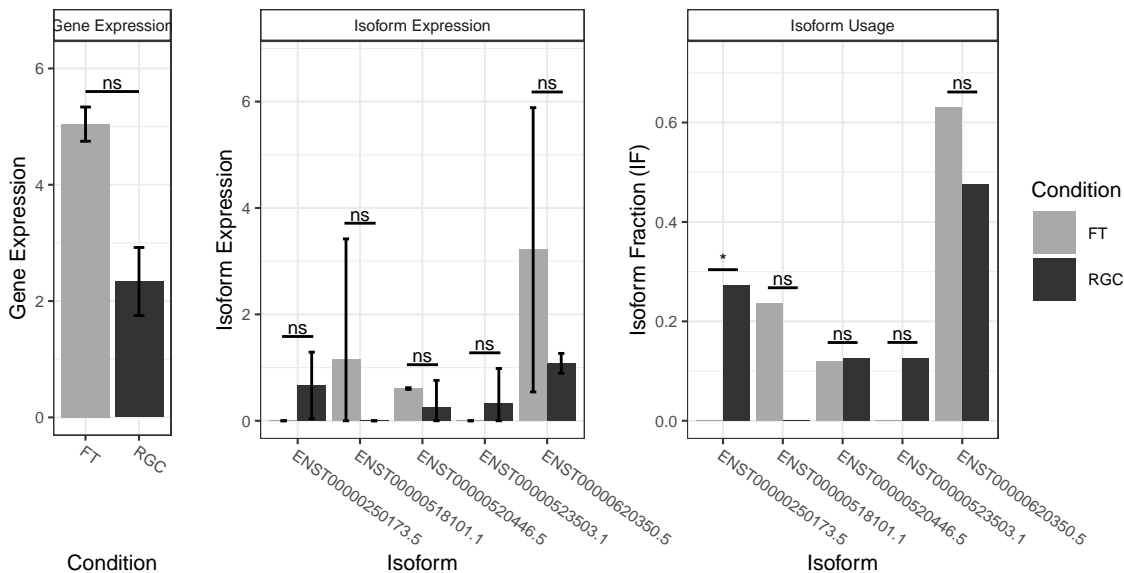
SWIRM

	Coding
	Non-coding
	NMD Sensitive

 LRR_9

Non-coding

NMD Sensitive

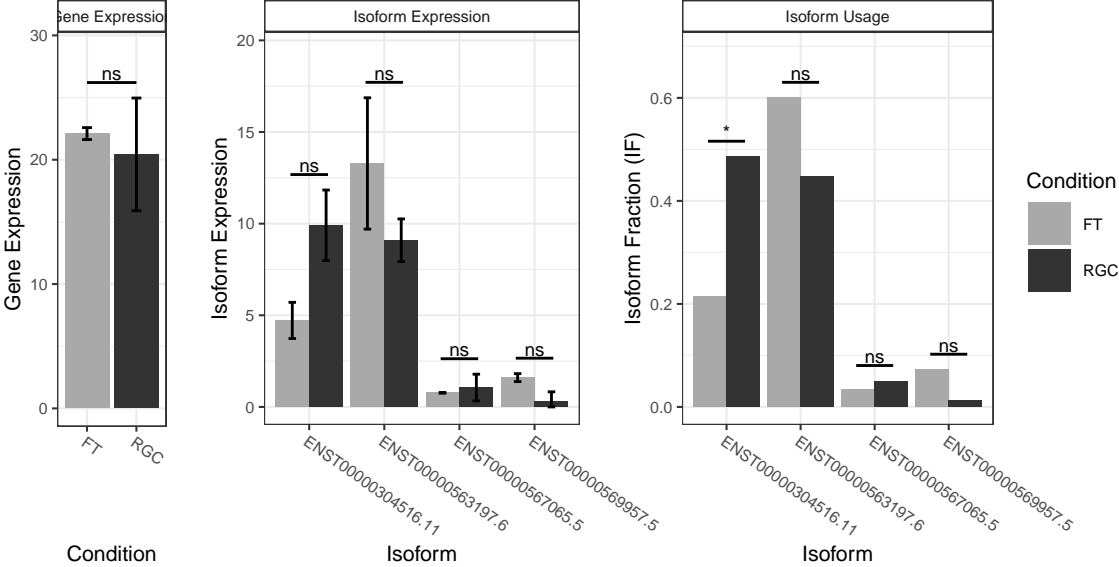
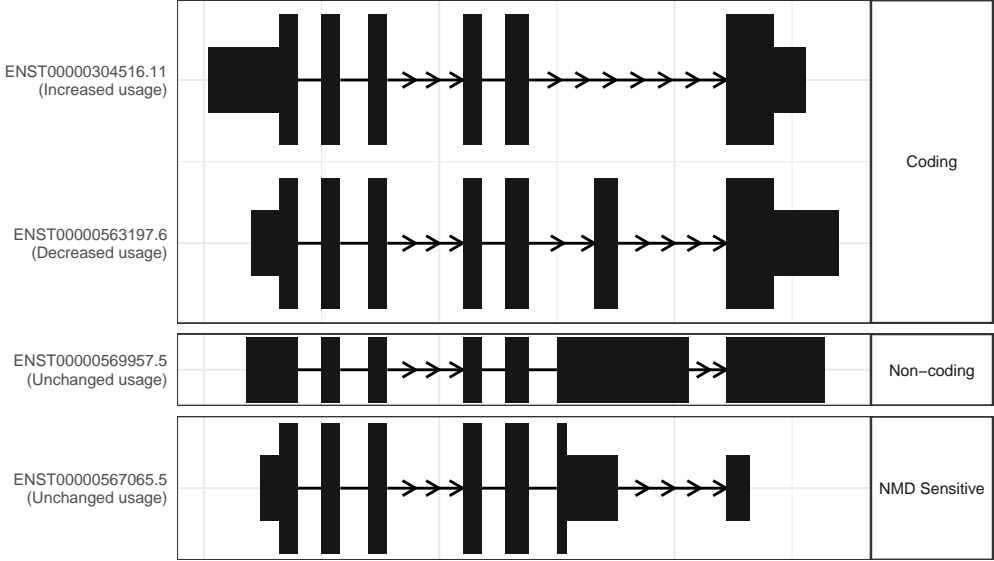


Domain

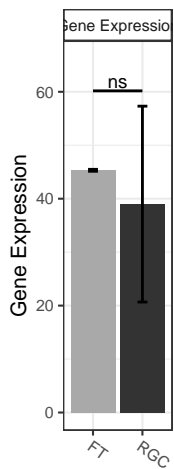
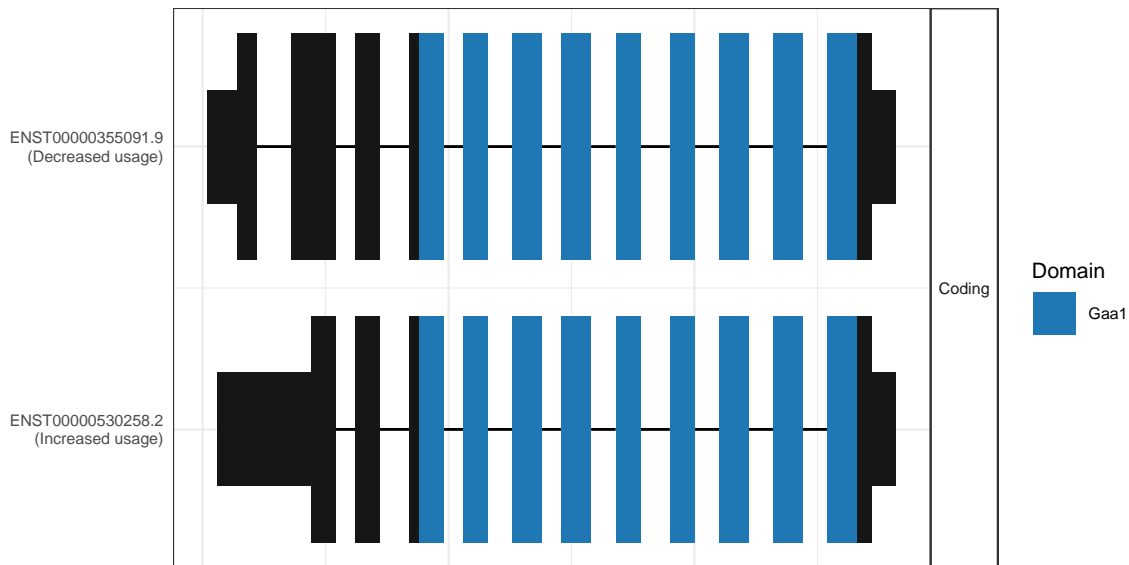


LRR_9

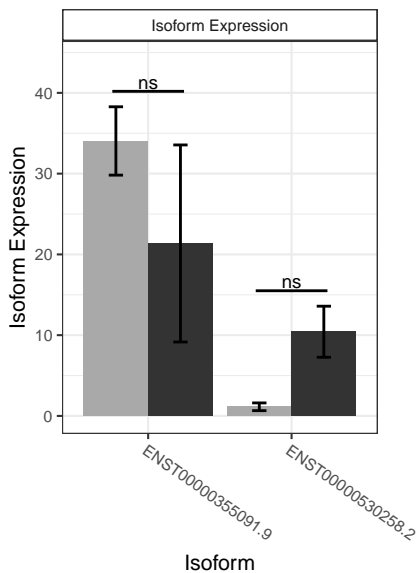
The isoform switch in INO80E (FT vs RGC)



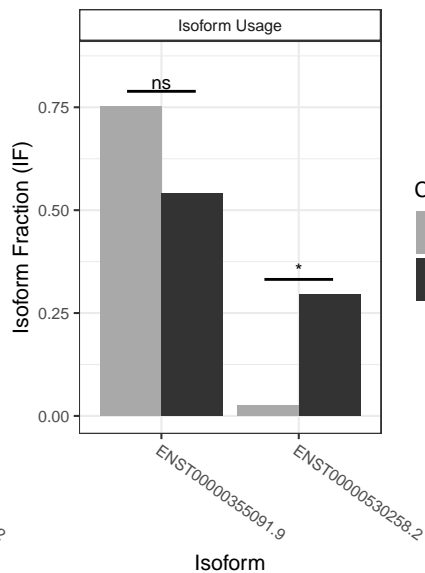
The isoform switch in GPAA1 (FT vs RGC)



Condition



Isoform



Isoform

Condition

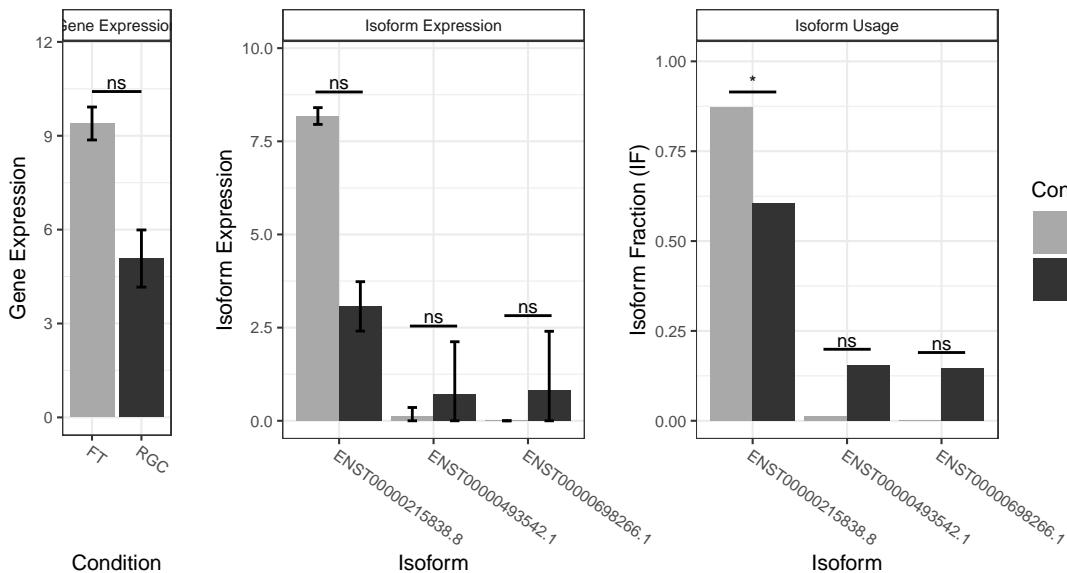
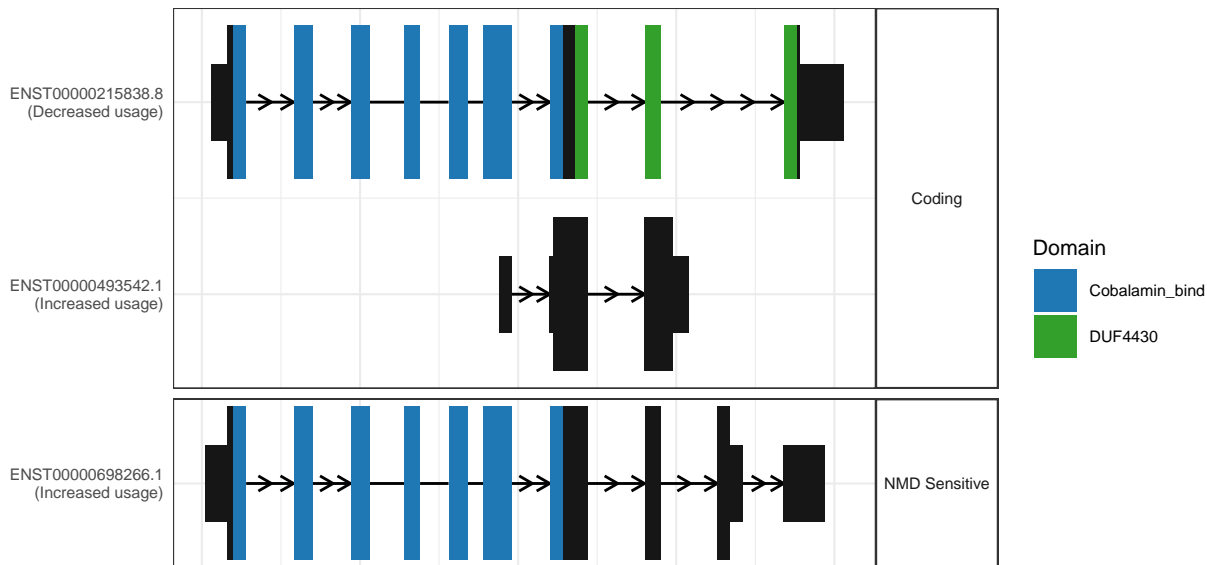


Domain



Gaa1

The isoform switch in TCN2 (FT vs RGC)



Domain

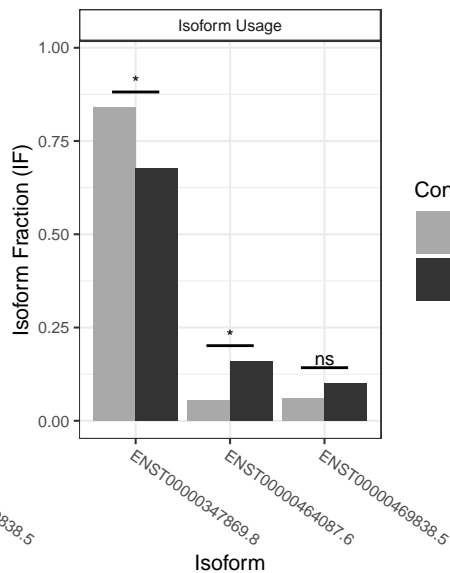
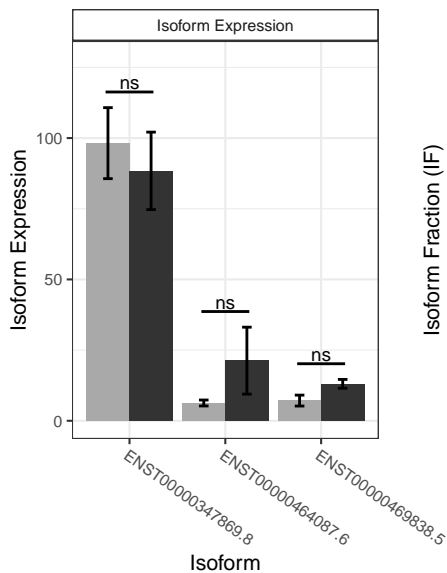
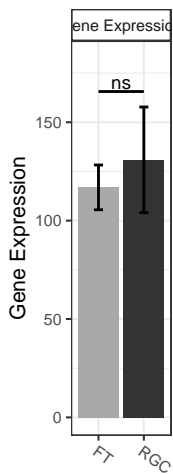
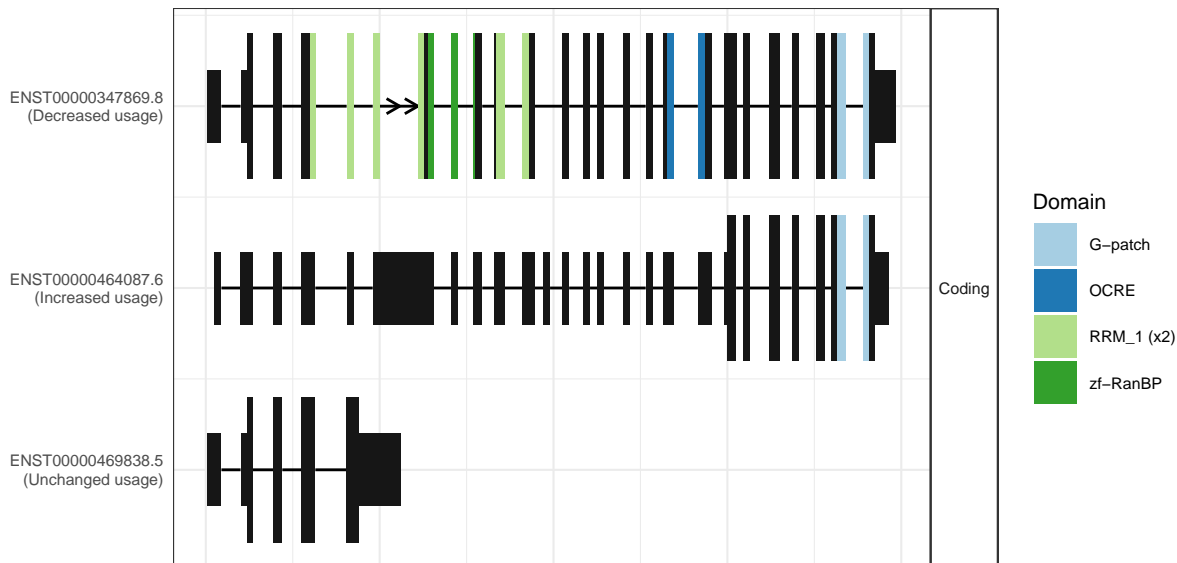


Cobalamin_bind



DUF4430

The isoform switch in RBM5 (FT vs RGC)



Domain



G-patch



OCRE

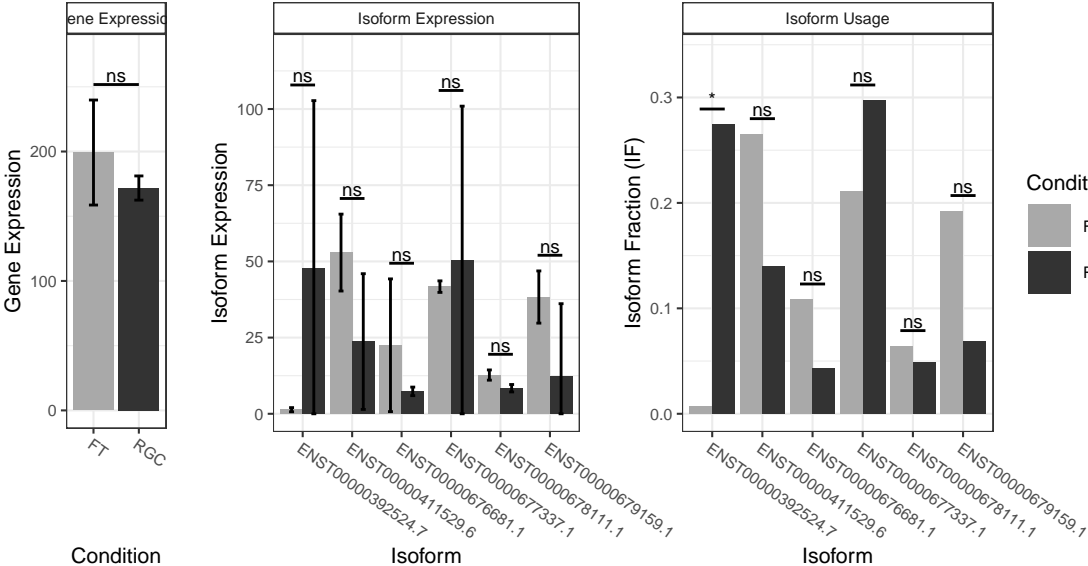
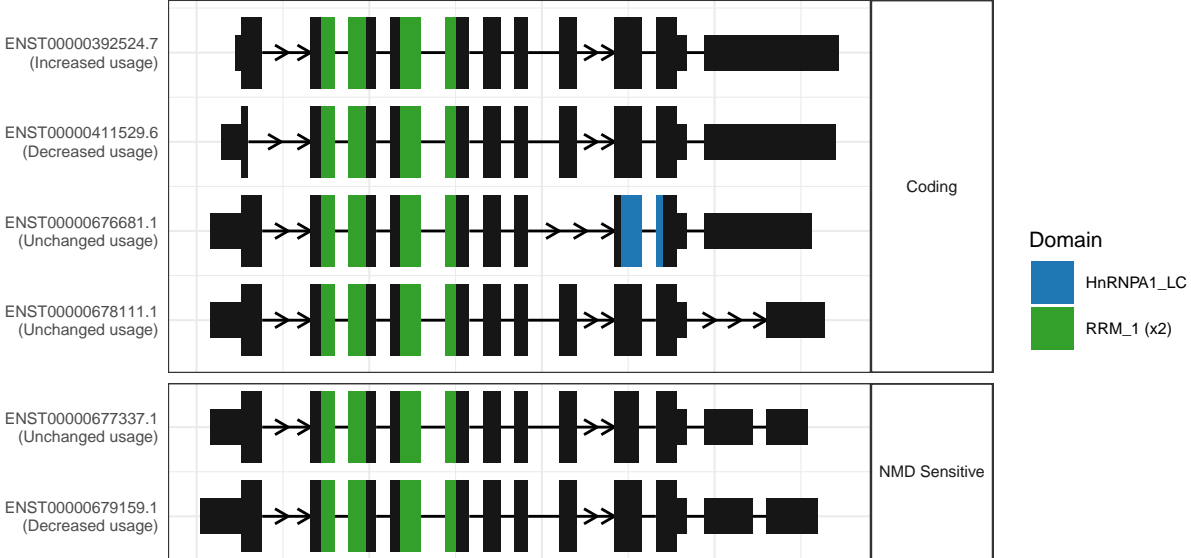


RRM_1 (x2)



zf-RanBP

The isoform switch in HNRNPA3 (FT vs RGC)



Domain

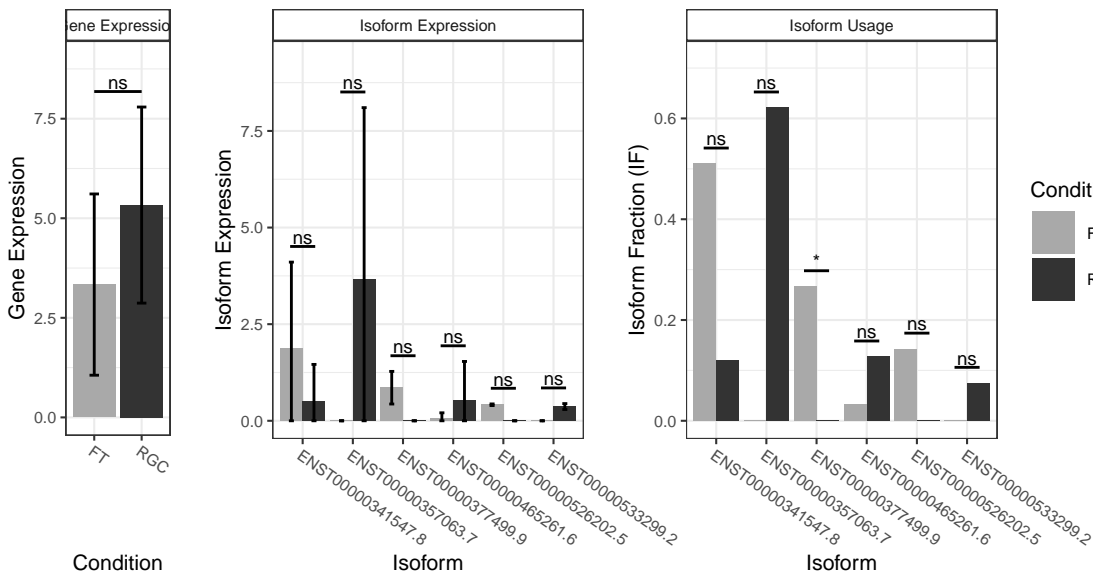
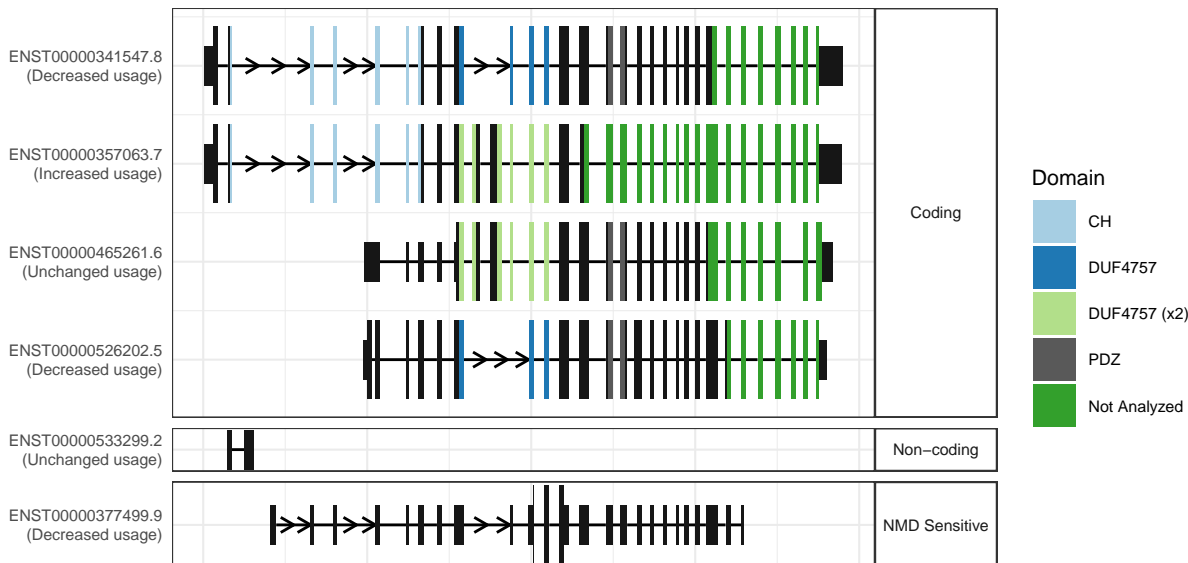


HnRNPA1_LC








RRM_1 (x2)

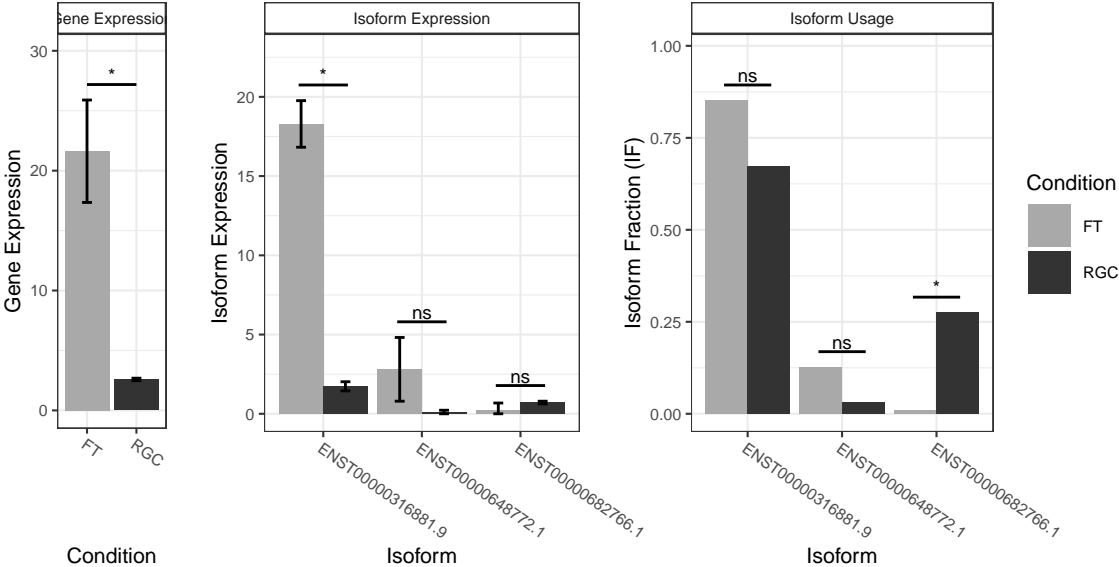
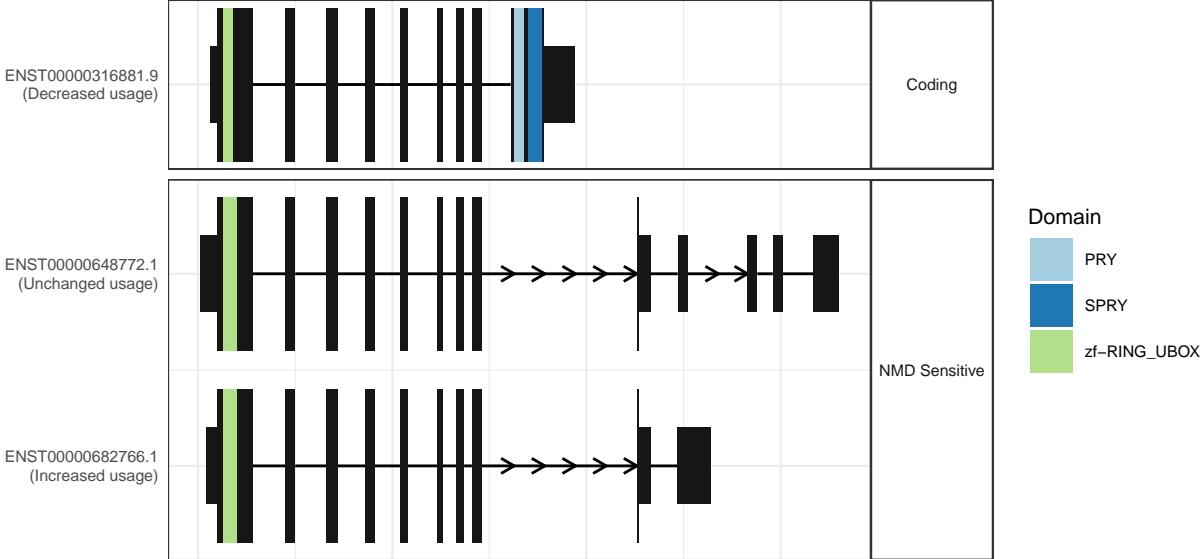
The isoform switch in LMO7 (FT vs RGC)



Domain

	CH
	DUF4757
	DUF4757 (x2)
	PDZ
	Not Analyzed

The isoform switch in TRIM25 (FT vs RGC)



Domain



PRY

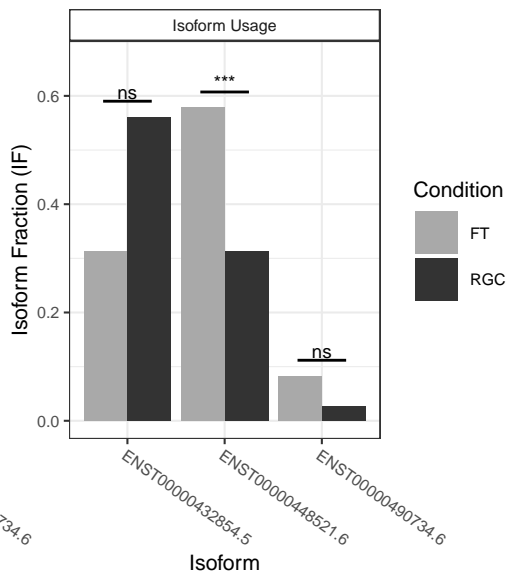
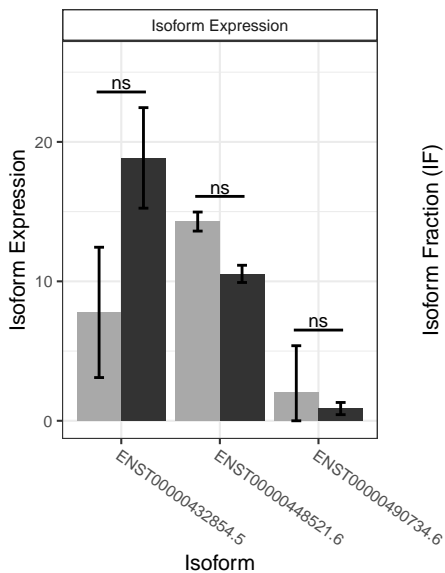
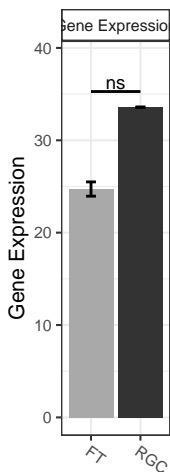
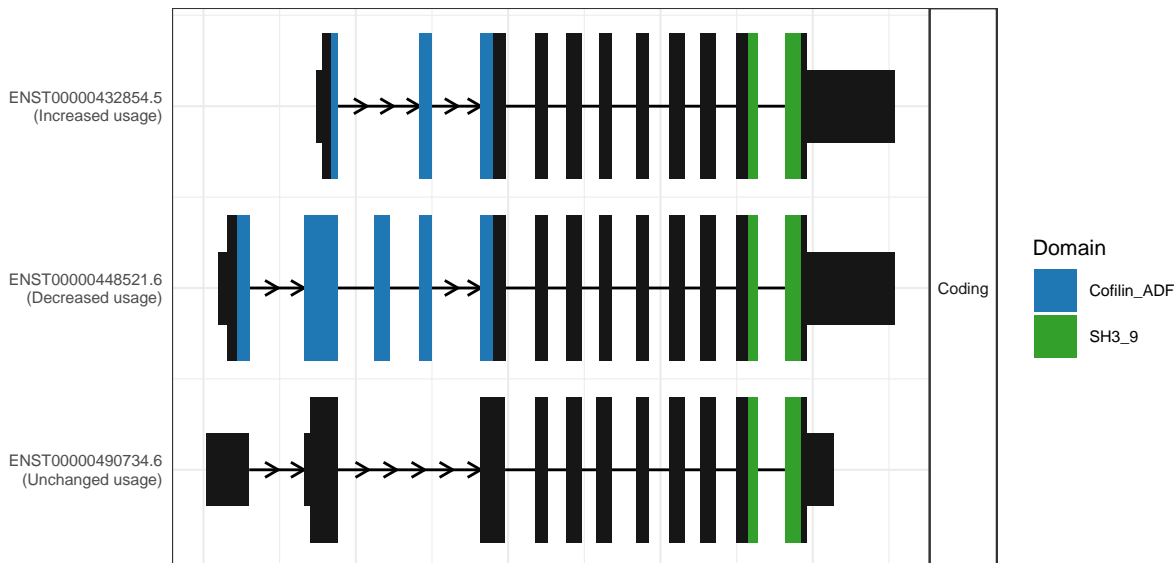


SPRY



zf-RING_UBOX

The isoform switch in DBNL (FT vs RGC)



Domain

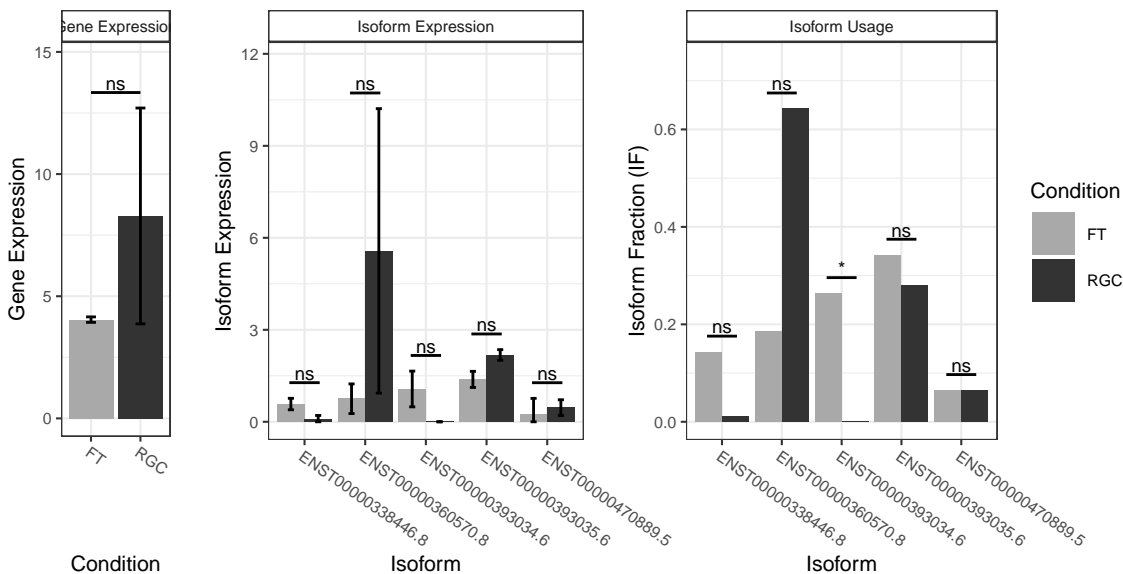
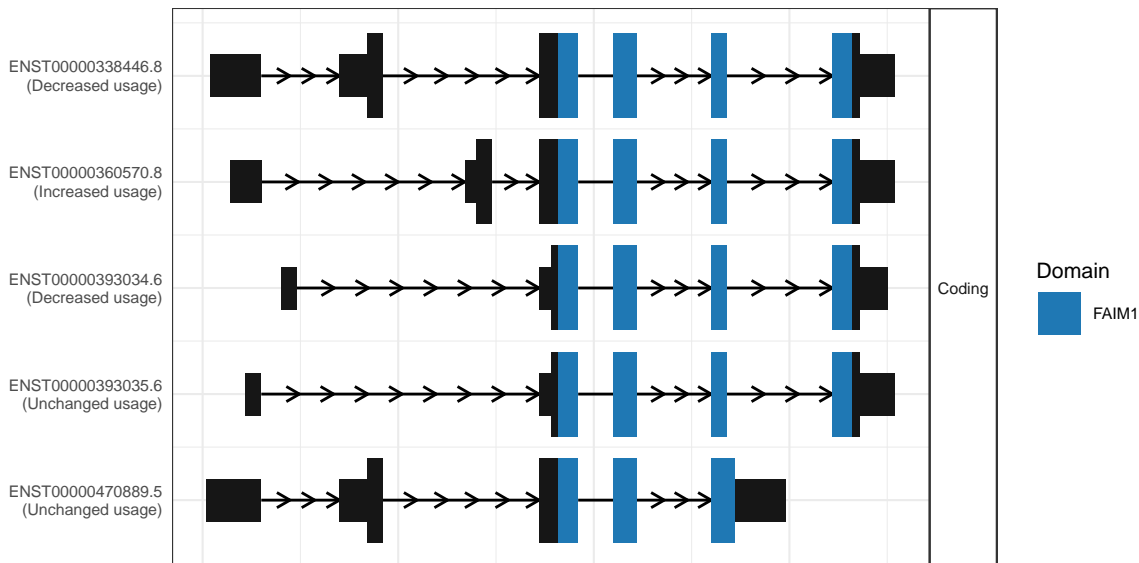


Cofilin_ADF



SH3_9

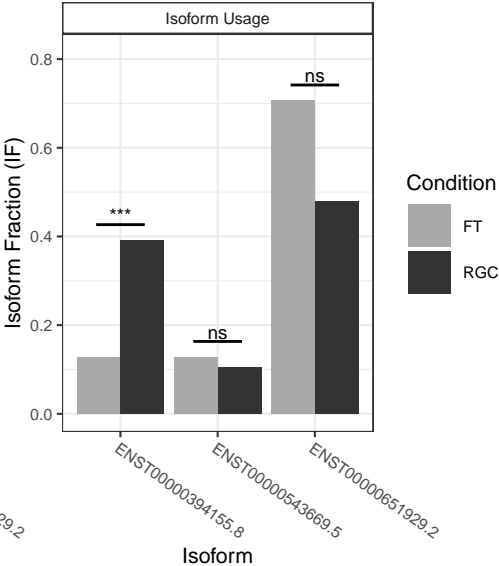
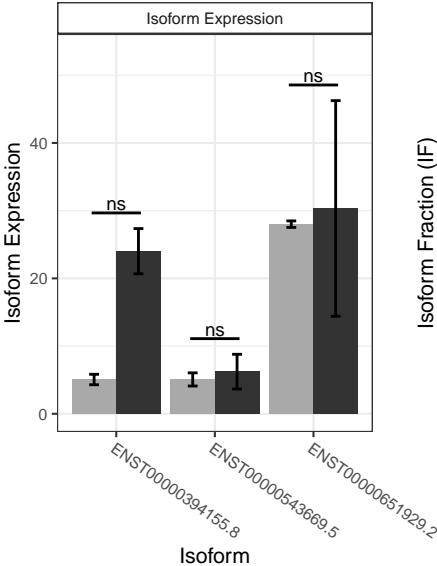
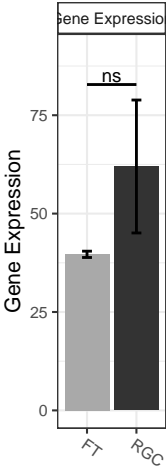
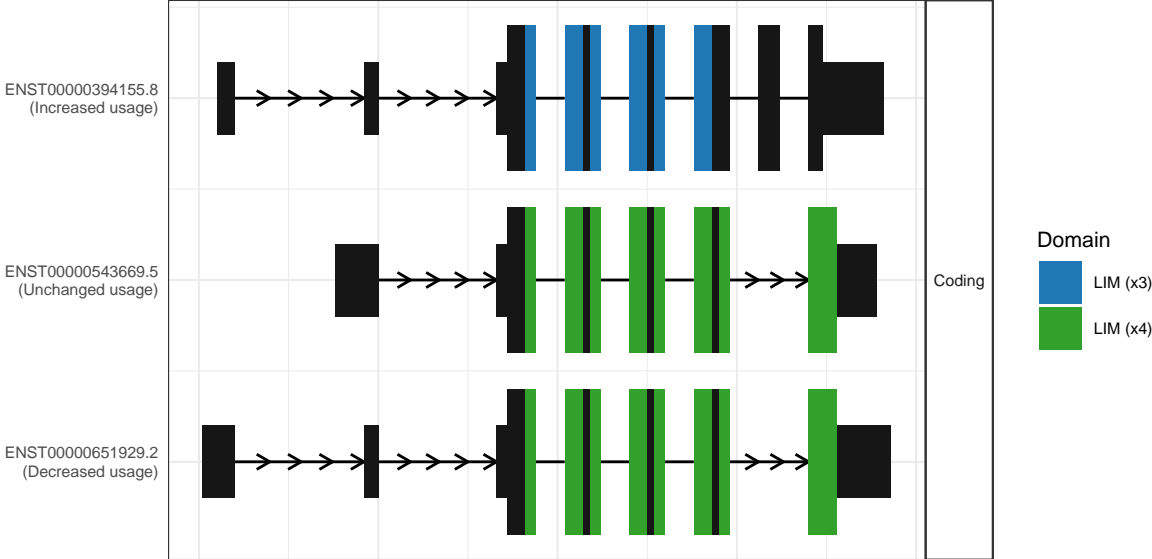
The isoform switch in FAIM (FT vs RGC)



Domain

 FAIM1

The isoform switch in FHL1 (FT vs RGC)



Domain

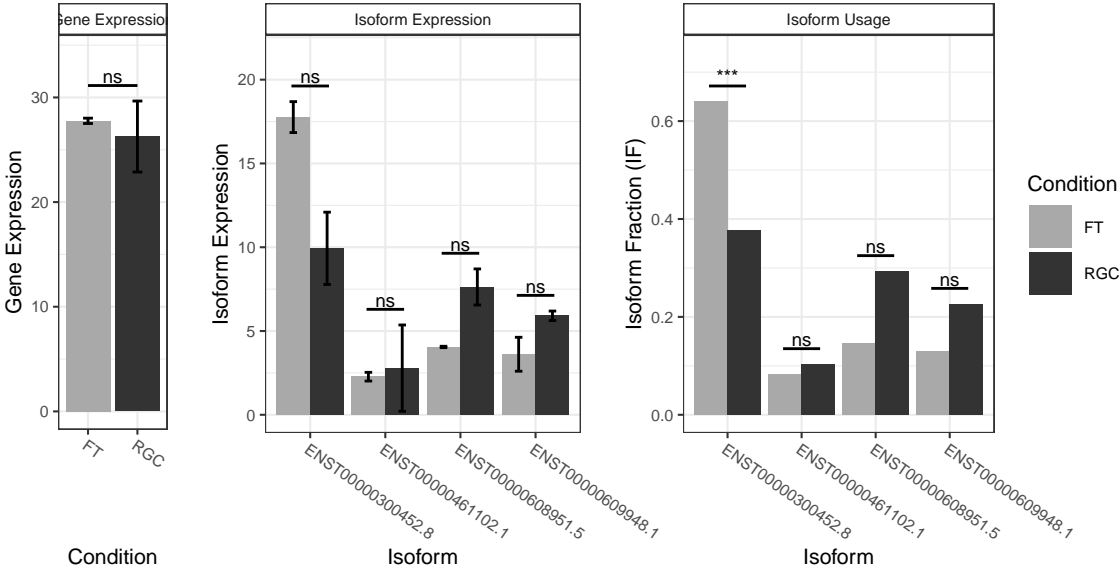
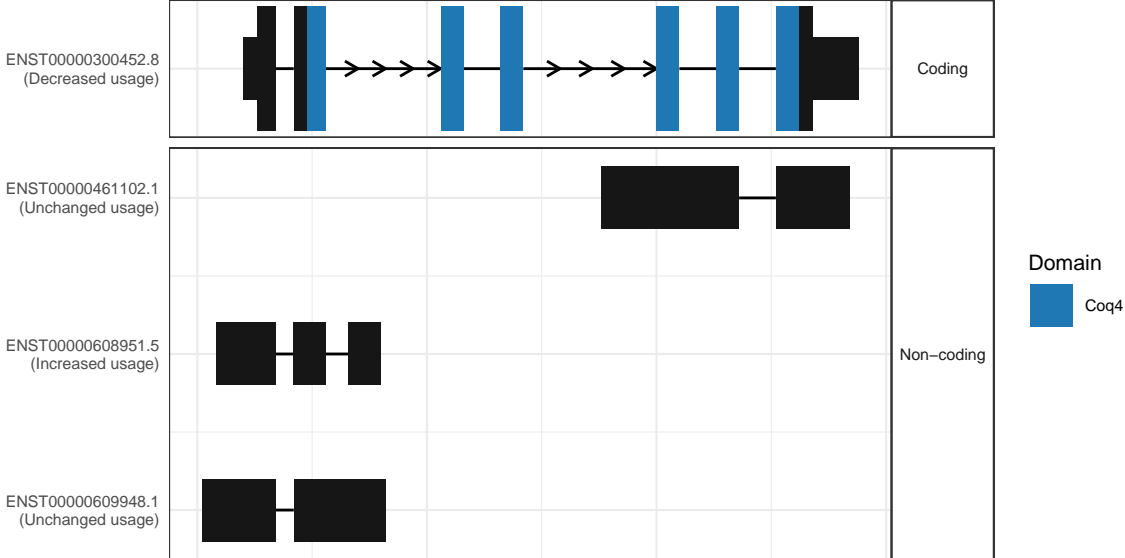


LIM (x3)



LIM (x4)

The isoform switch in COQ4 (FT vs RGC)

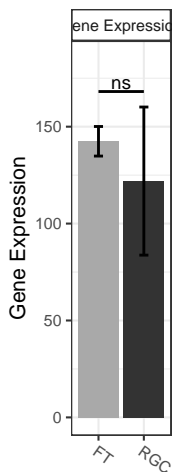
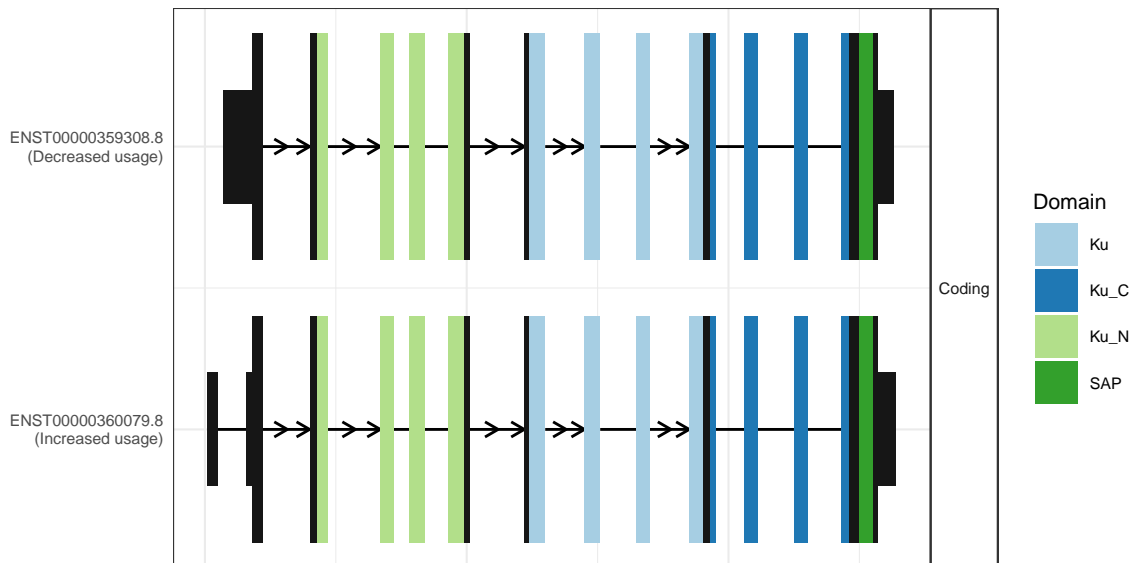


Domain

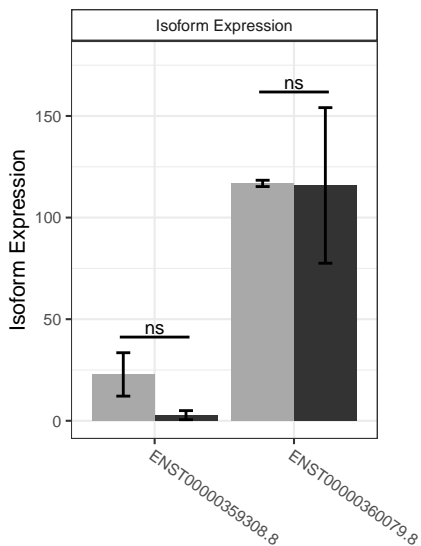


Coq4

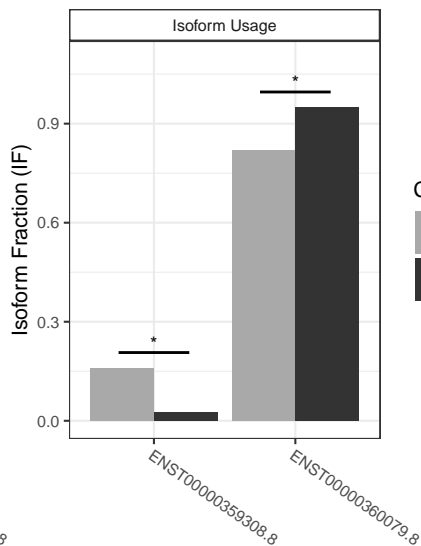
The isoform switch in XRCC6 (FT vs RGC)



Condition



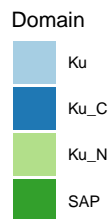
Isoform



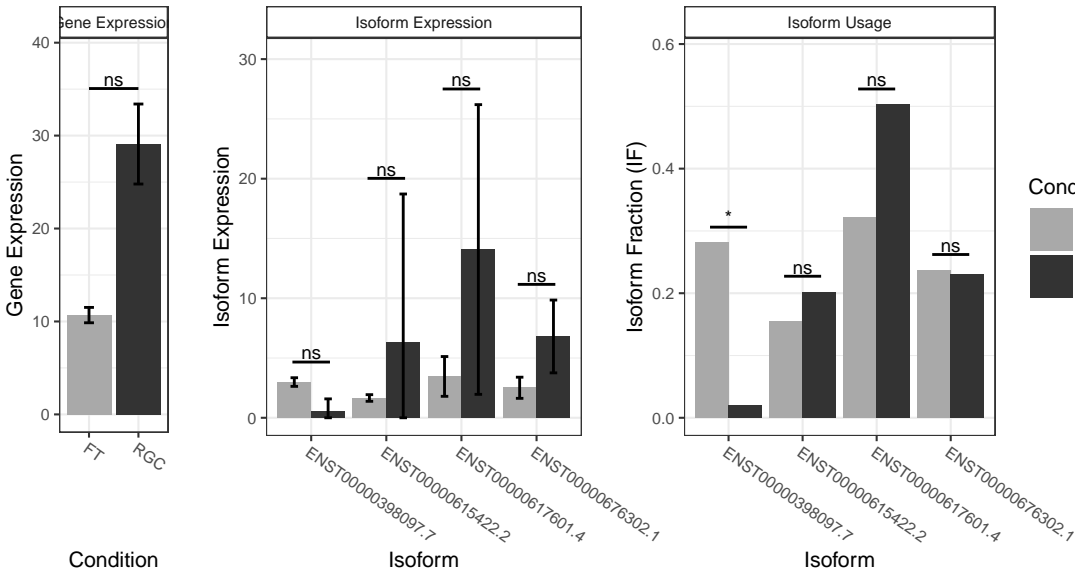
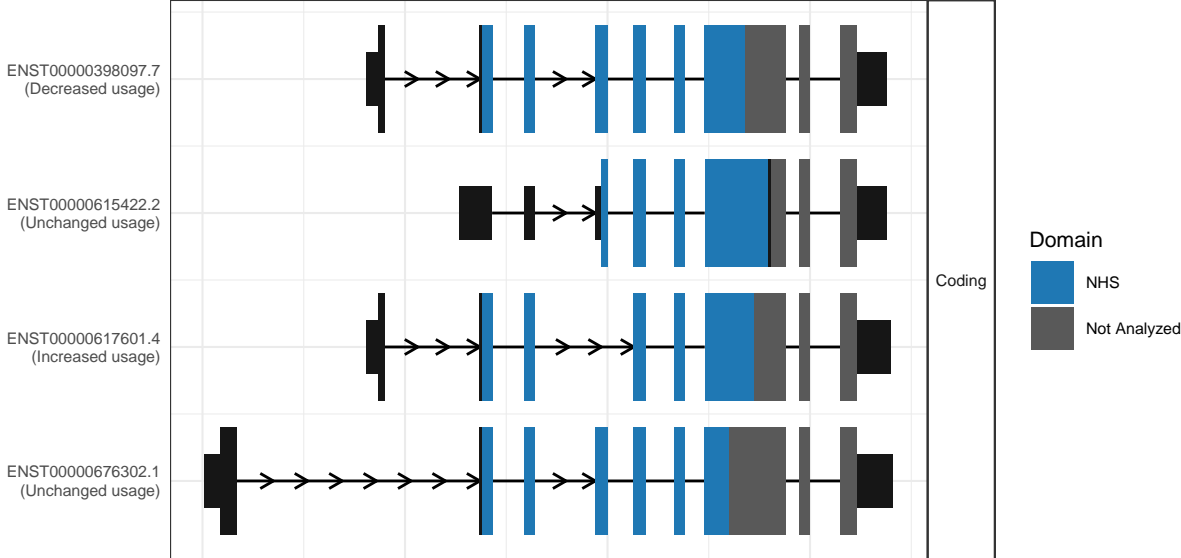
Isoform

Condition





The isoform switch in NHS (FT vs RGC)



Domain



NHS

Not Analyzed

Domain

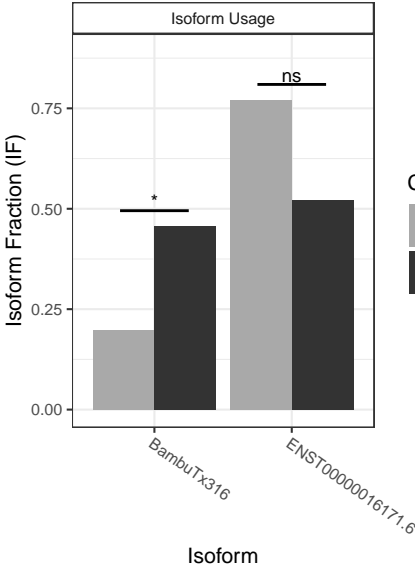
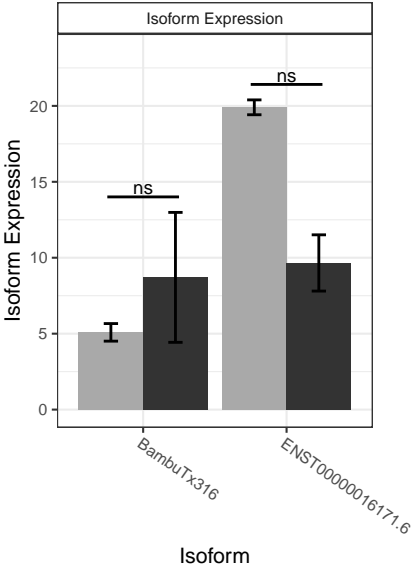
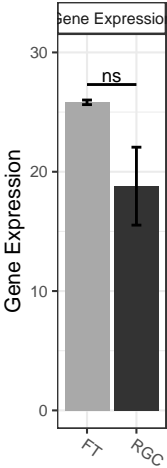
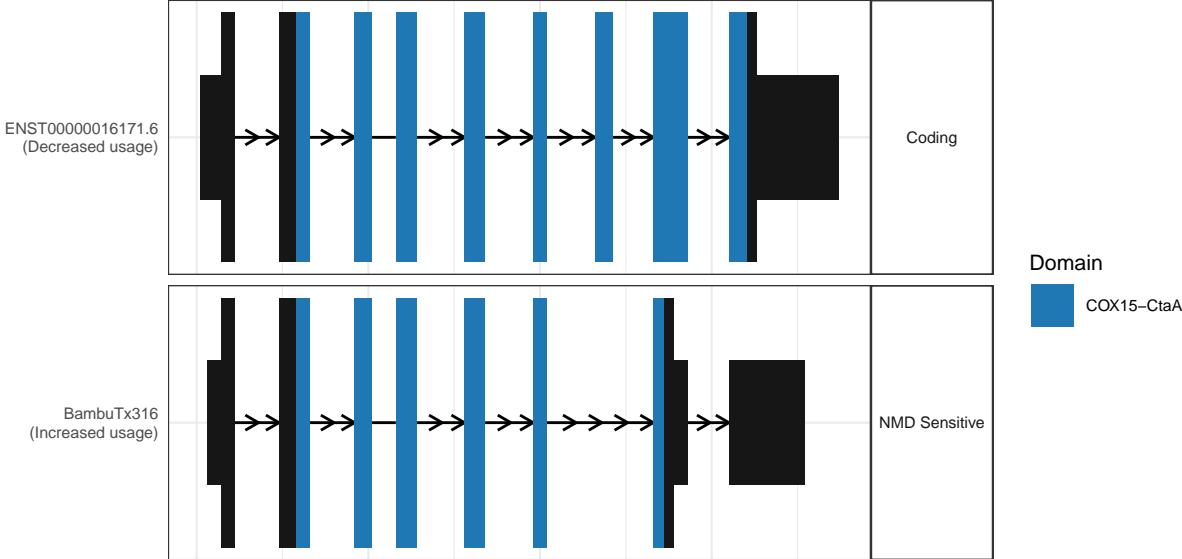


Pkinase (x2)



Pkinase_C

The isoform switch in COX15 (FT vs RGC)

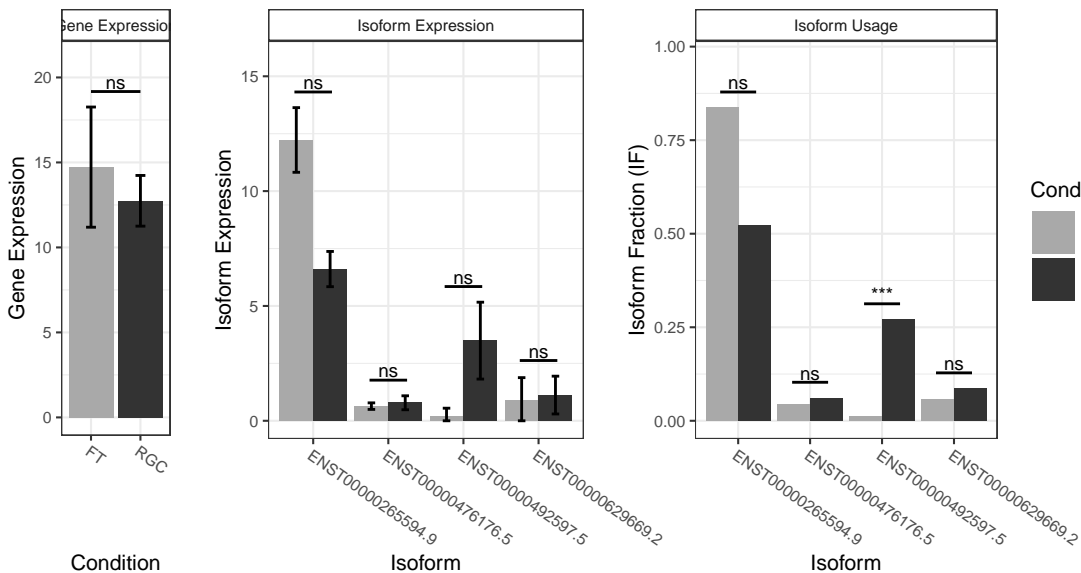
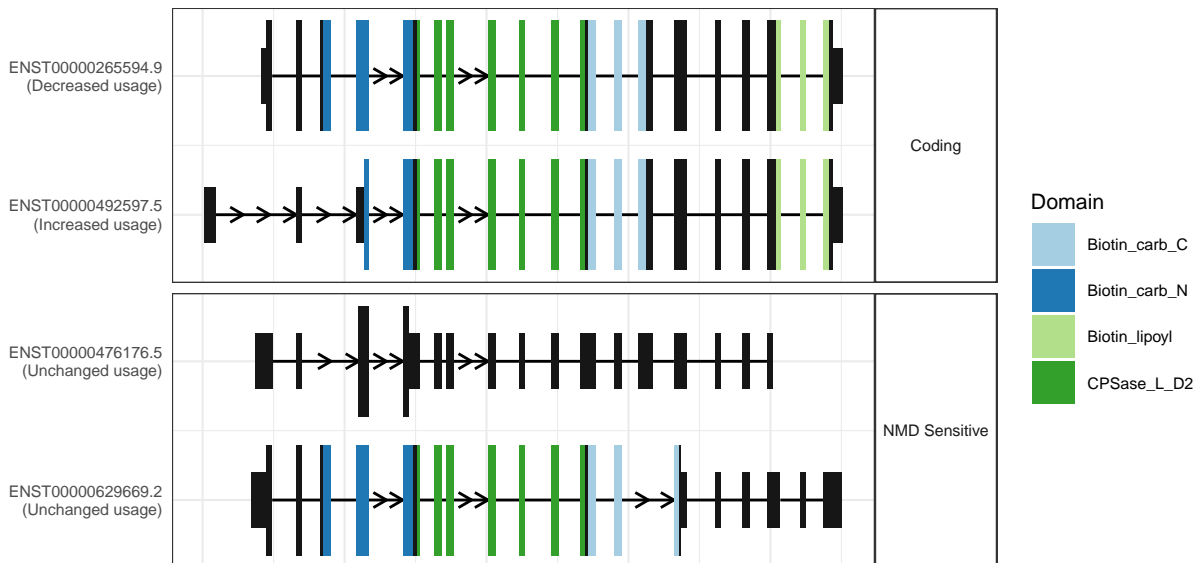


Domain







COX15-CtaA

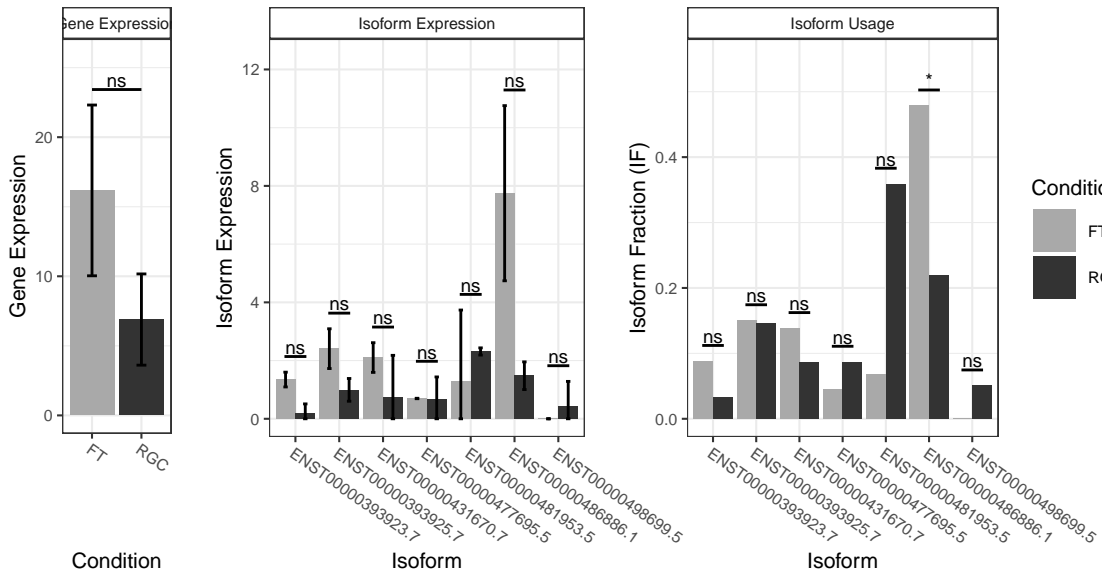
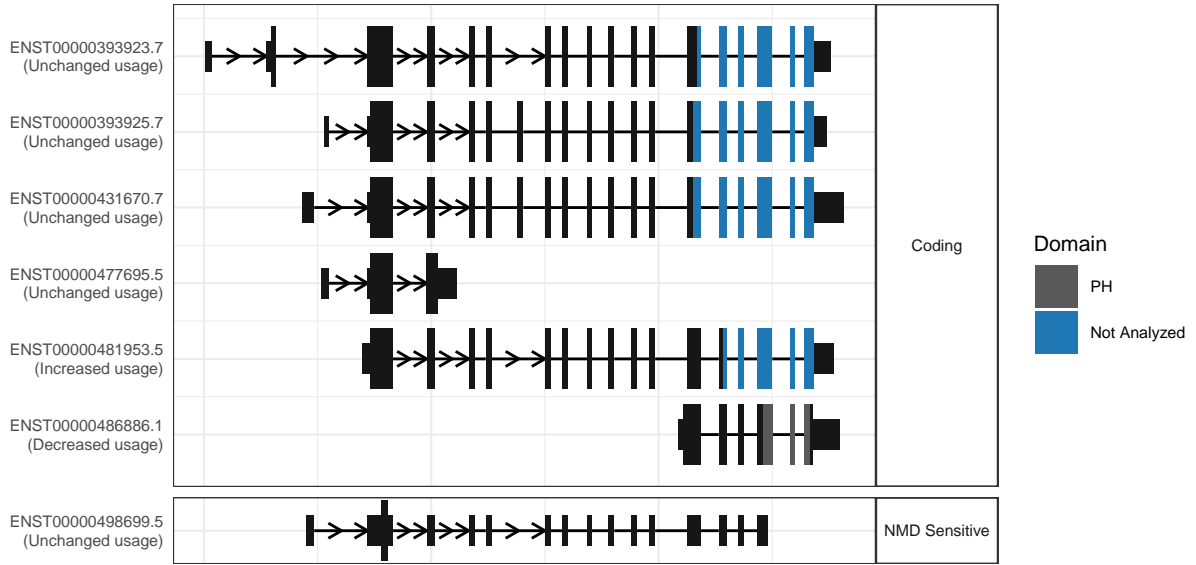
The isoform switch in MCCC1 (FT vs RGC)



Domain

	Biotin_carb_C
	Biotin_carb_N
	Biotin_lipoyl
	CPSase_L_D2

The isoform switch in PHLDB2 (FT vs RGC)



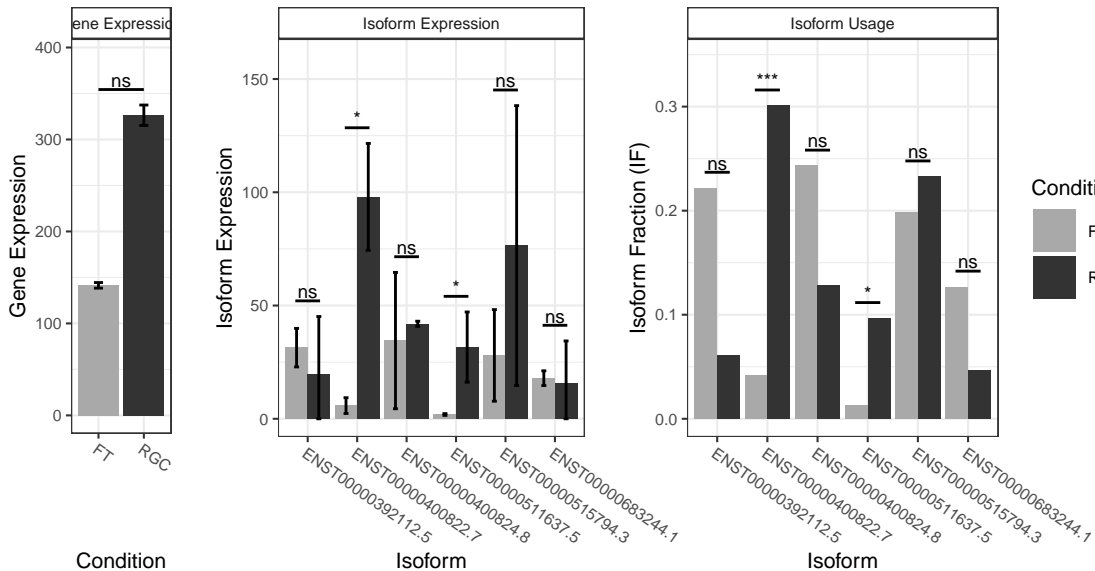
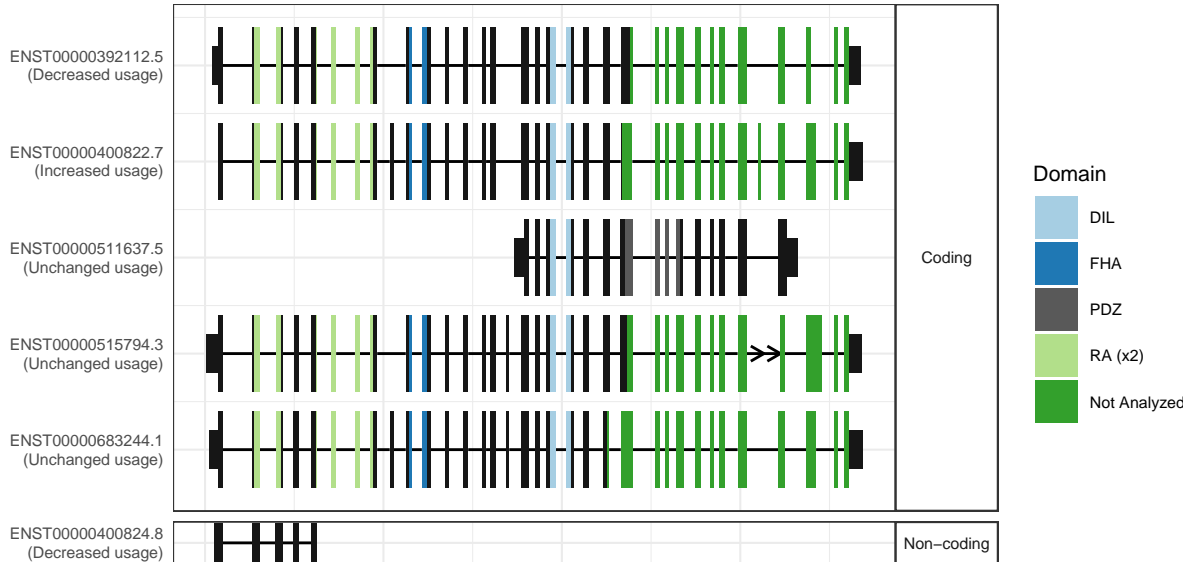
Domain



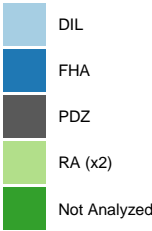
PH

Not Analyzed

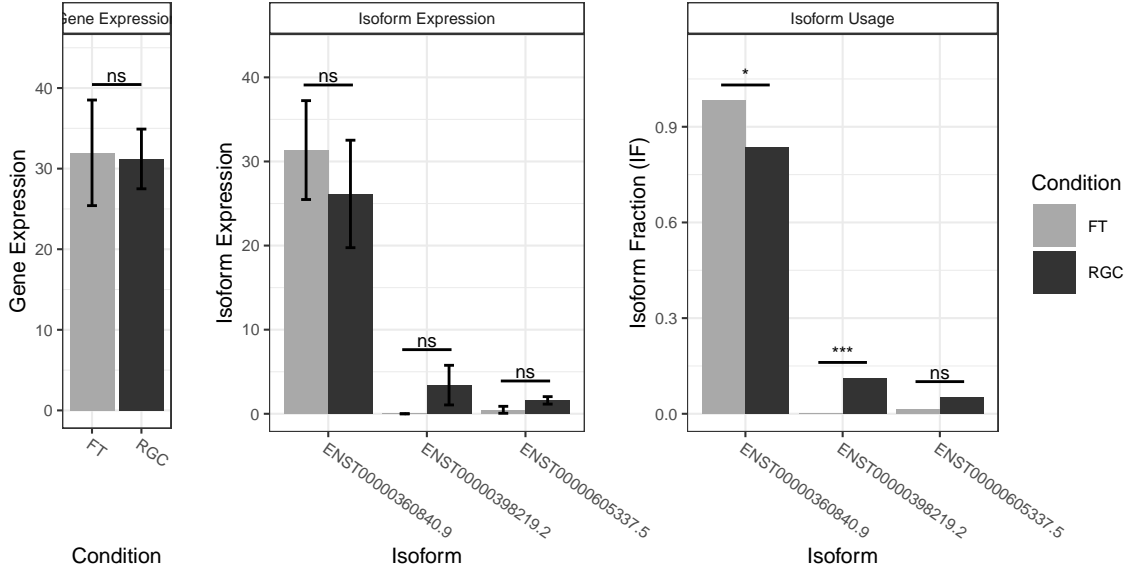
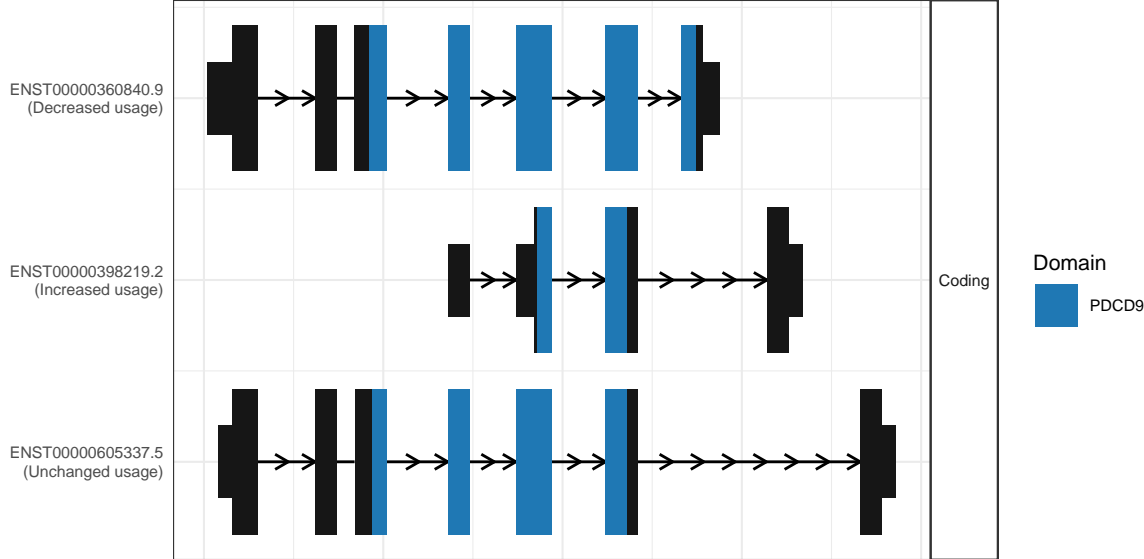
The isoform switch in AFDN (FT vs RGC)



Domain



The isoform switch in MRPL37 (FT vs RGC)

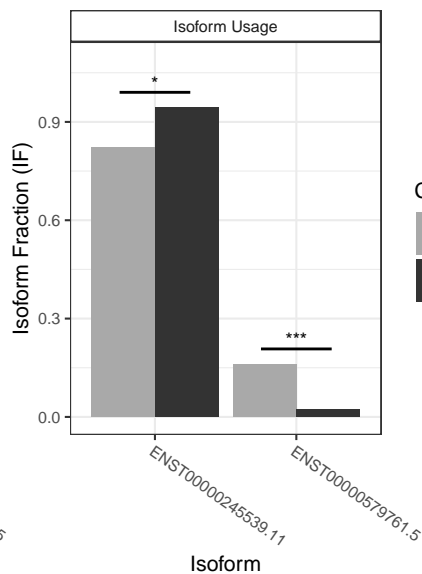
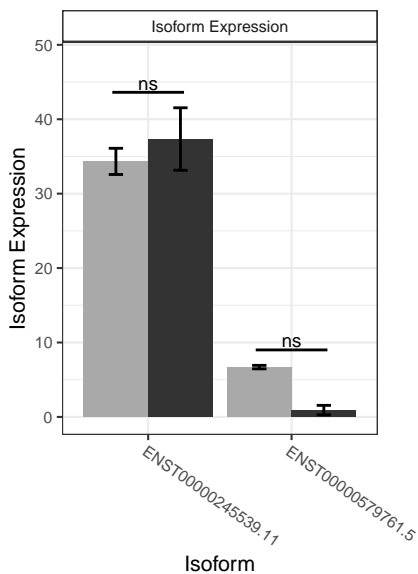
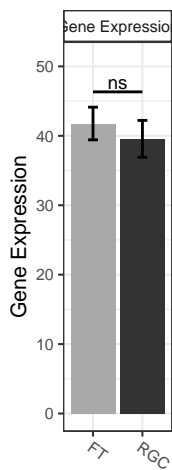
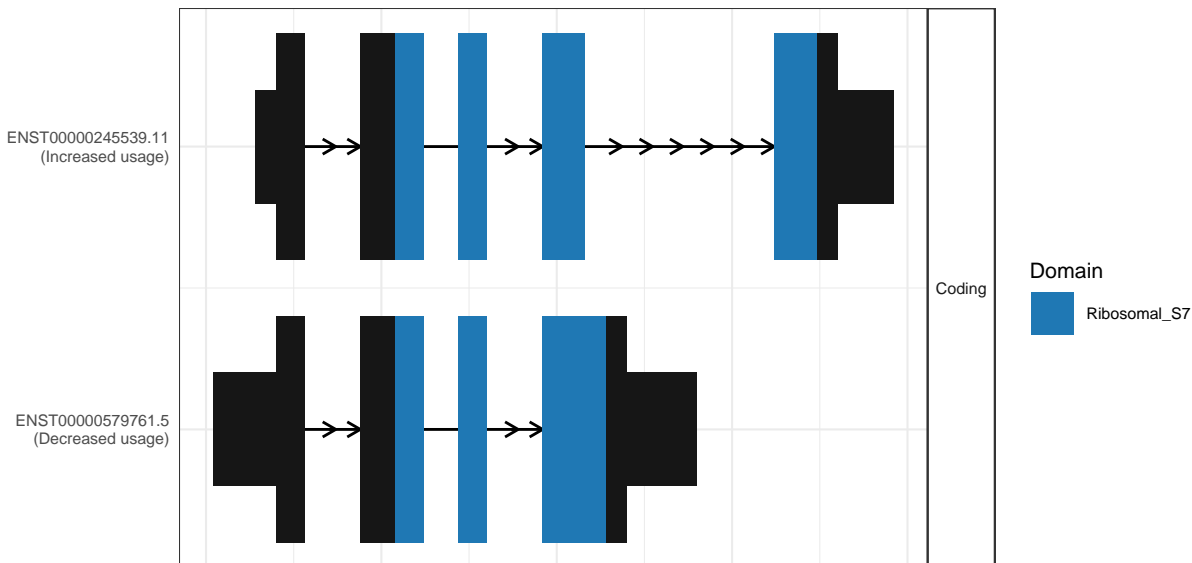


Domain



PDCD9

The isoform switch in MRPS7 (FT vs RGC)

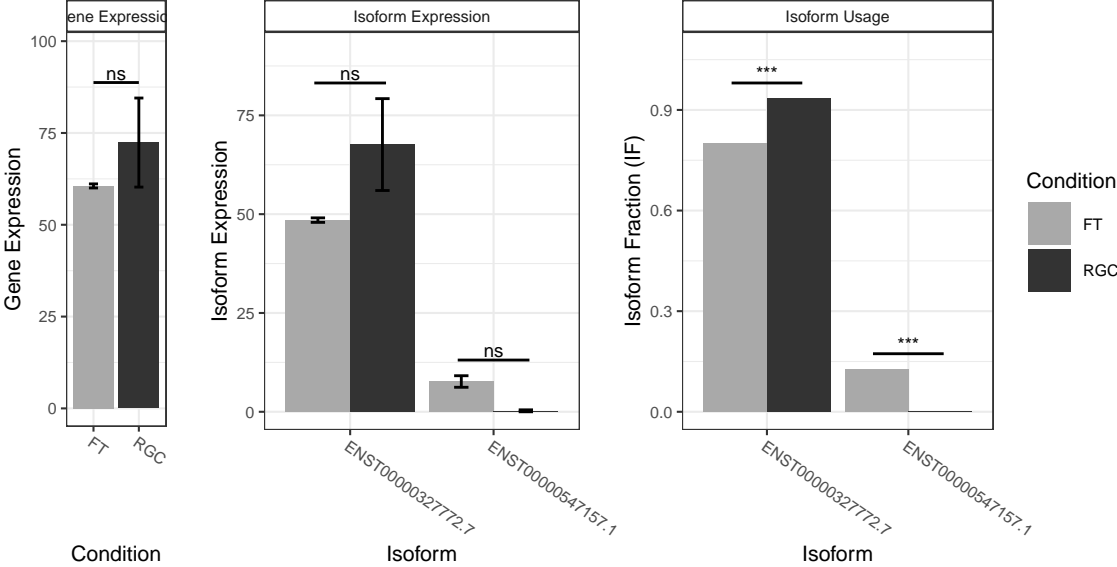
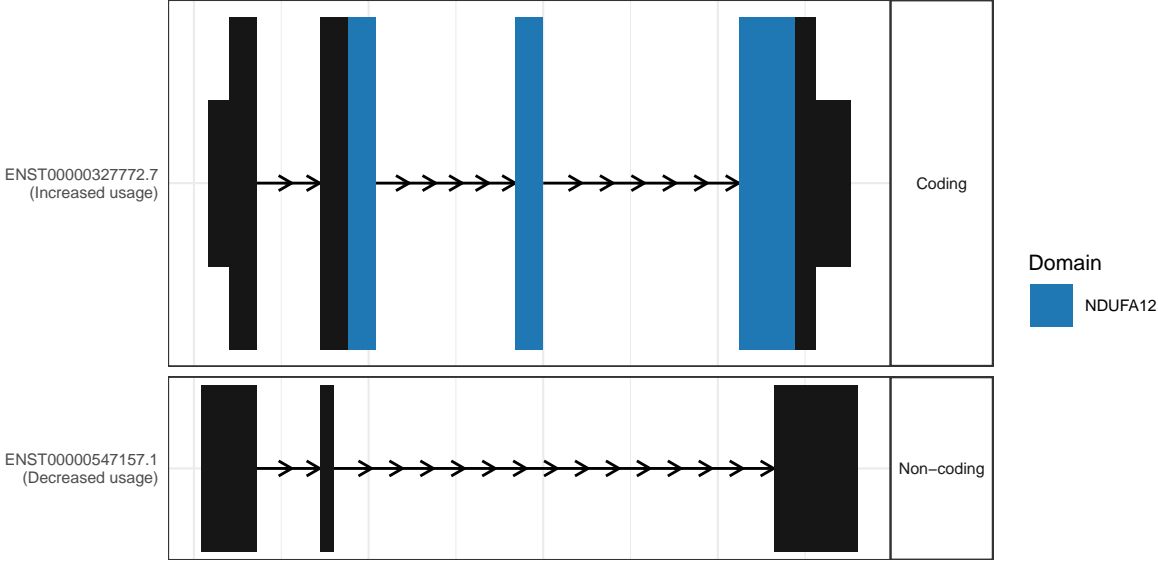


Domain



Ribosomal_S7

The isoform switch in NDUFA12 (FT vs RGC)

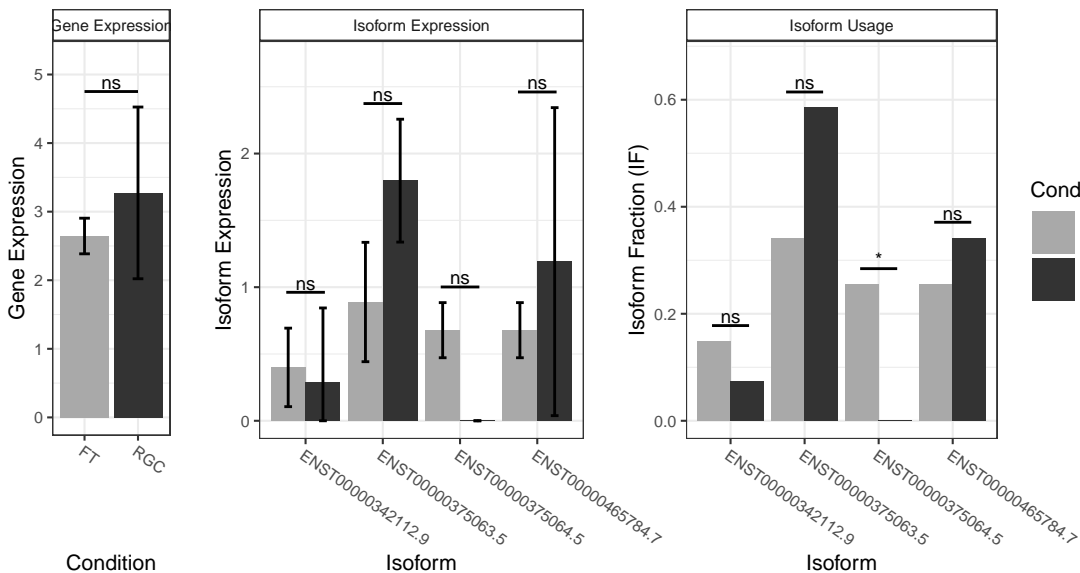
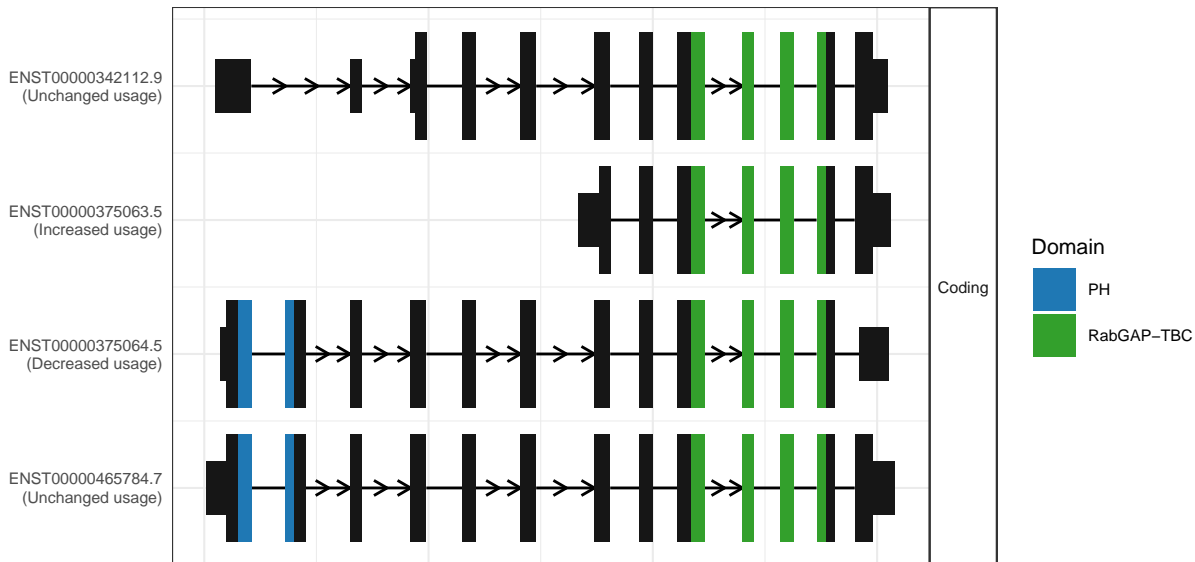


Domain



NDUFA12

The isoform switch in TBC1D2 (FT vs RGC)



Domain

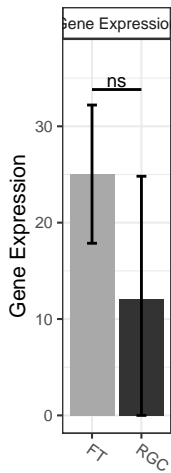
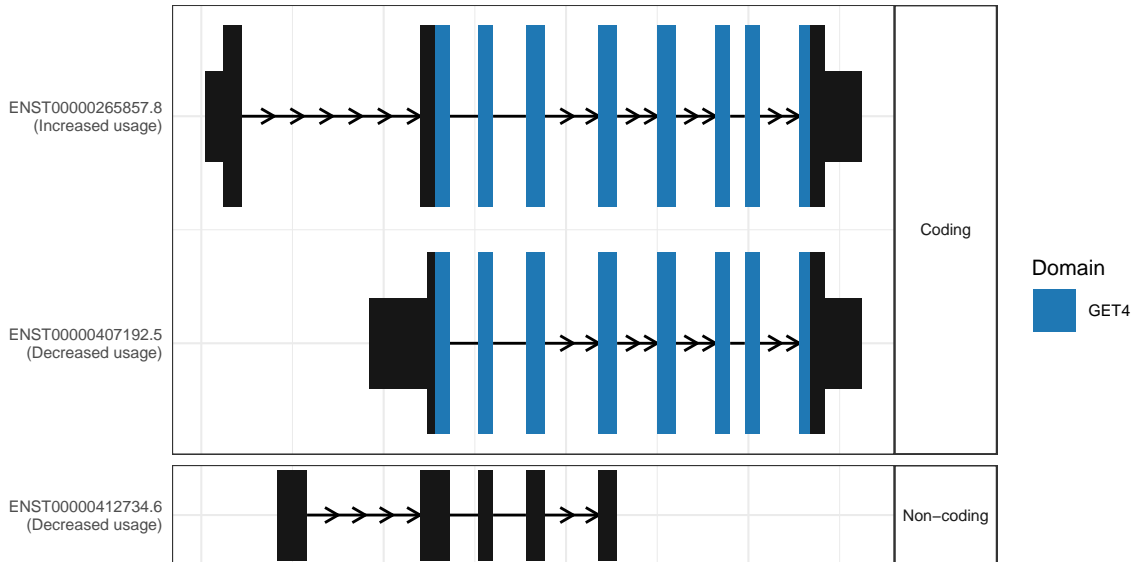


PH

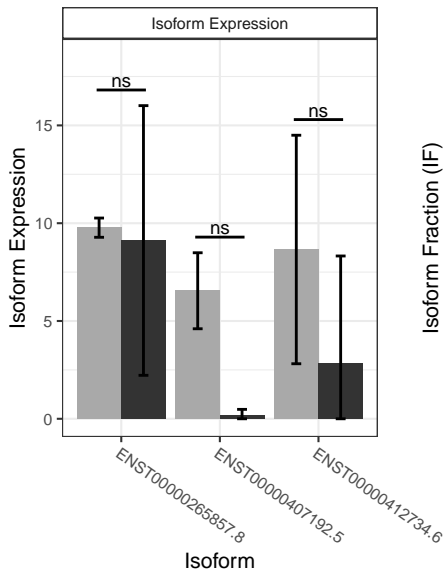


RabGAP-TBC

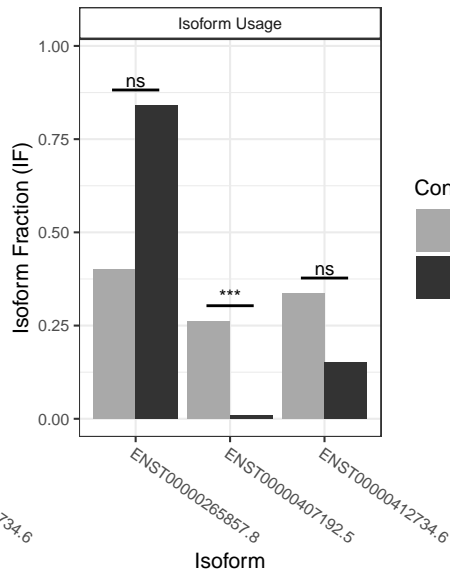
The isoform switch in GET4 (FT vs RGC)



Condition



Isoform



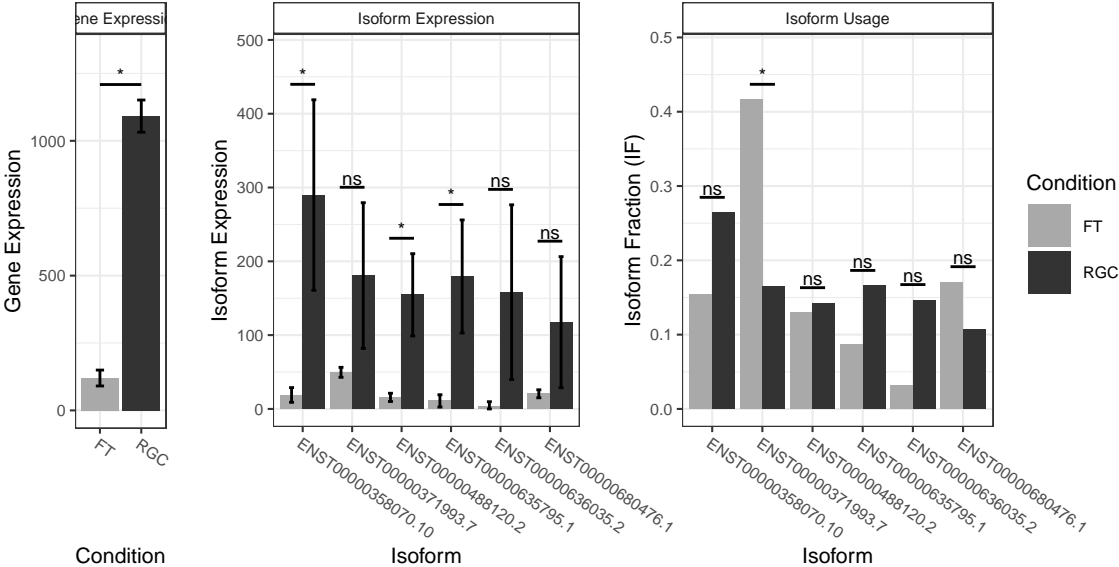
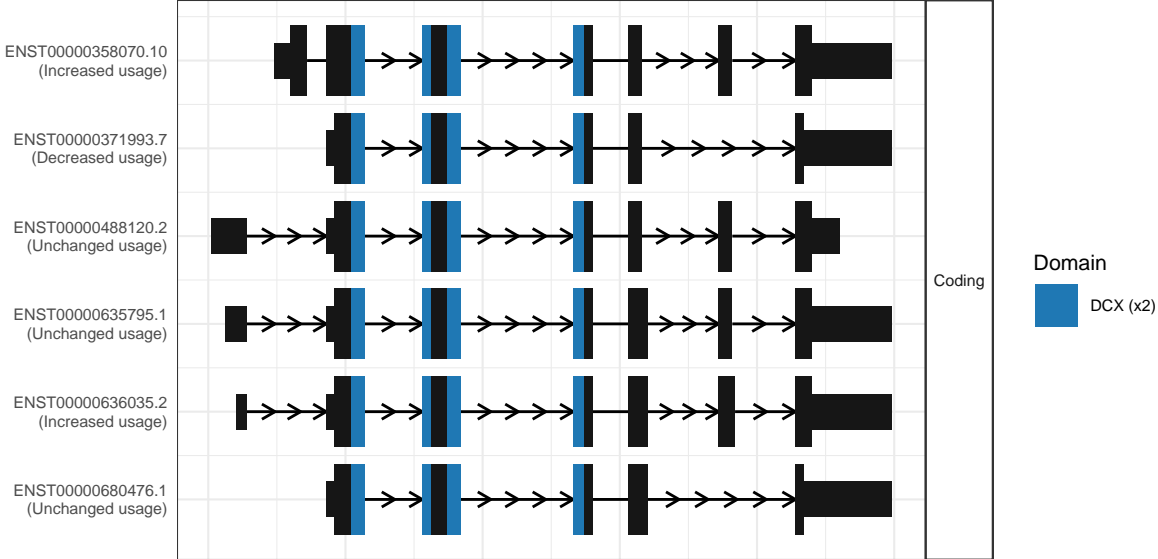
Isoform

Domain



GET4

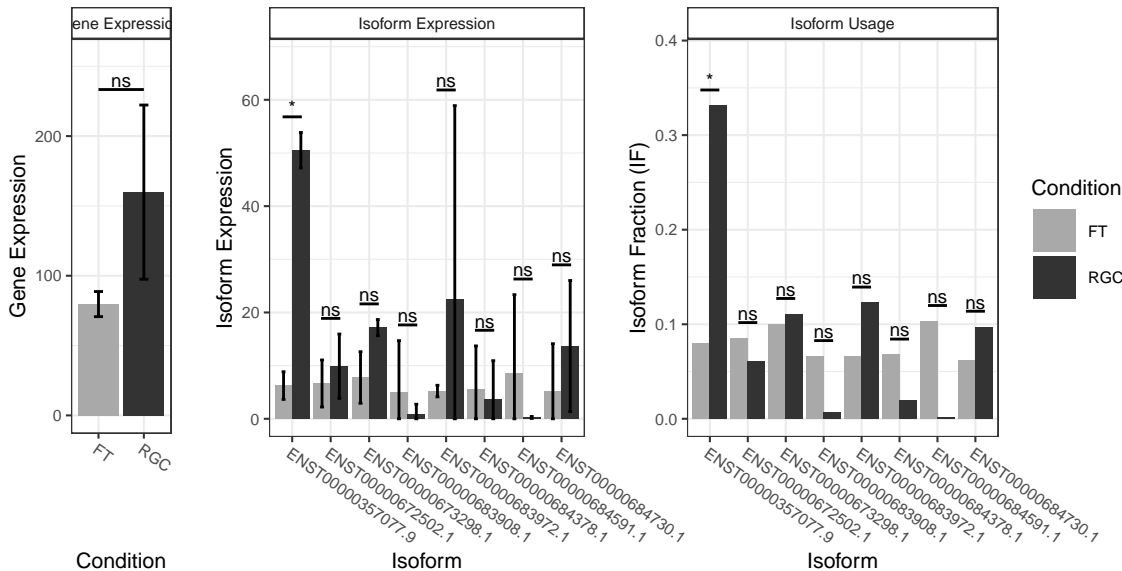
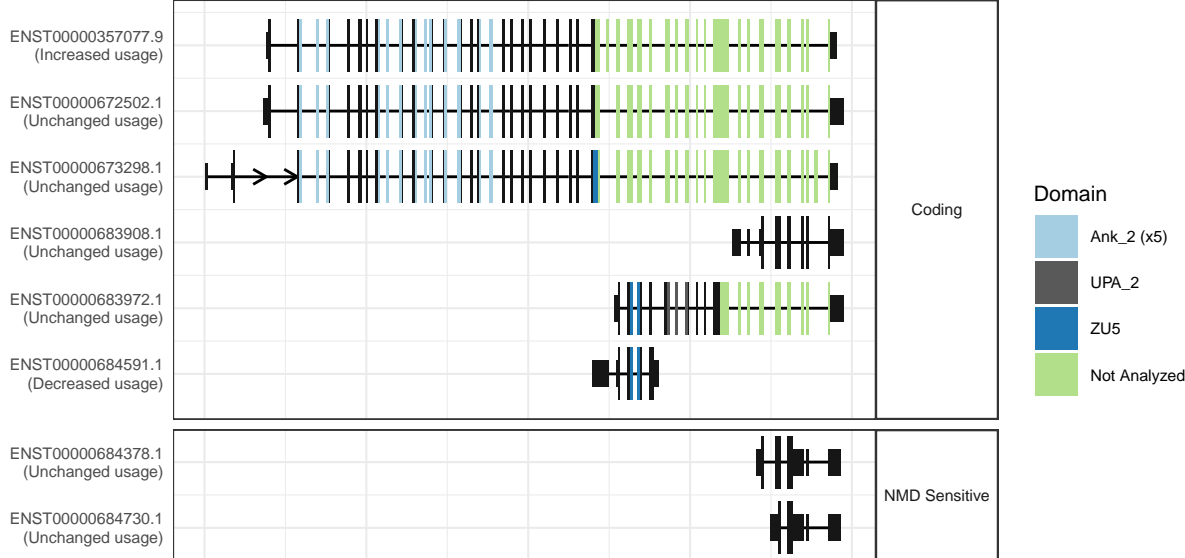
The isoform switch in DCX (FT vs RGC)



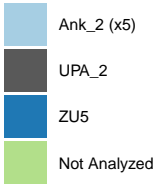
Domain

 DCX (x2)

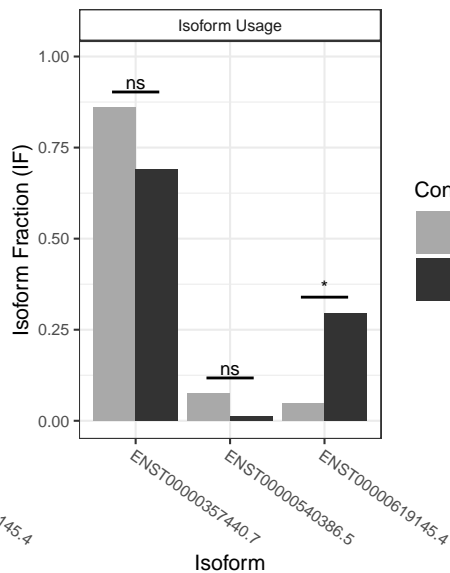
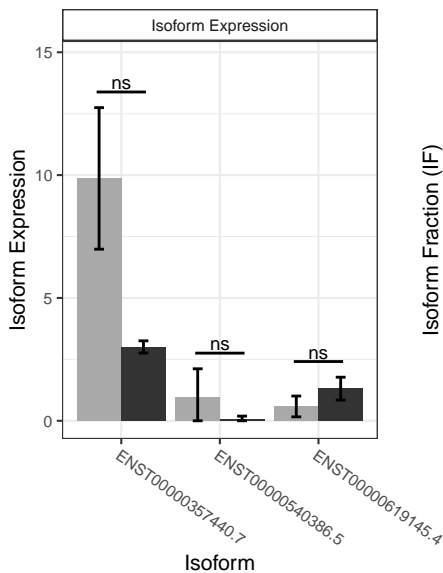
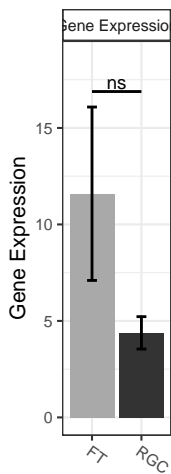
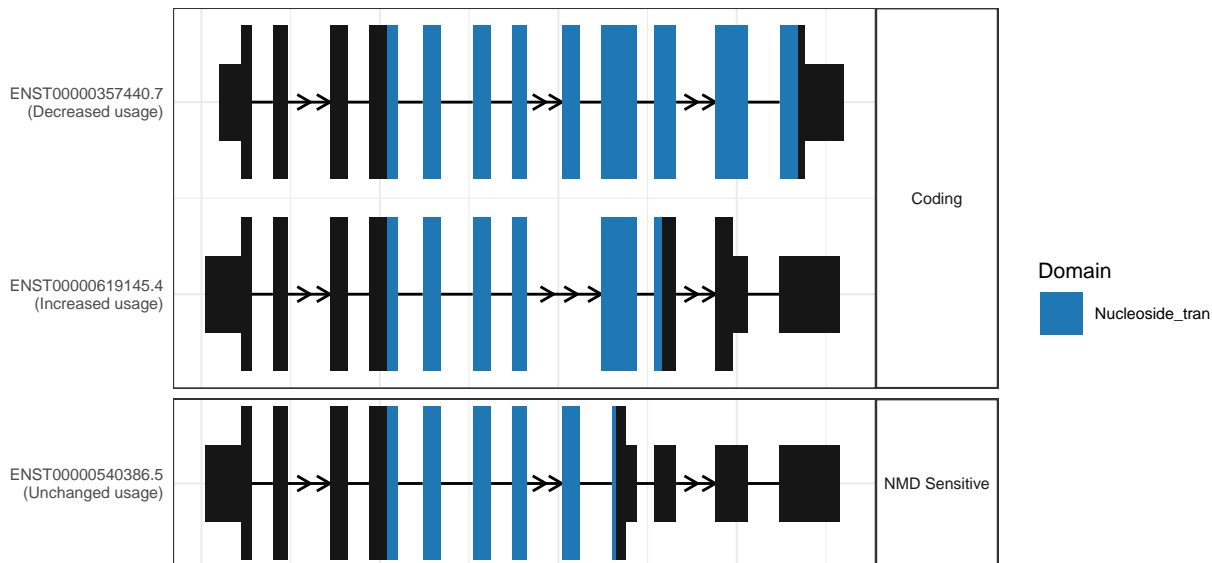
The isoform switch in ANK2 (FT vs RGC)



Domain



The isoform switch in SLC29A2 (FT vs RGC)

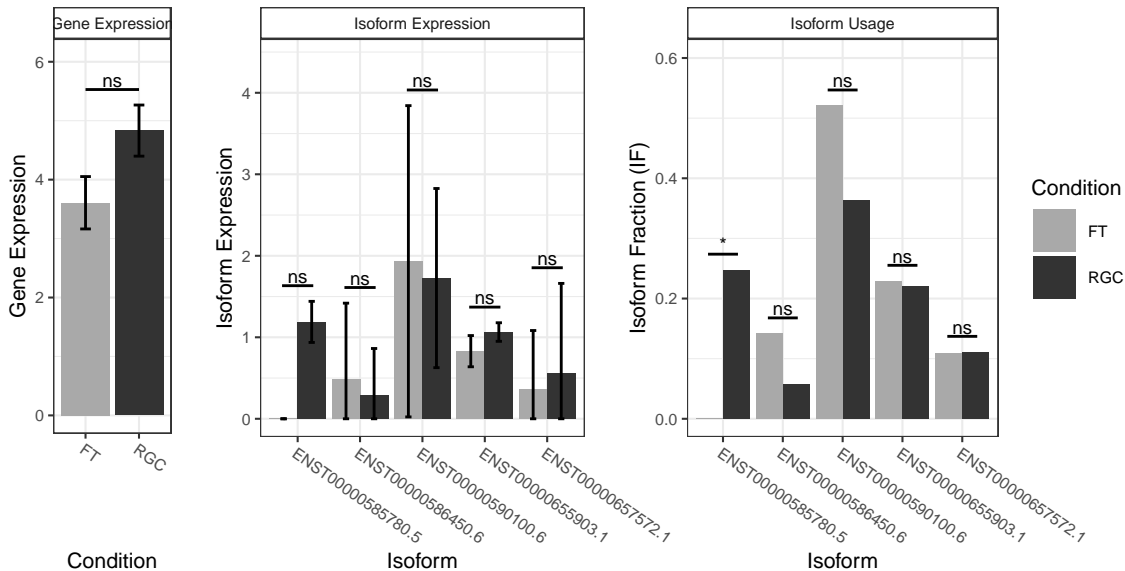
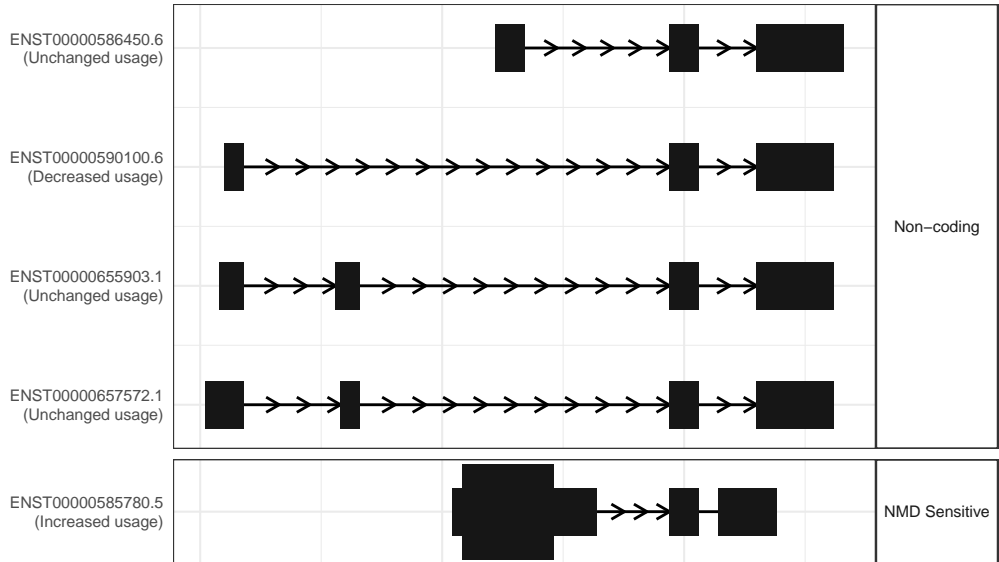


Domain

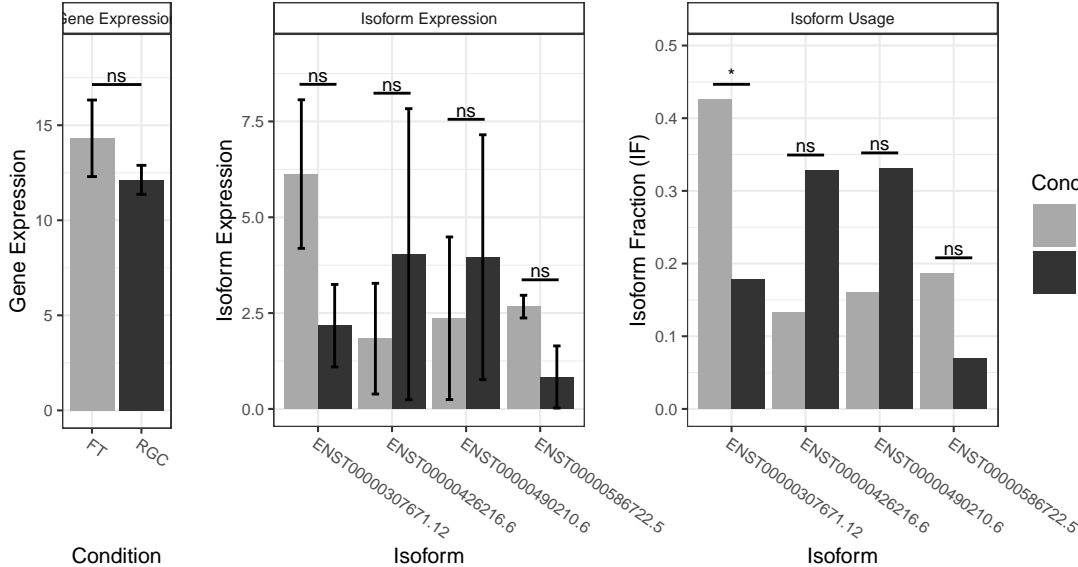
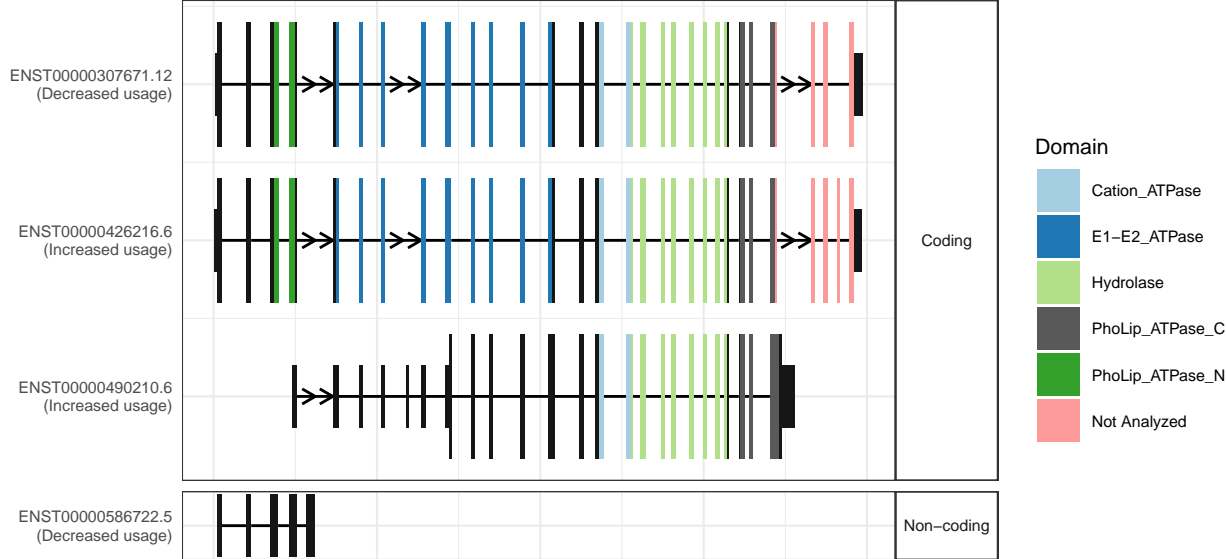


Nucleoside_tran







The isoform switch in MAP3K14-AS1 (FT vs RGC)



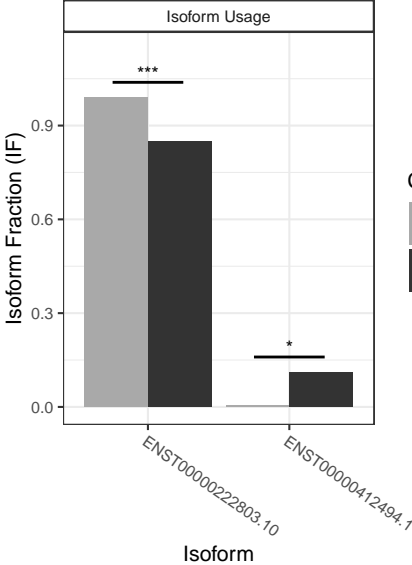
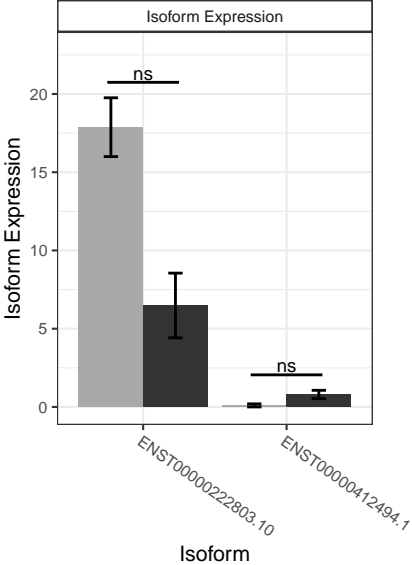
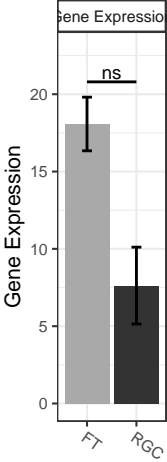
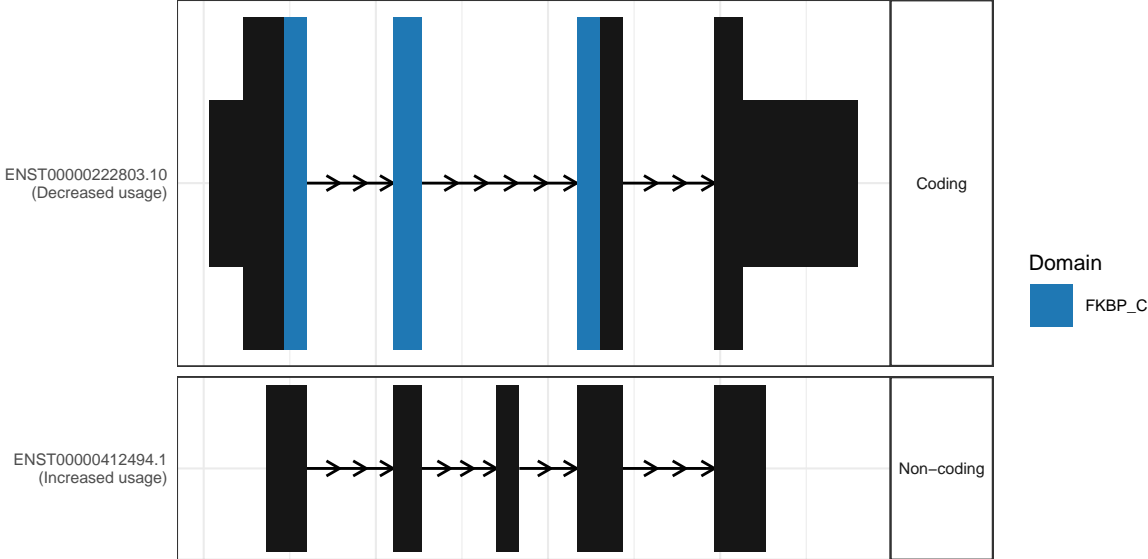
The isoform switch in ATP9B (FT vs RGC)



Domain

	Cation_ATPase
	E1-E2_ATPase
	Hydrolase
	PhoLip_ATPase_C
	PhoLip_ATPase_N
	Not Analyzed

The isoform switch in FKBP14 (FT vs RGC)

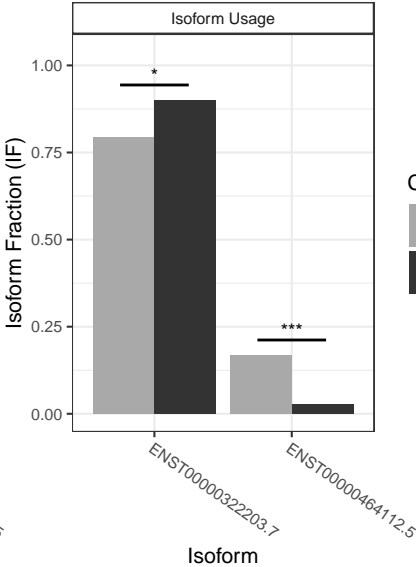
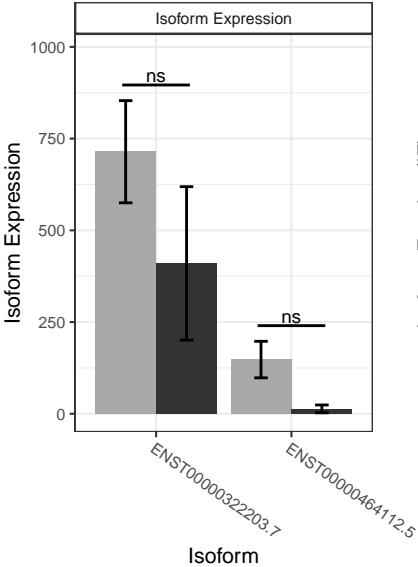
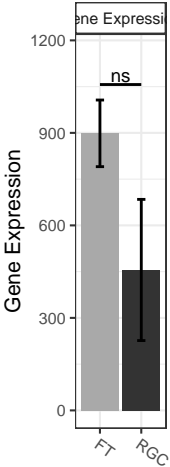
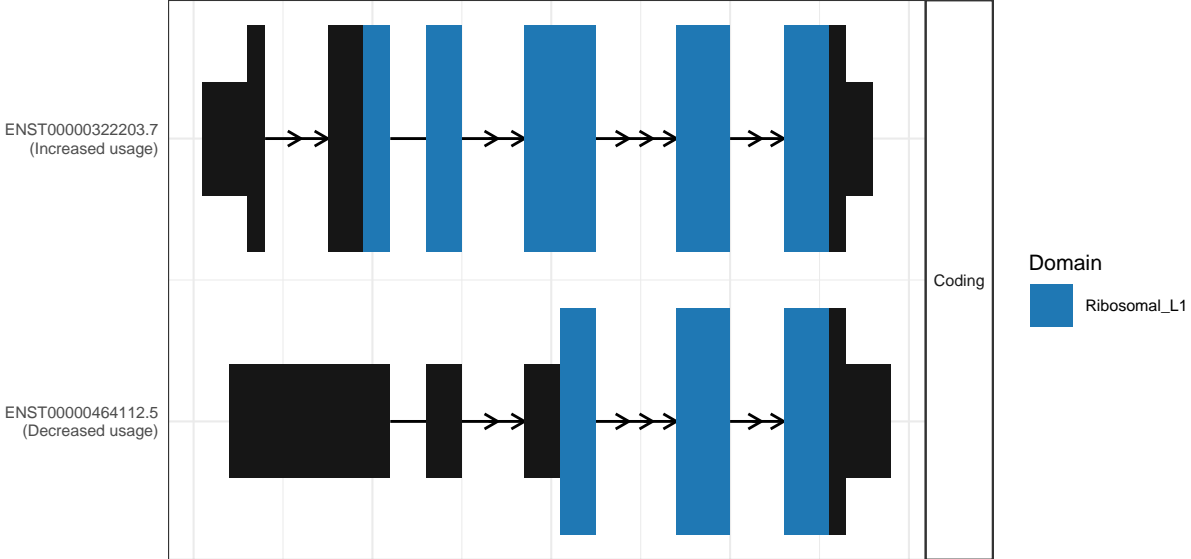


Domain



FKBP_C

The isoform switch in RPL10A (FT vs RGC)

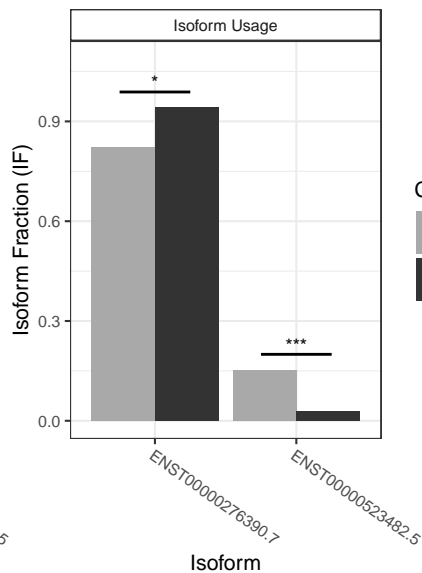
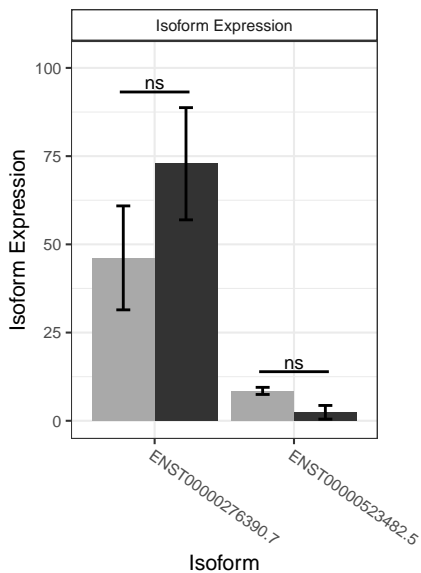
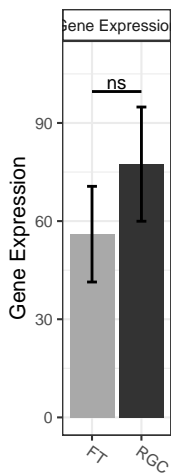
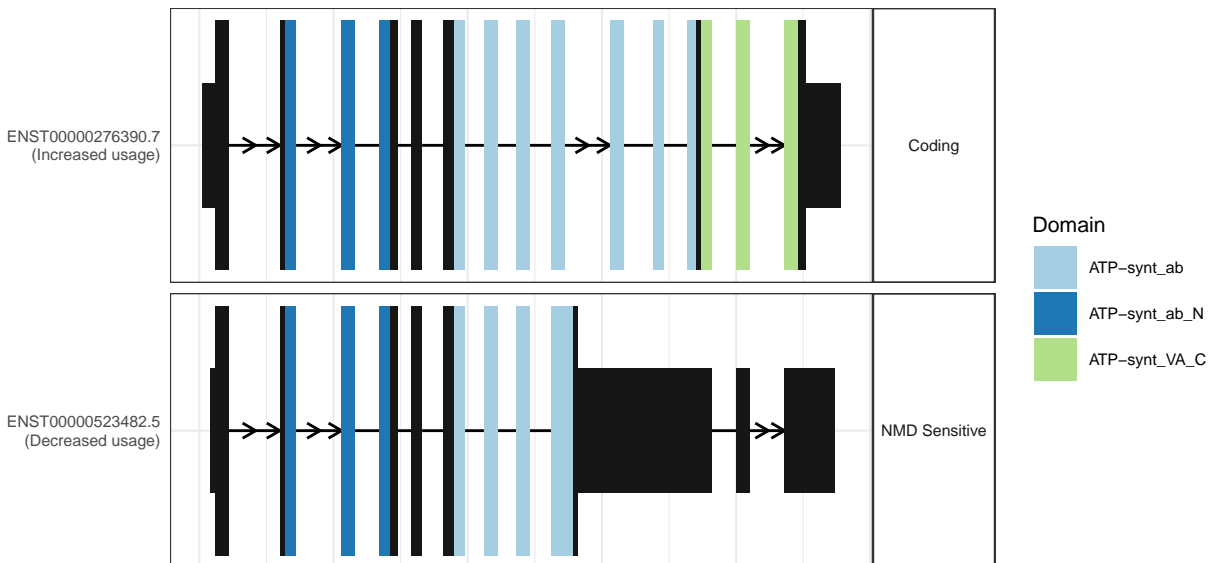


Domain



Ribosomal_L1

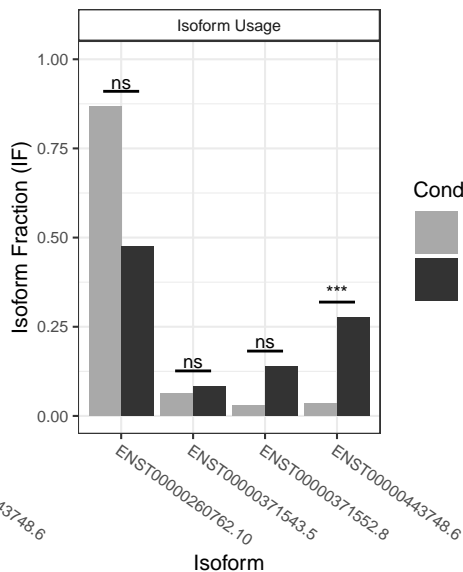
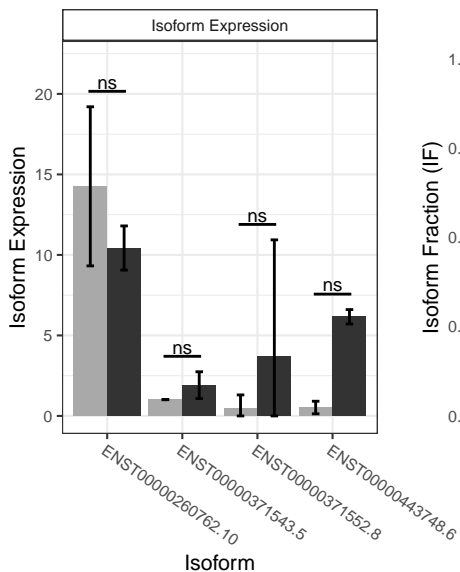
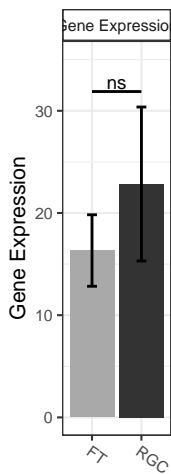
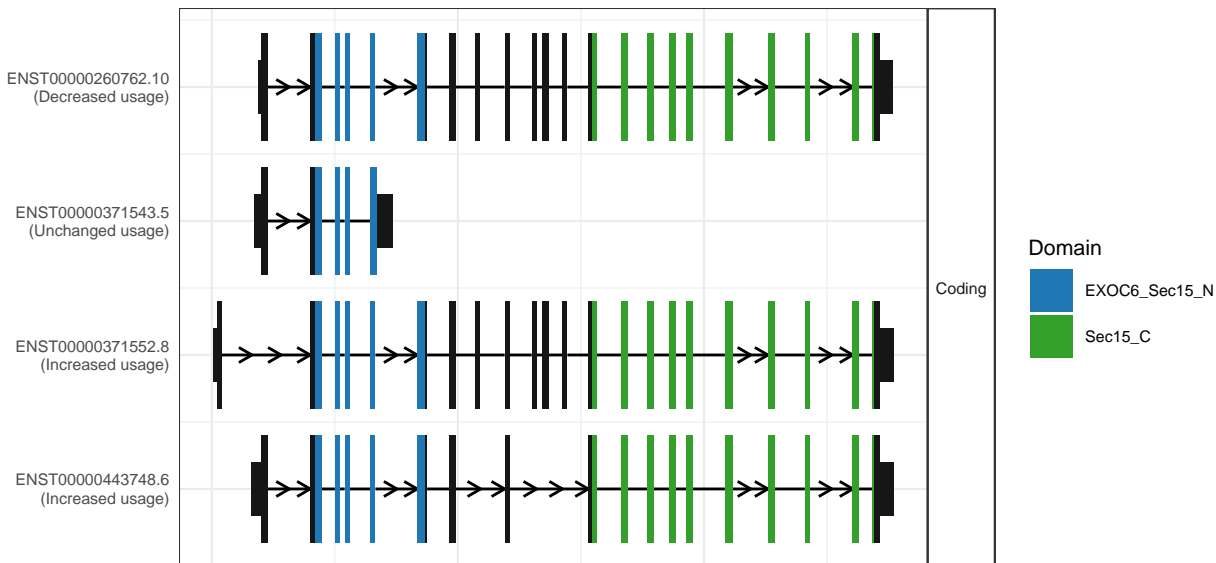
The isoform switch in ATP6V1B2 (FT vs RGC)



Domain

	ATP-synt_ab
	ATP-synt_ab_N
	ATP-synt_VA_C

The isoform switch in EXOC6 (FT vs RGC)



Condition

Isoform

Isoform

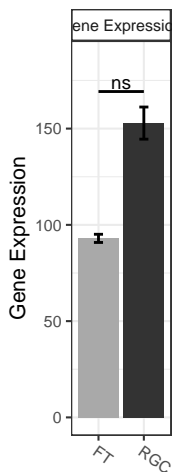
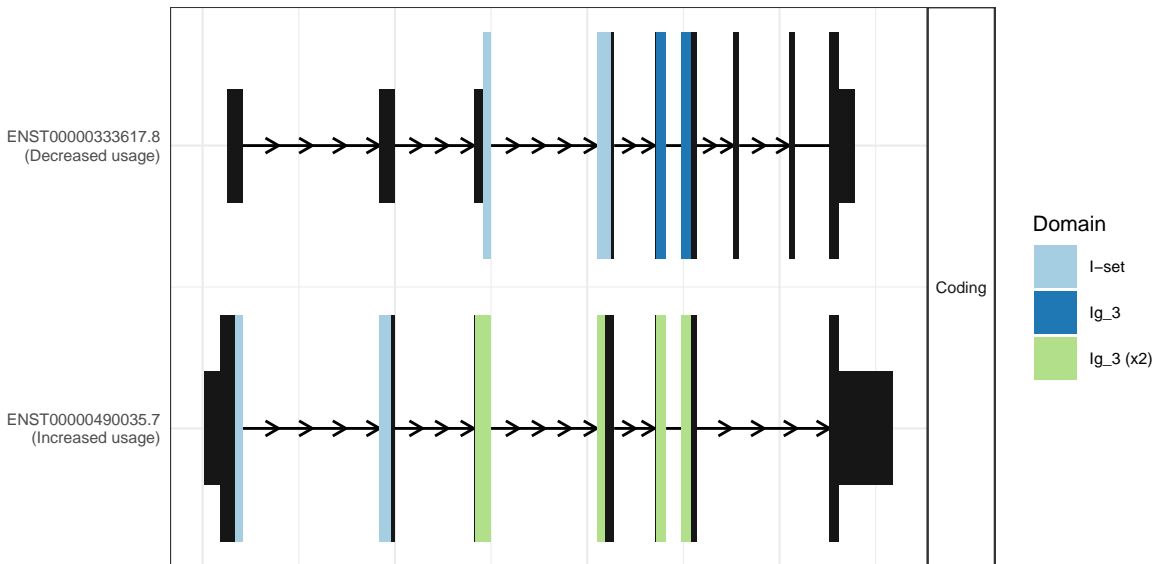
Domain



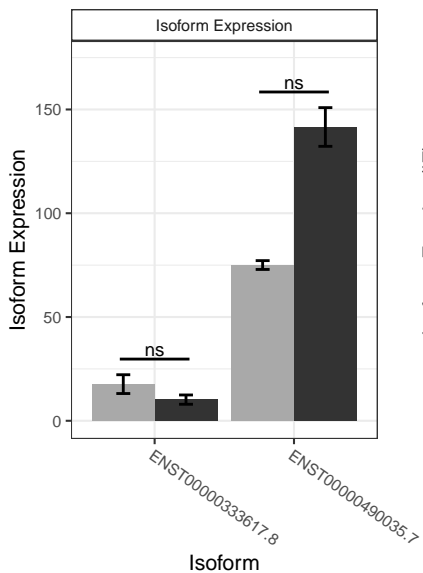
EXOC6_Sec15_N

Sec15_C

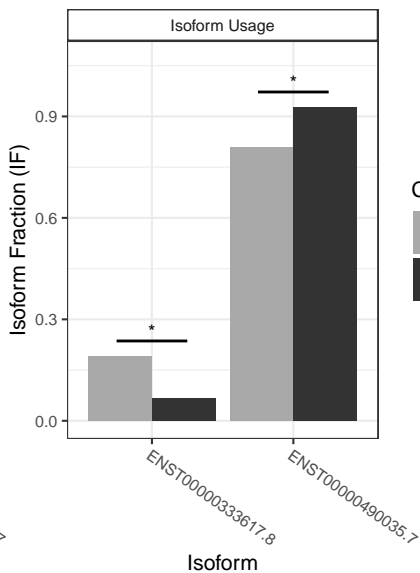
The isoform switch in LSAMP (FT vs RGC)



Condition



Isoform



Isoform

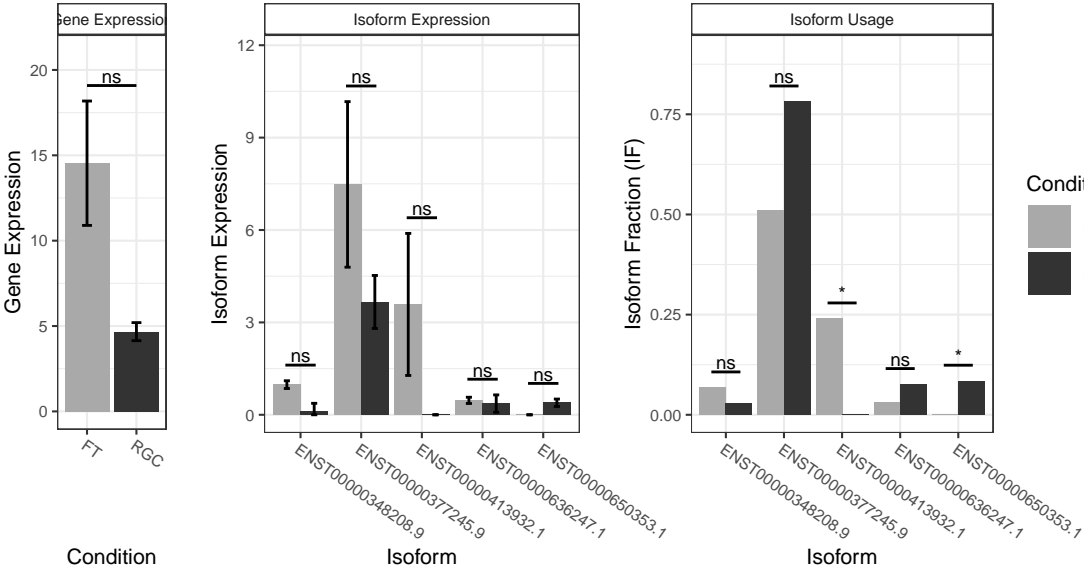
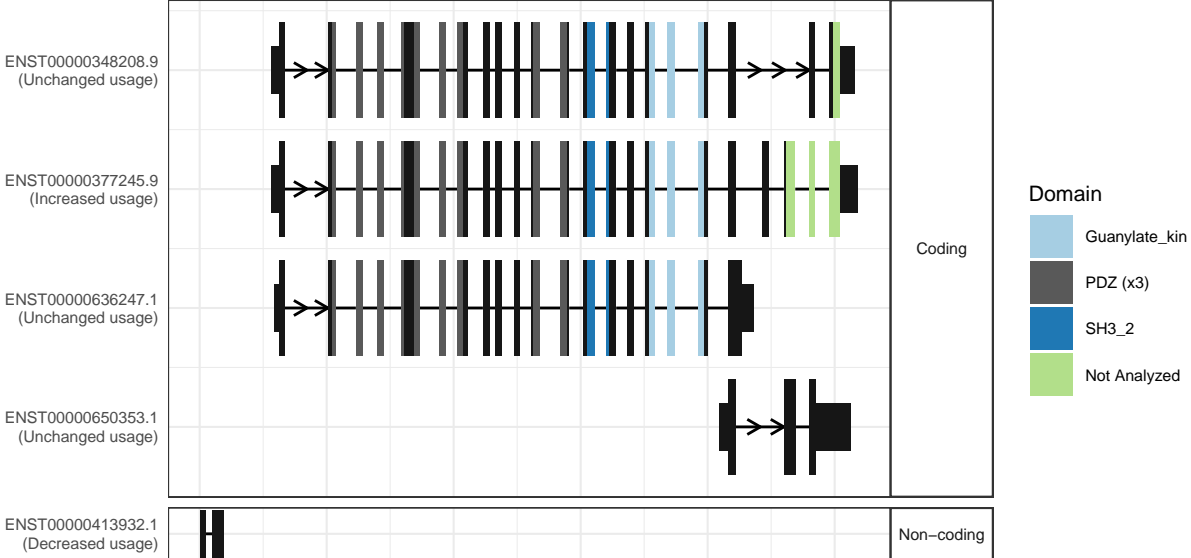
Condition







Domain



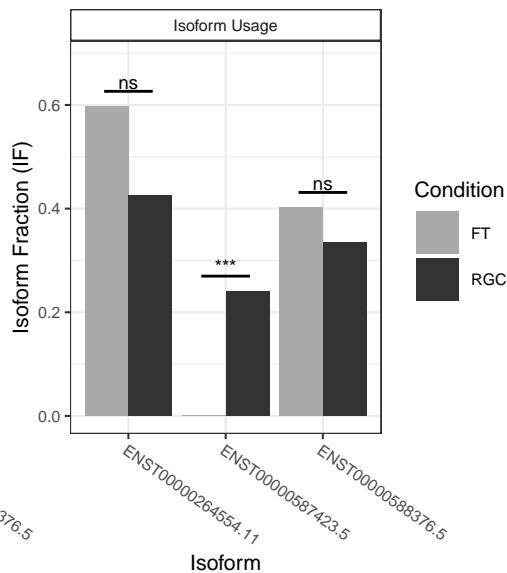
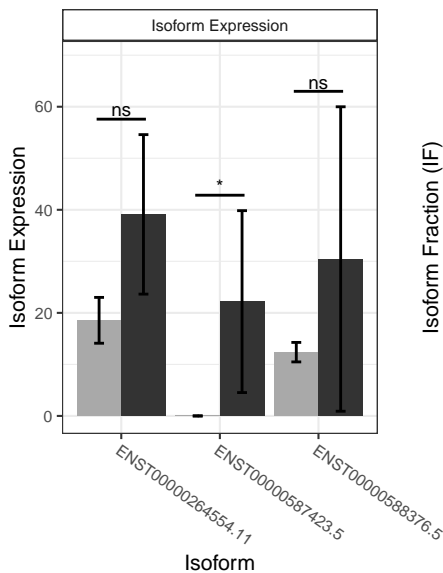
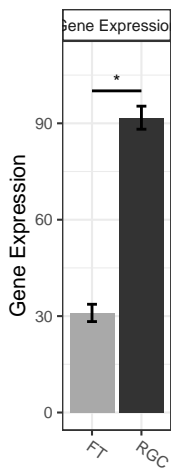
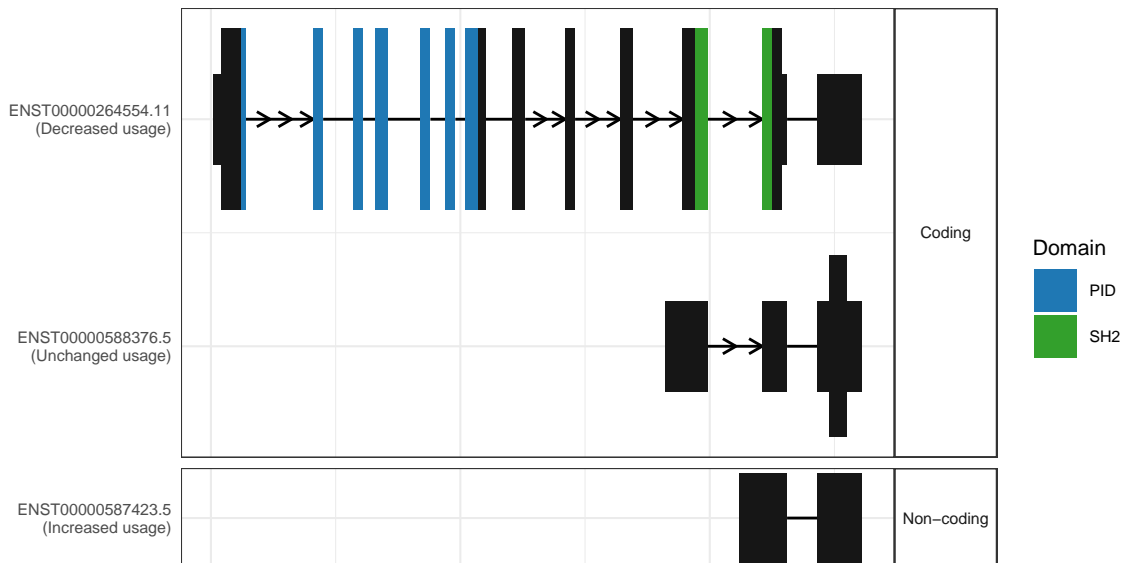
The isoform switch in TJP2 (FT vs RGC)



Domain

	Guanylate_kin
	PDZ (x3)
	SH3_2
	Not Analyzed

The isoform switch in SHC2 (FT vs RGC)



Domain

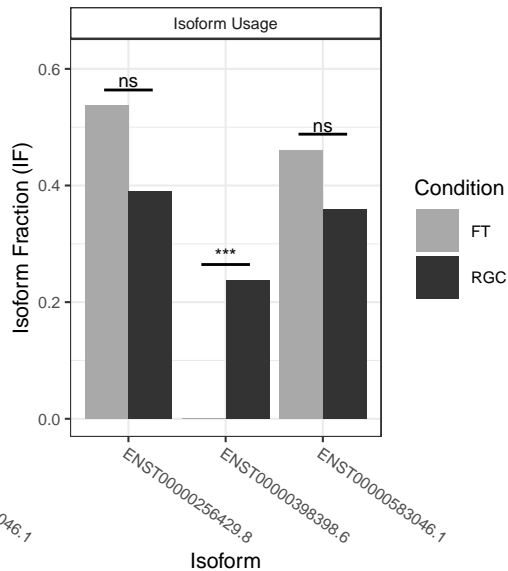
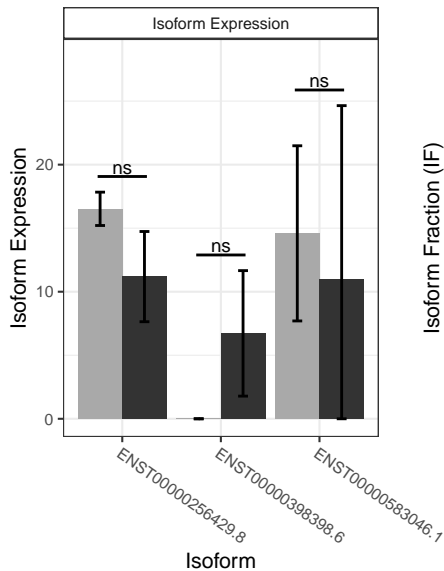
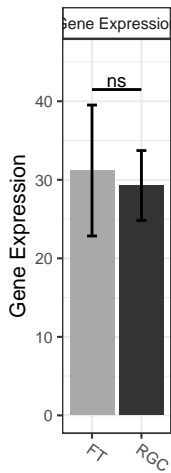
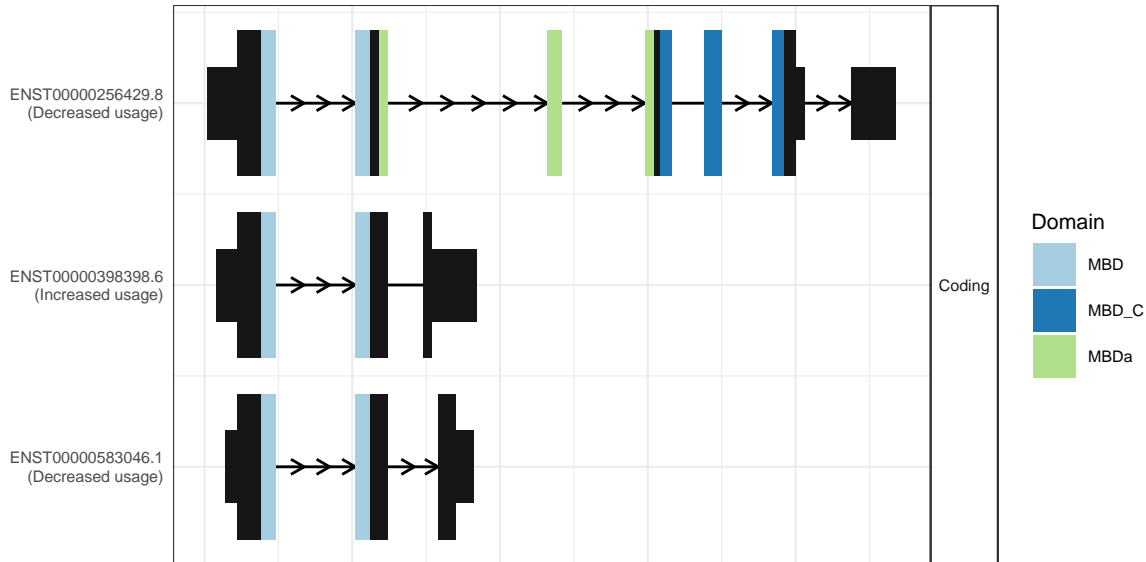


PID



SH2

The isoform switch in MBD2 (FT vs RGC)



Domain



MBD

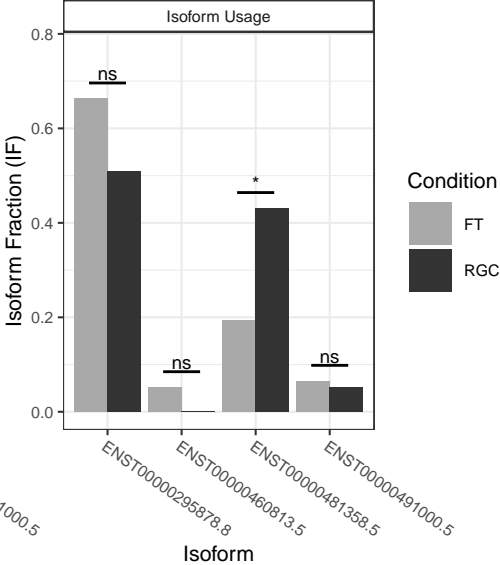
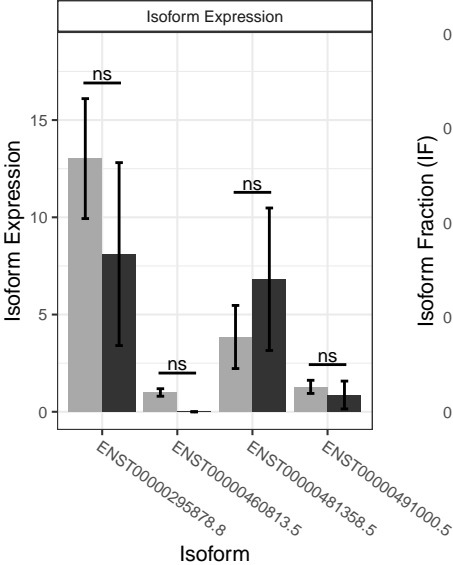
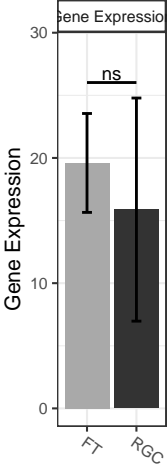
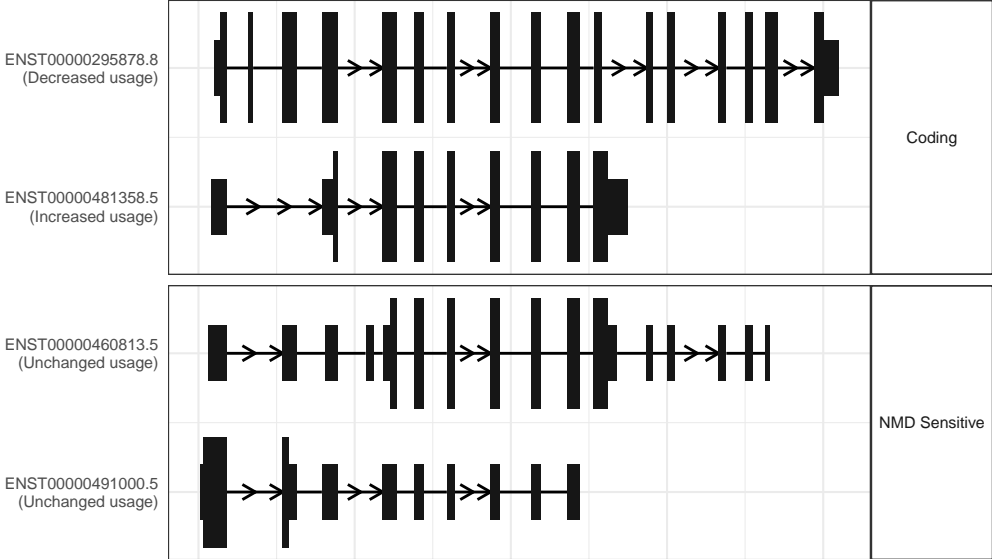


MBD_C

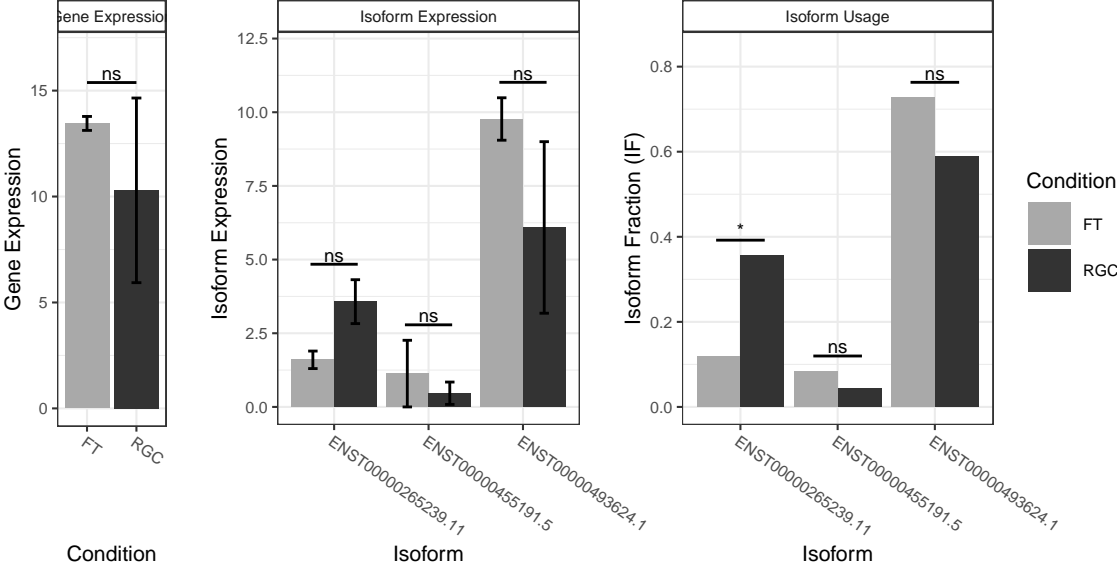
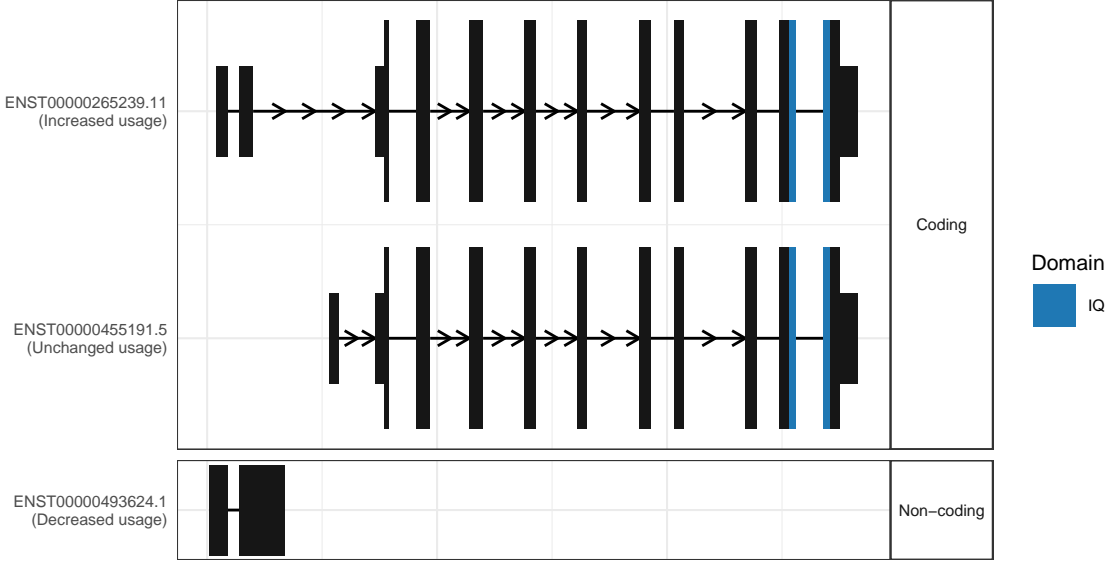


MBDa

The isoform switch in CCDC191 (FT vs RGC)



The isoform switch in IQCG (FT vs RGC)

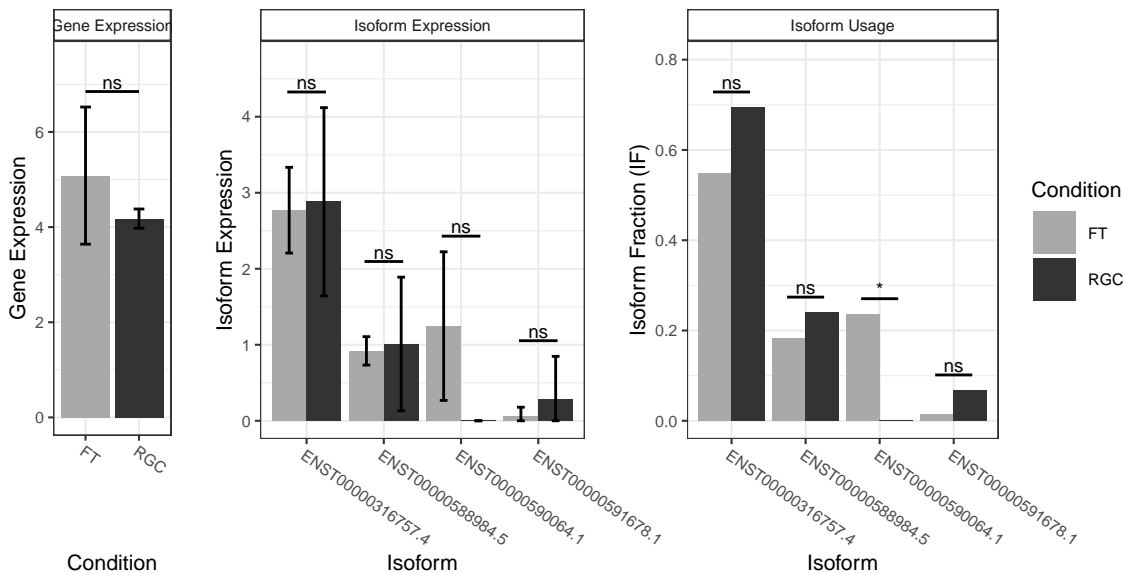
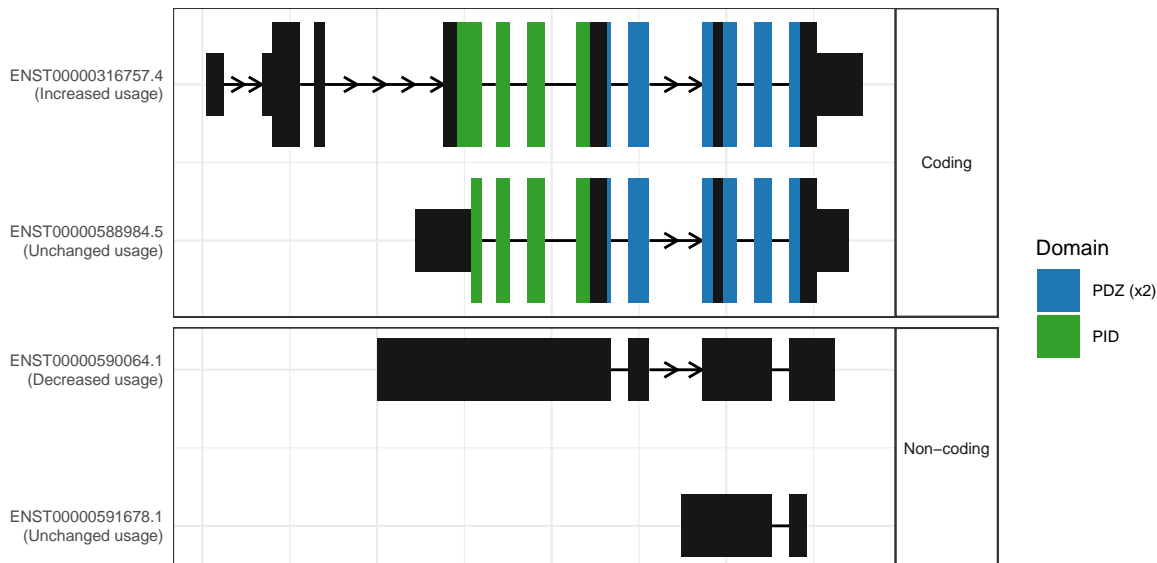


Domain





IQ

The isoform switch in APBA3 (FT vs RGC)

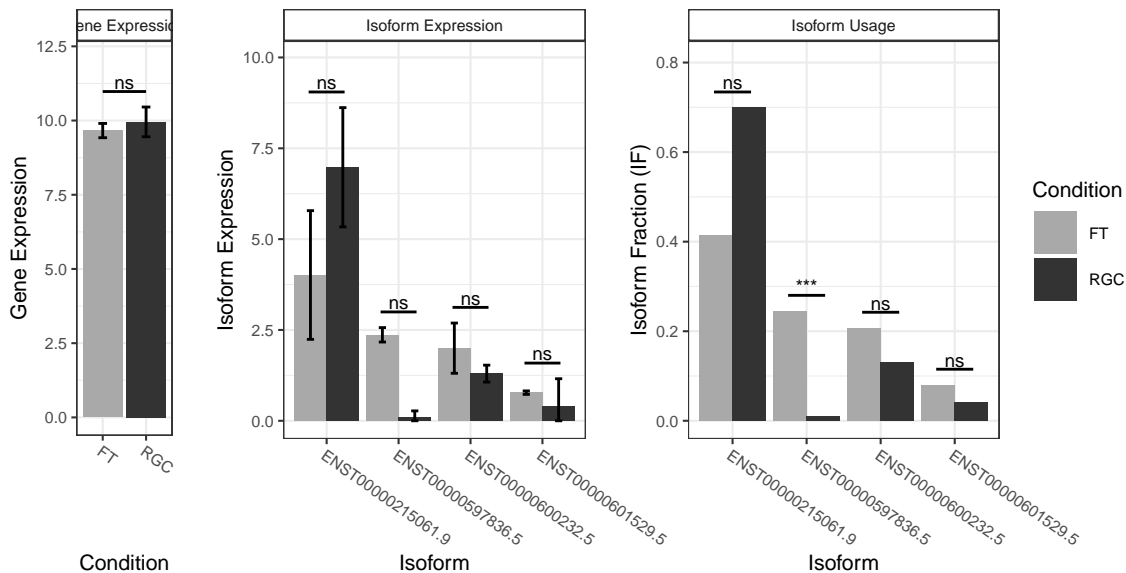
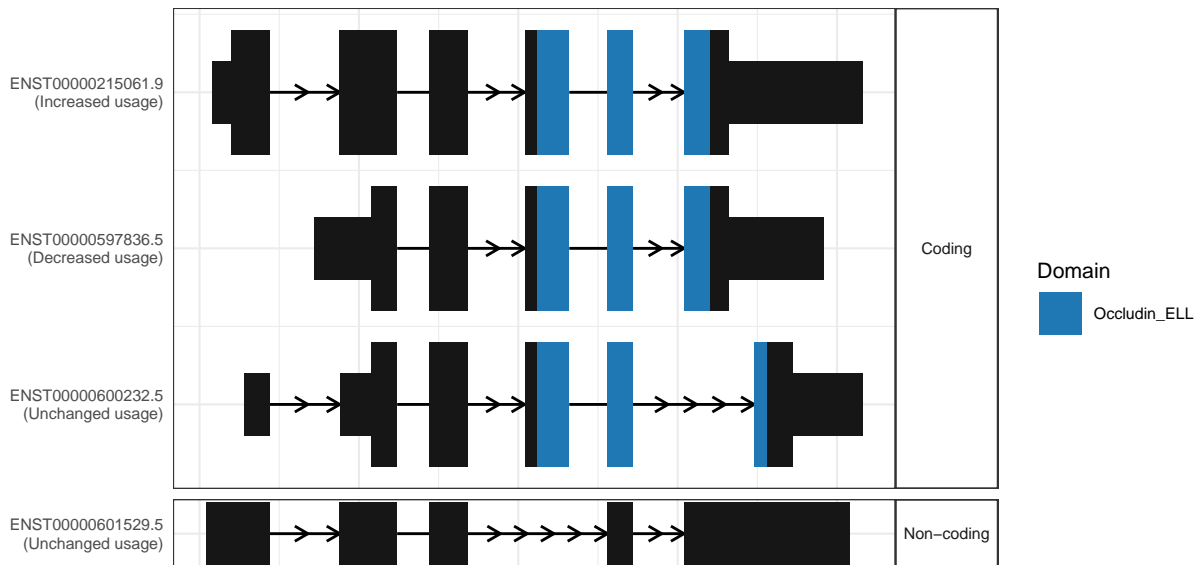


Domain

 PDZ (x2)

 PID

The isoform switch in OCEL1 (FT vs RGC)

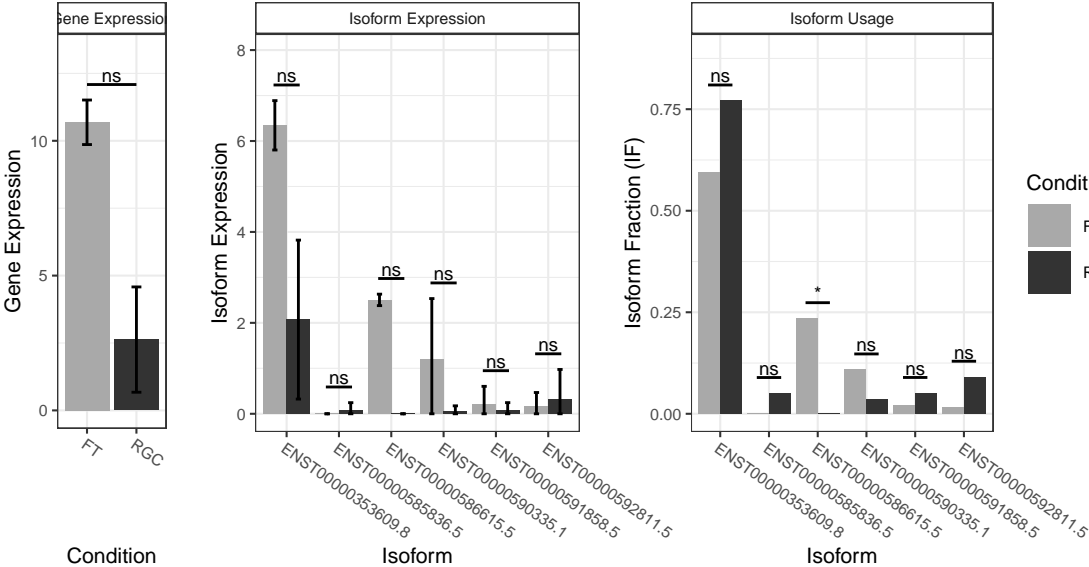
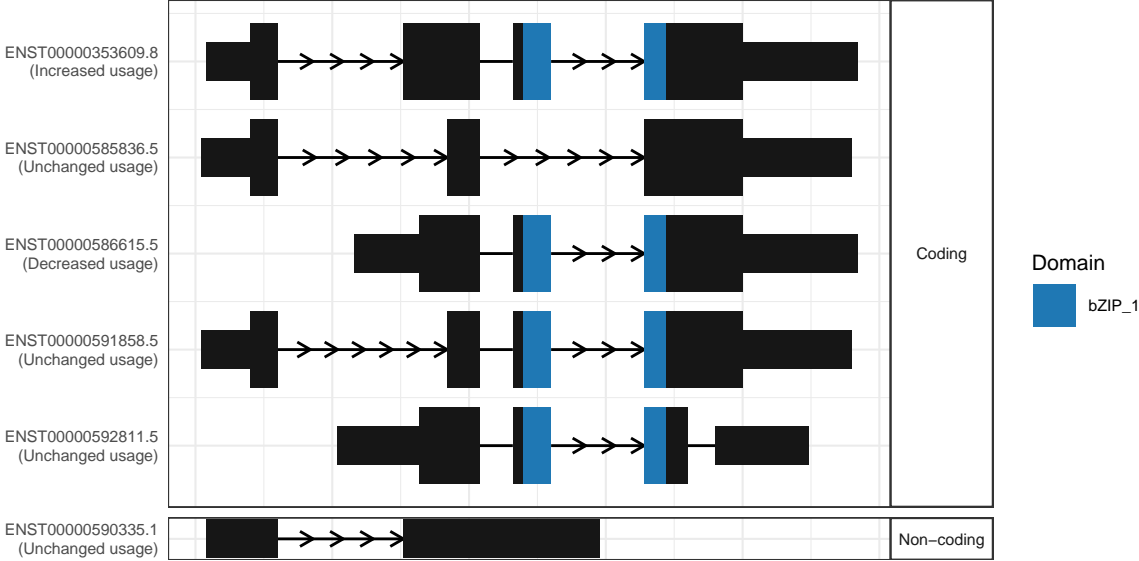


Domain



Occludin_ELL

The isoform switch in FOSB (FT vs RGC)

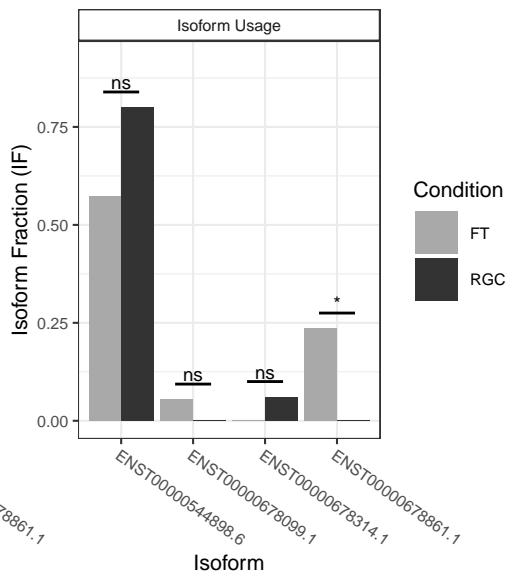
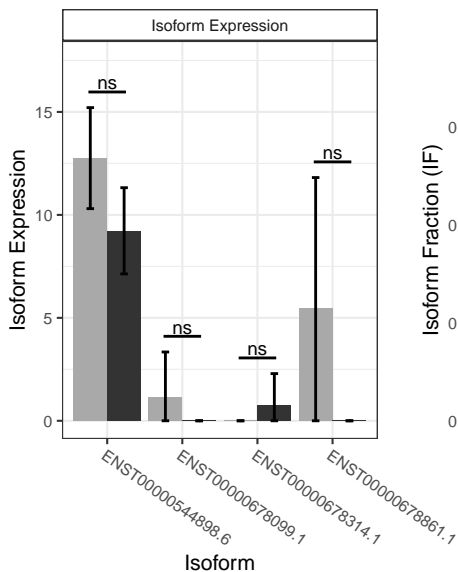
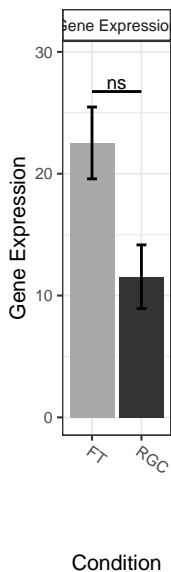
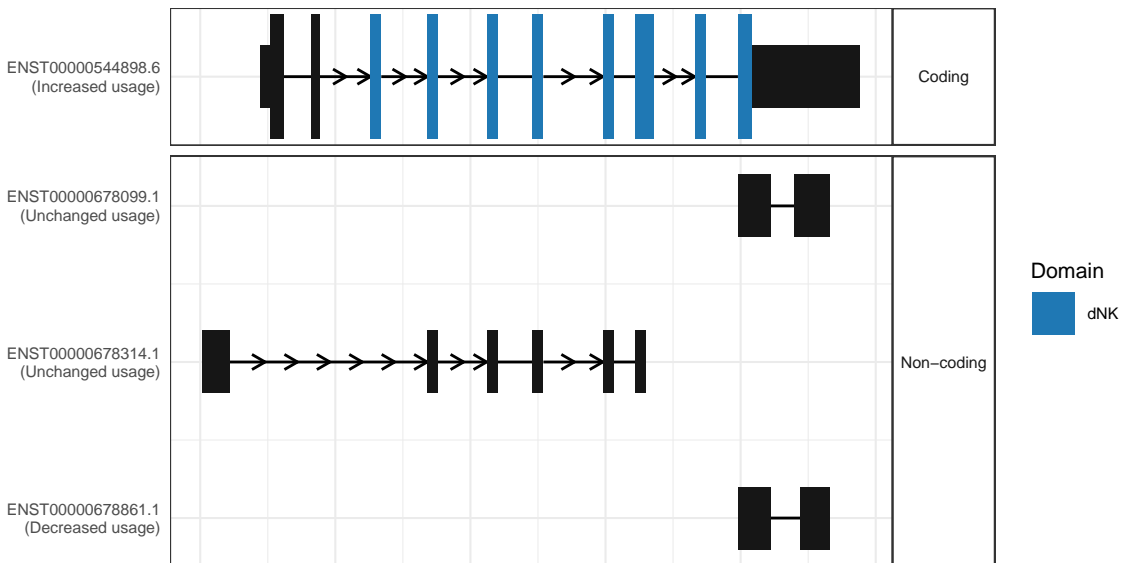


Domain



bZIP_1

The isoform switch in TK2 (FT vs RGC)

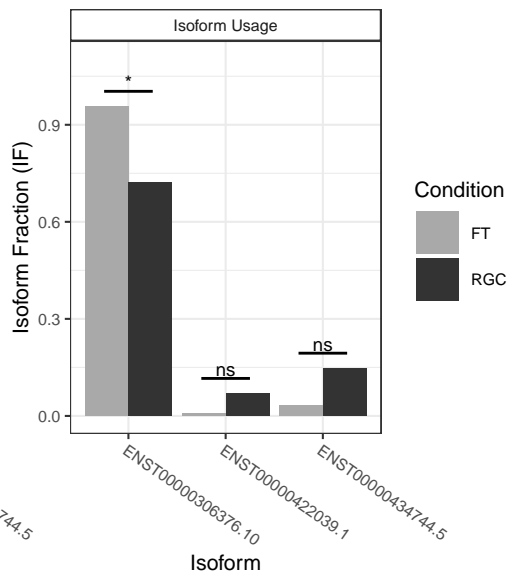
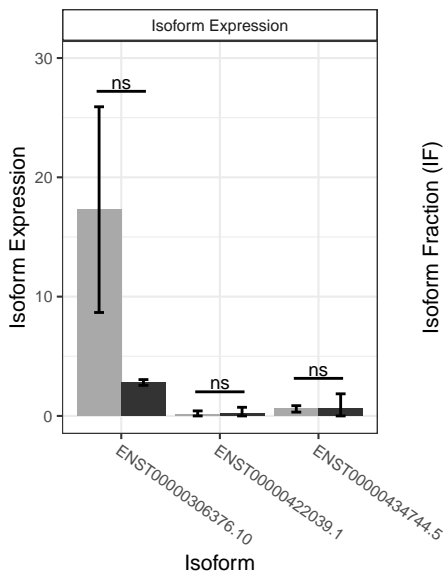
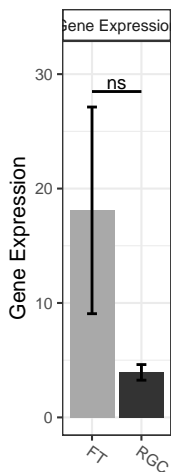
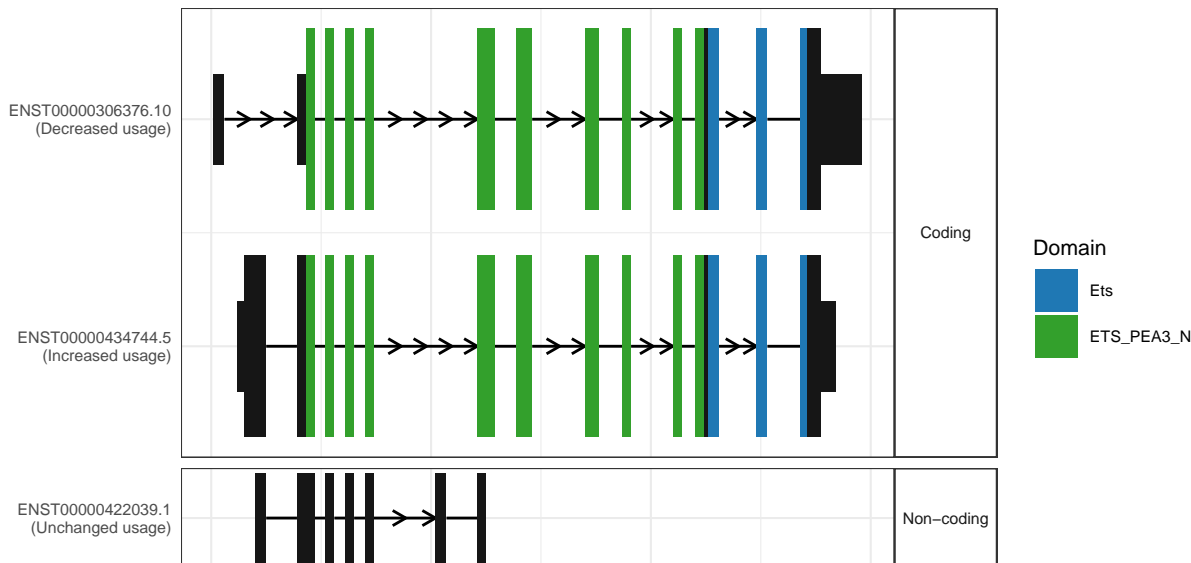


Domain



dNK

The isoform switch in ETV5 (FT vs RGC)



Domain

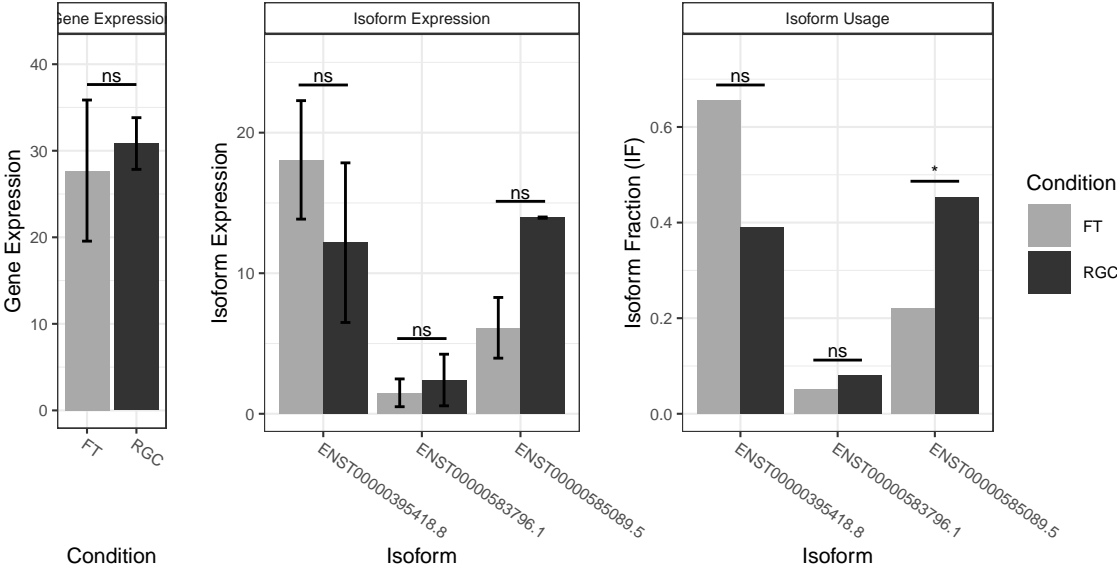
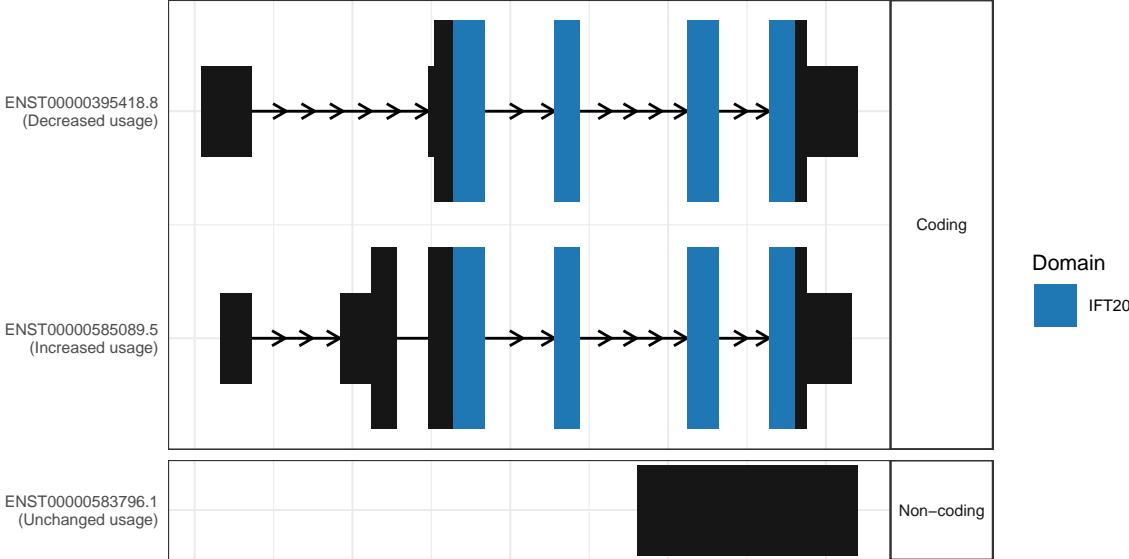


Ets



ETS_PEA3_N

The isoform switch in IFT20 (FT vs RGC)

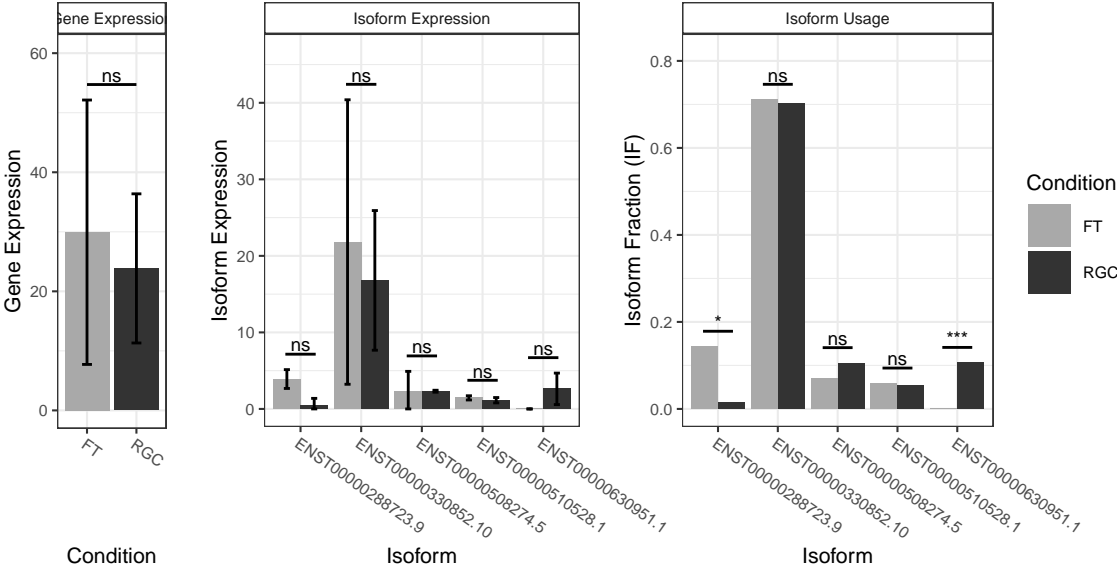
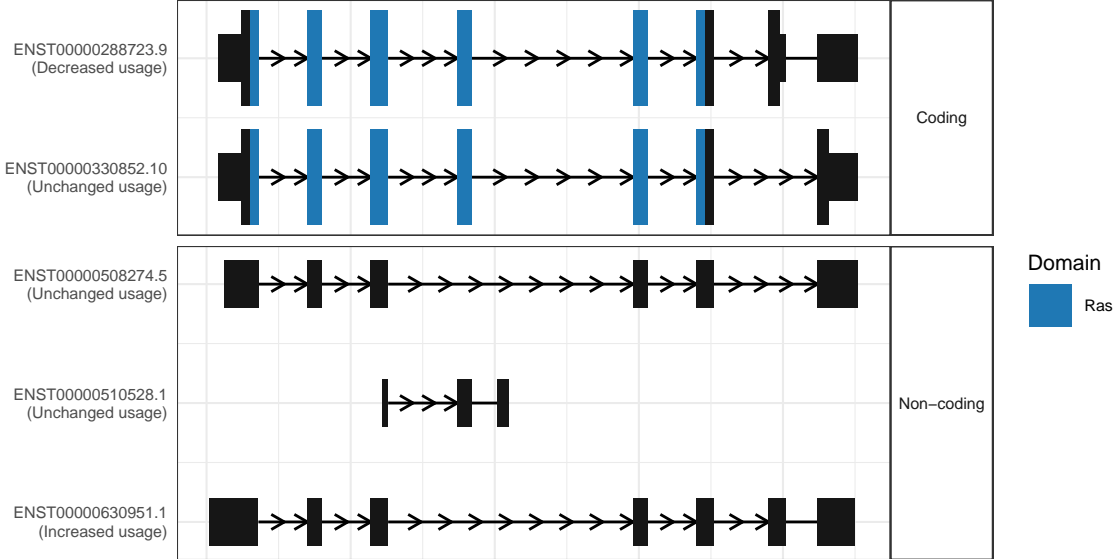


Domain



IFT20

The isoform switch in RAB28 (FT vs RGC)

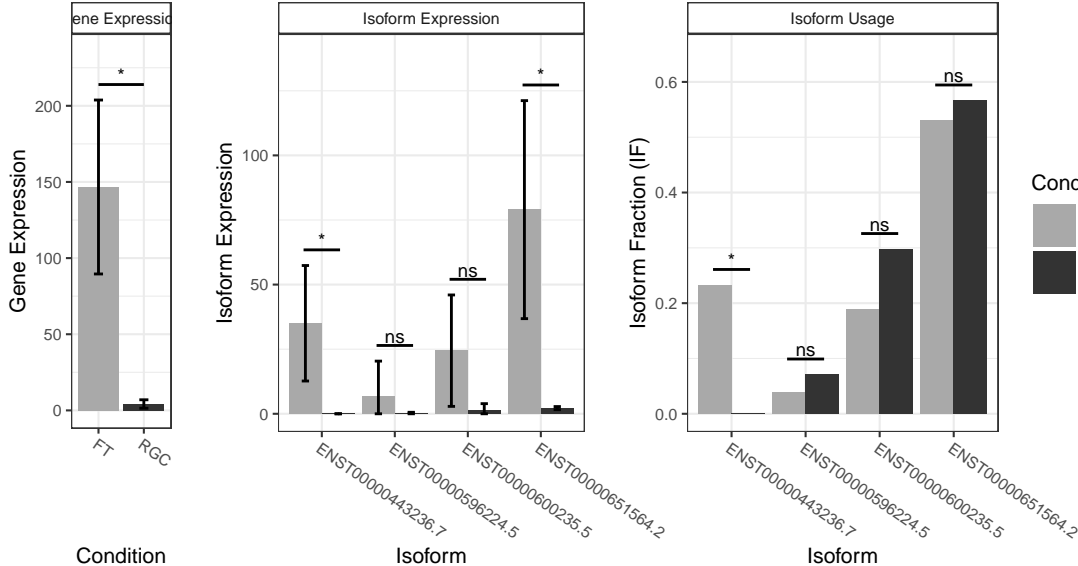
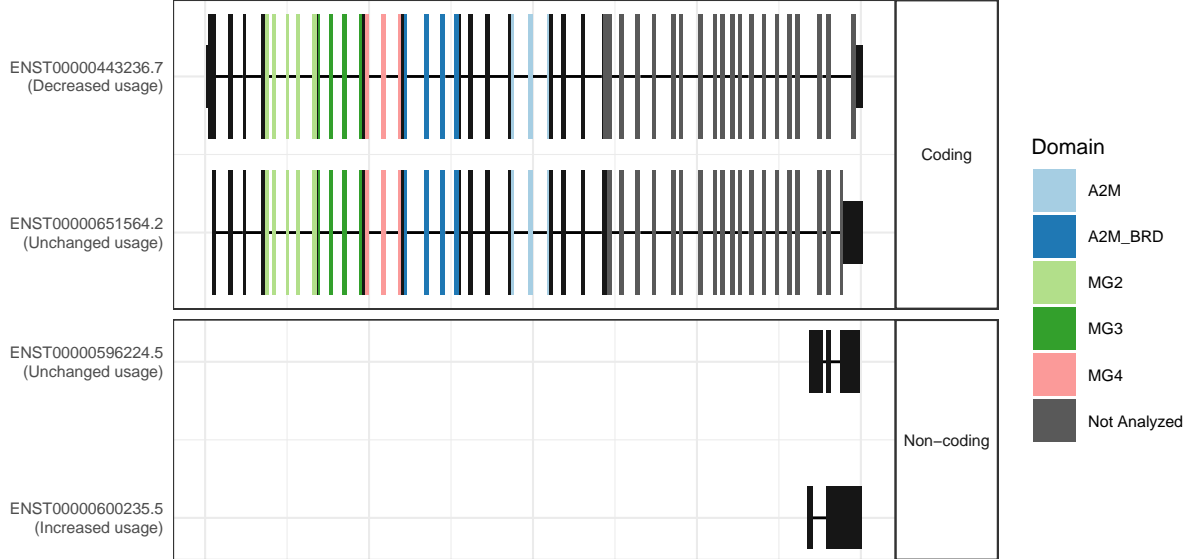


Domain

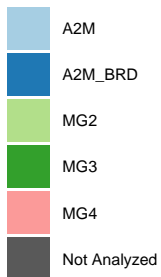


Ras

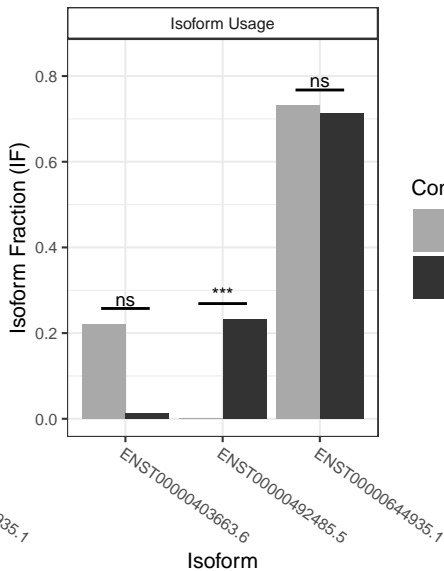
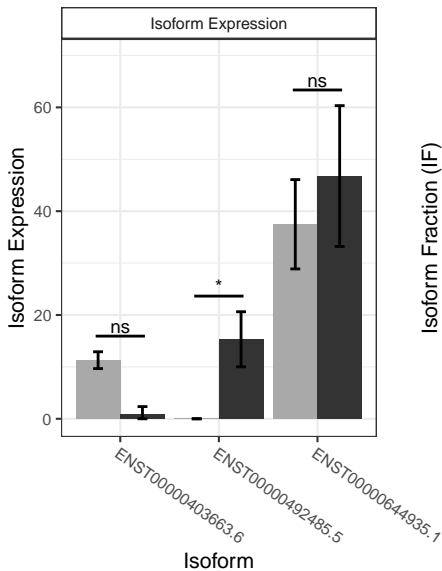
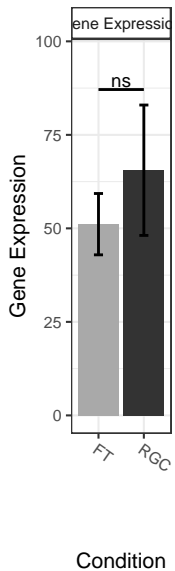
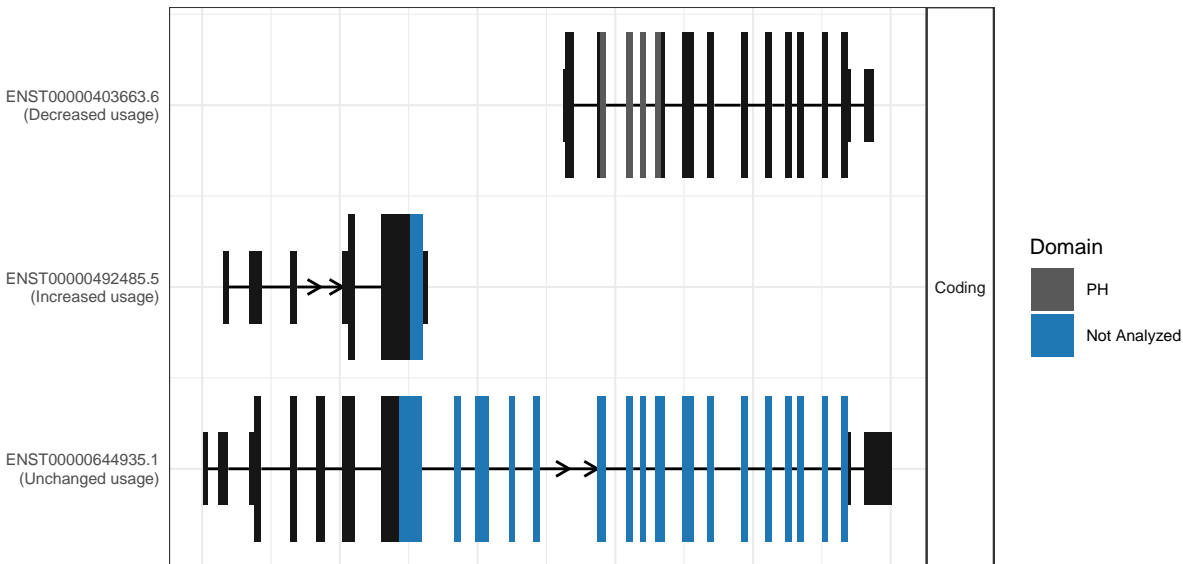
The isoform switch in CPAMD8 (FT vs RGC)



Domain



The isoform switch in TRIOBP (FT vs RGC)



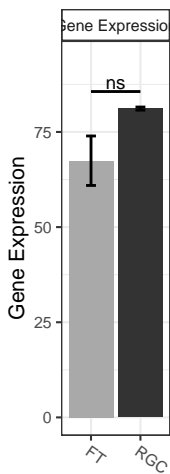
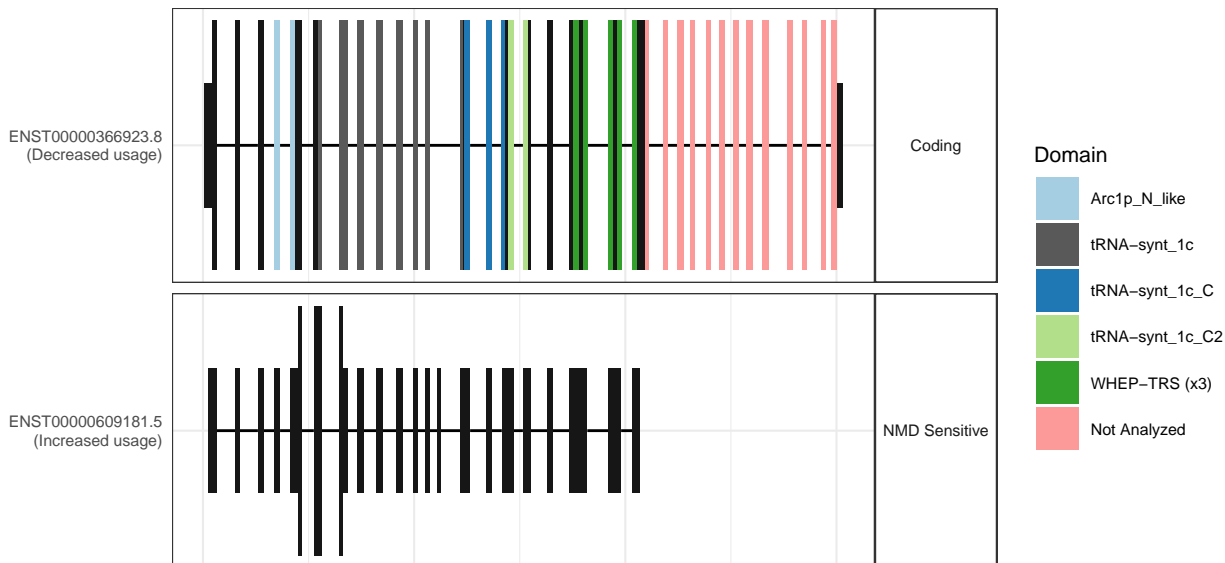
Domain



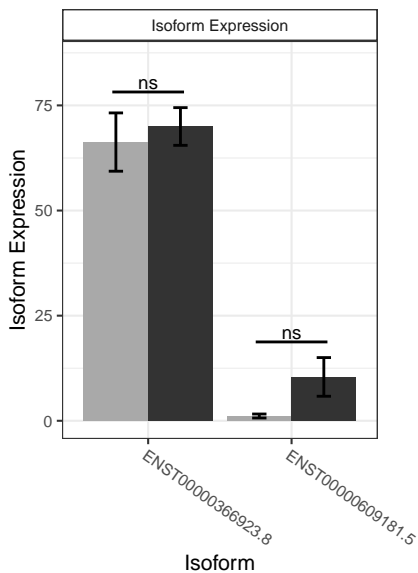
PH

Not Analyzed

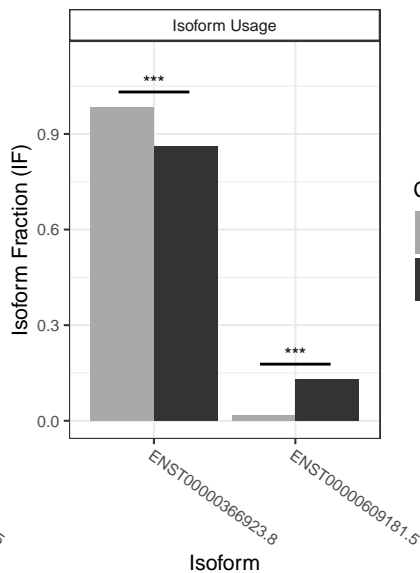
The isoform switch in EPRS1 (FT vs RGC)



Condition



Isoform









Isoform

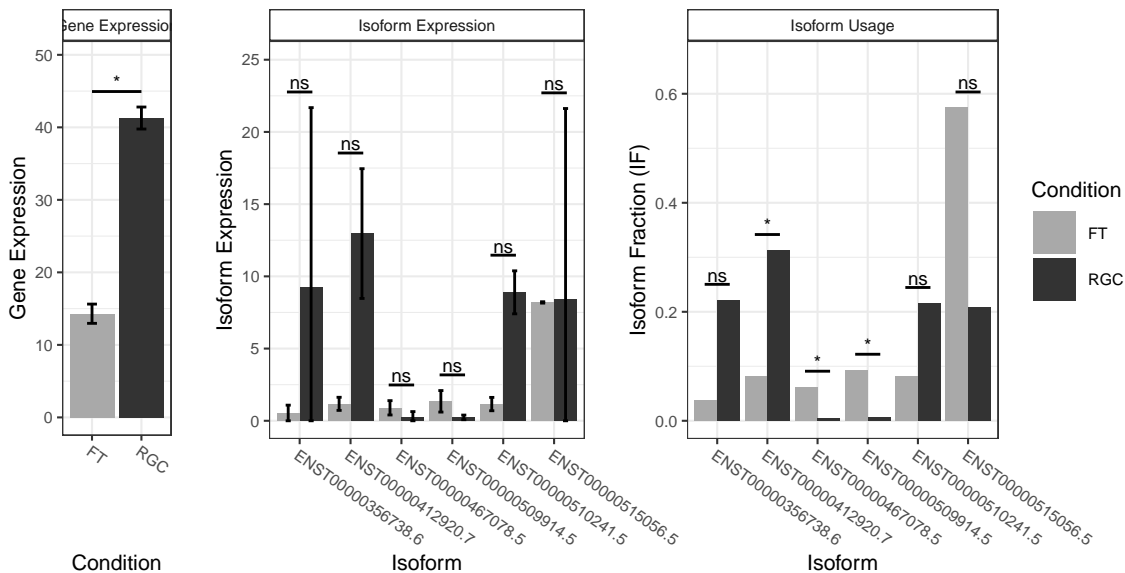
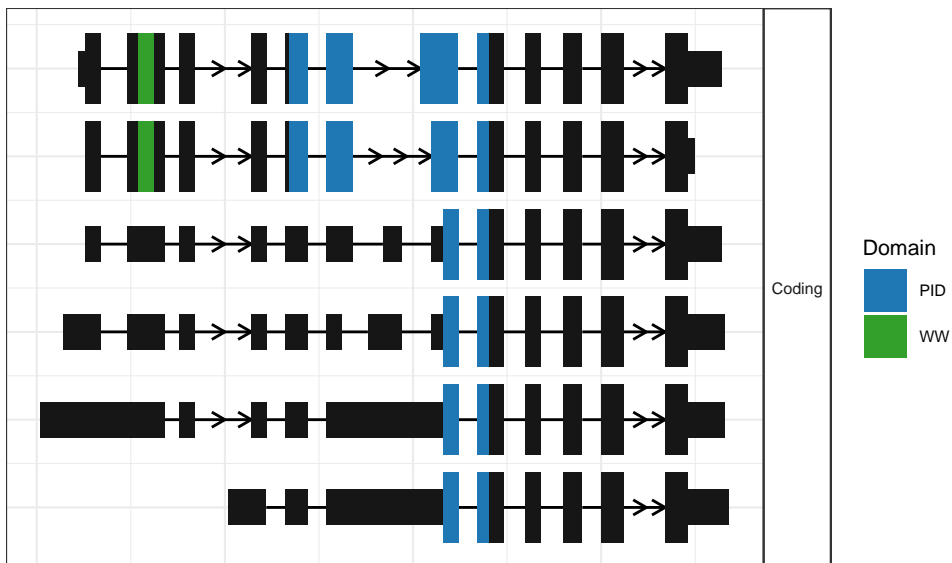
Condition



Domain

	Arc1p_N_like
	tRNA-synt_1c
	tRNA-synt_1c_C
	tRNA-synt_1c_C2
	WHEP-TRS (x3)
	Not Analyzed

The isoform switch in APBB3 (FT vs RGC)



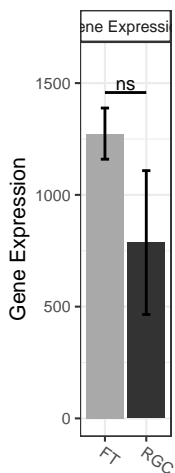
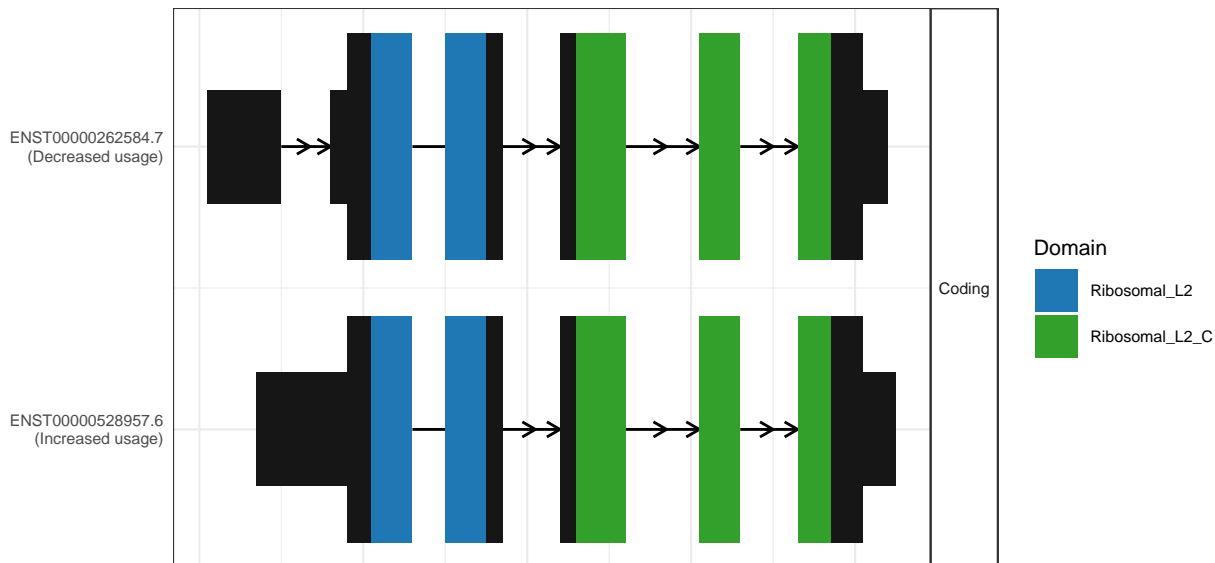
Domain



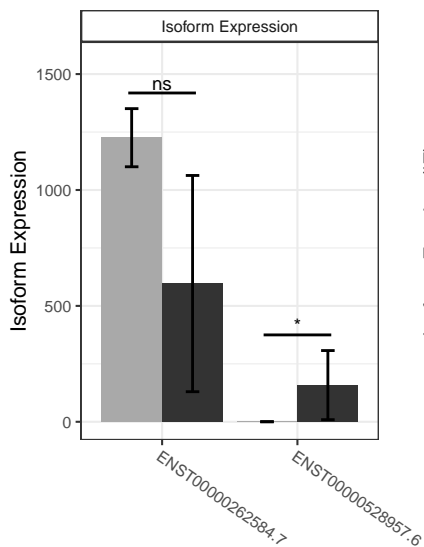
PID

WW

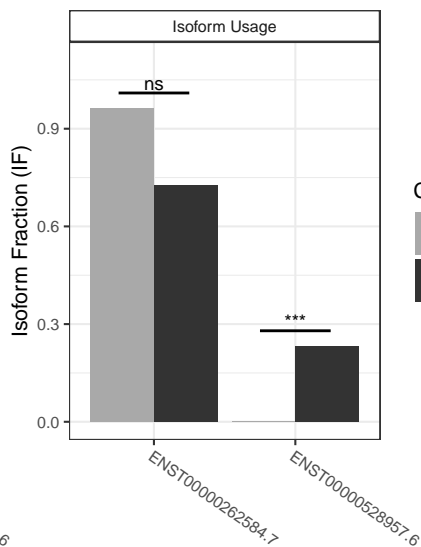
The isoform switch in RPL8 (FT vs RGC)



Condition



Isoform



Isoform

Condition



Domain

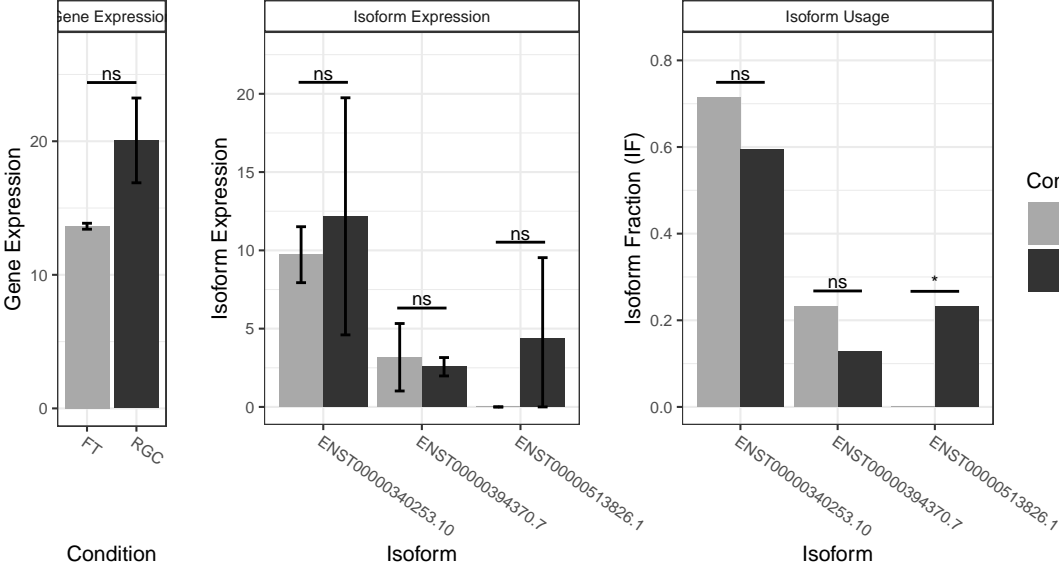
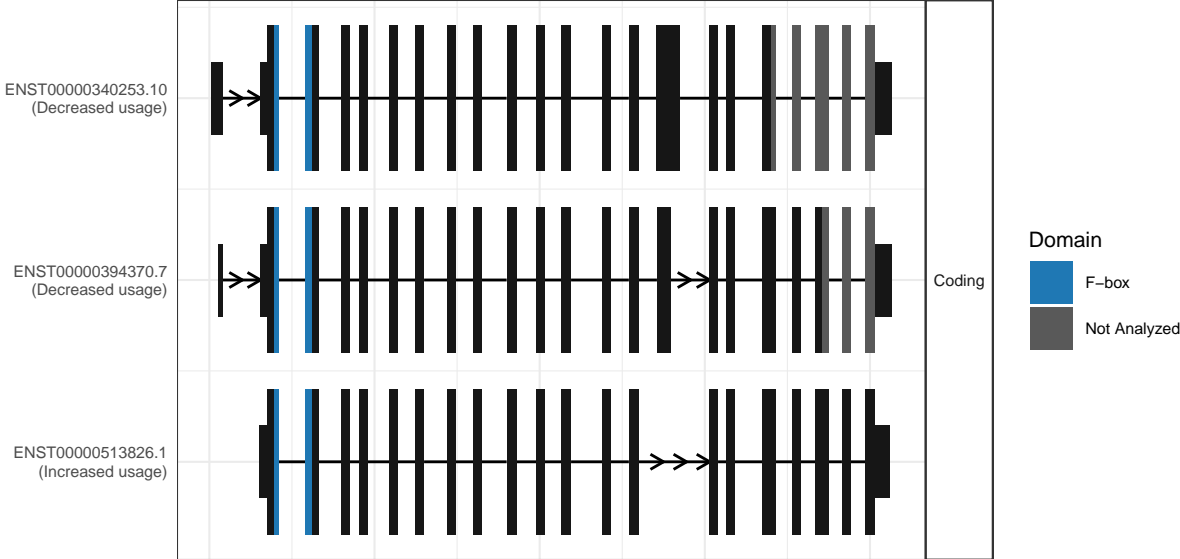


Ribosomal_L2



Ribosomal_L2_C

The isoform switch in FBXO38 (FT vs RGC)



Domain

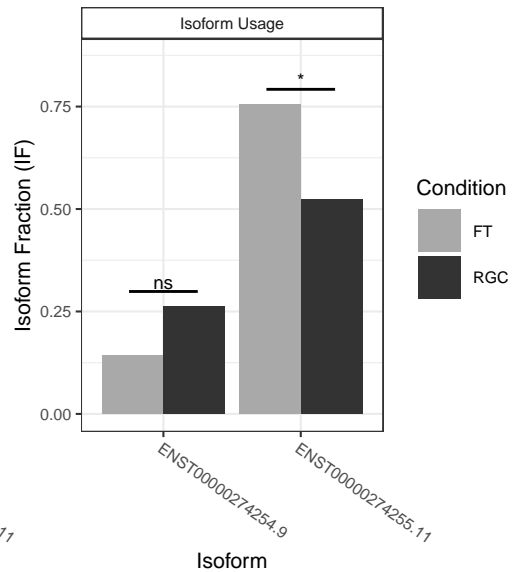
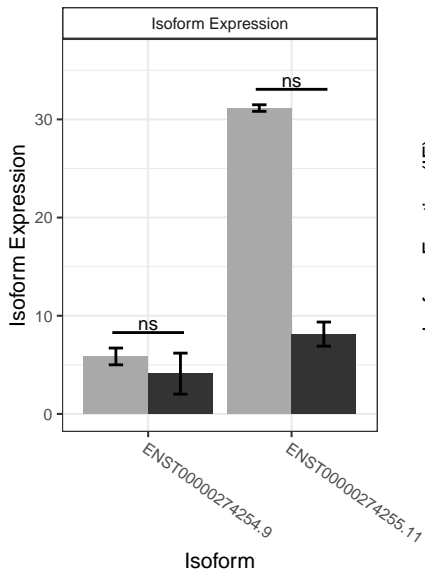
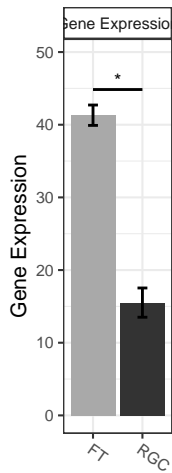
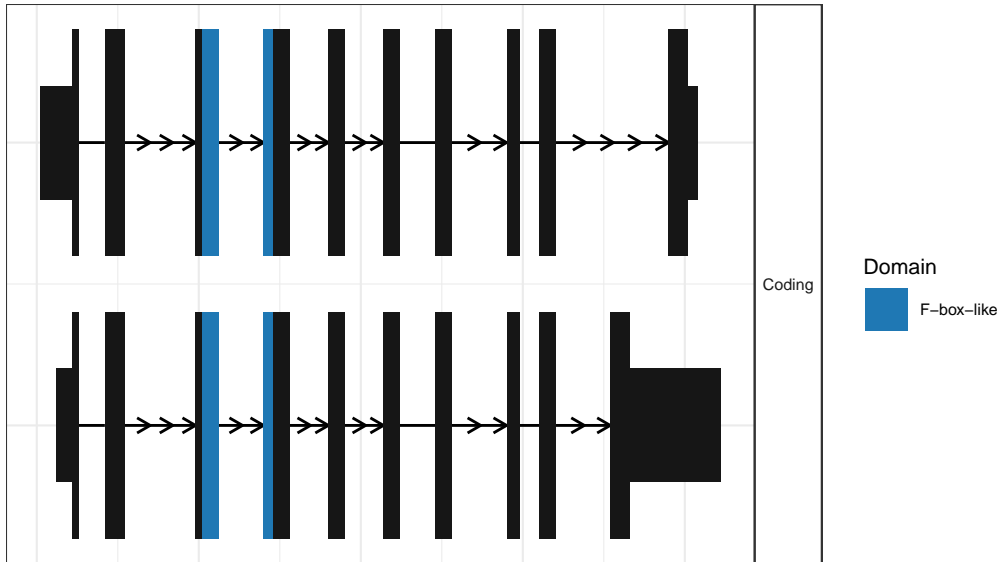


F-box



Not Analyzed

The isoform switch in SKP2 (FT vs RGC)

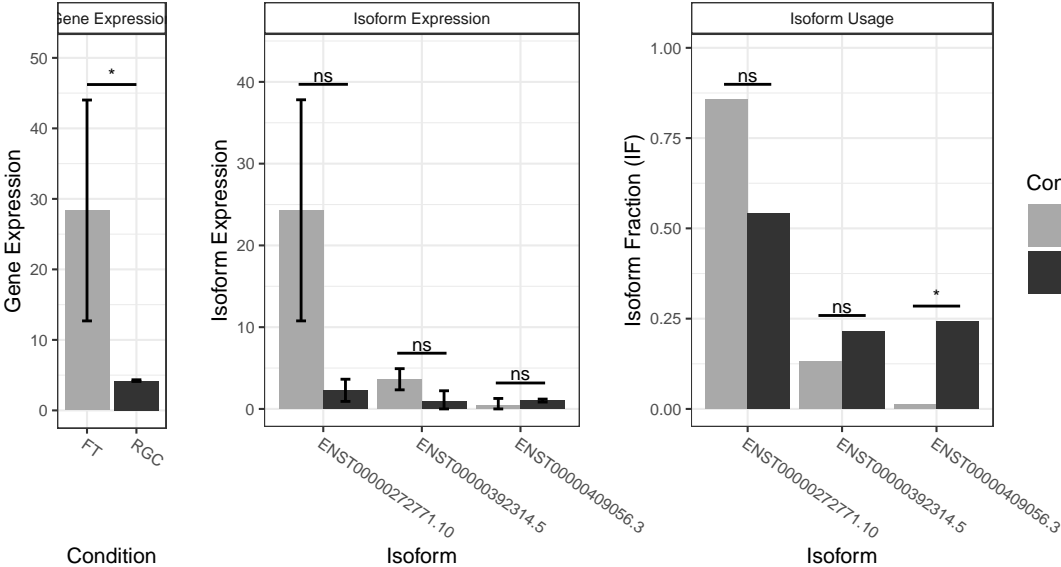
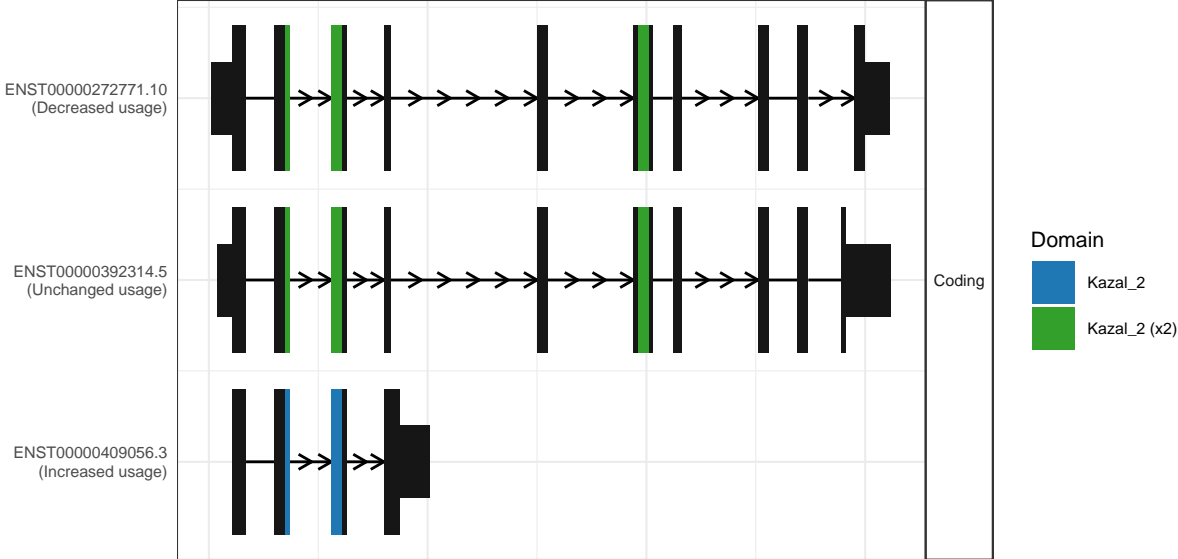


Domain



F-box-like

The isoform switch in TMEFF2 (FT vs RGC)



Domain

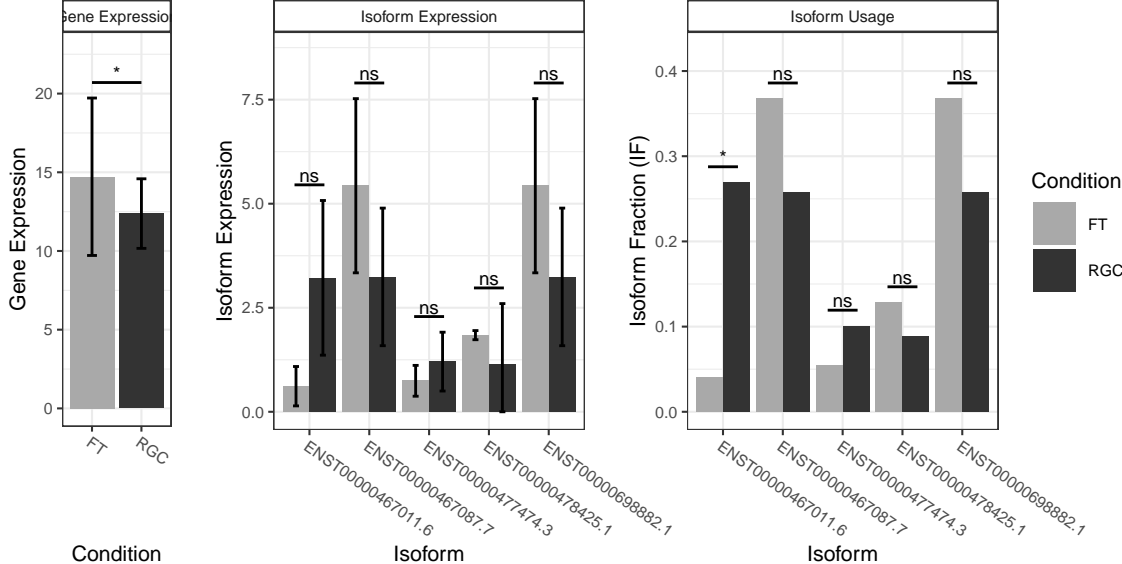
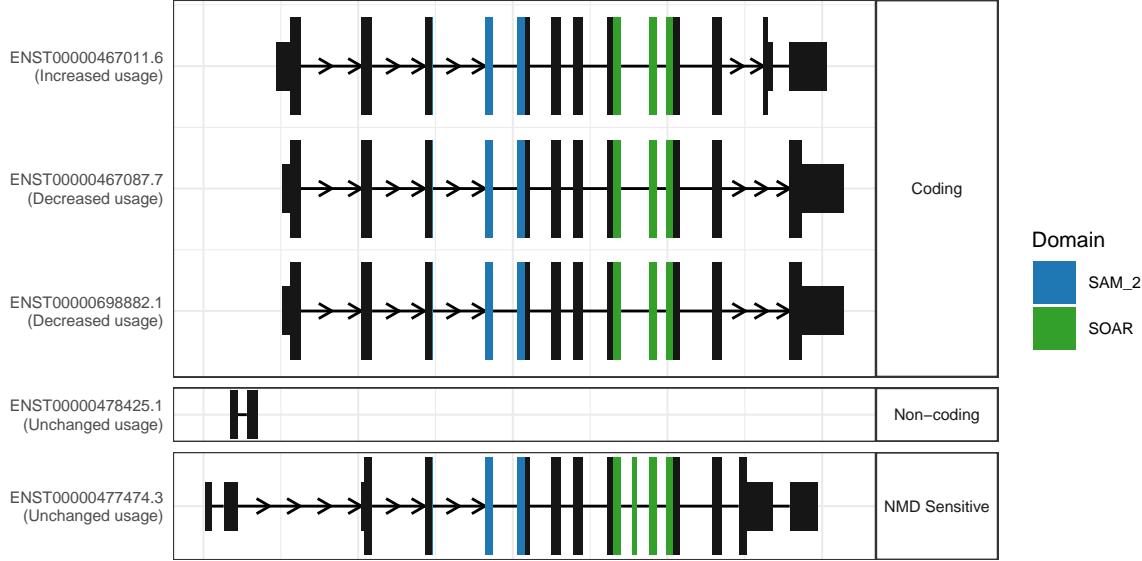


Kazal_2



Kazal_2 (x2)

The isoform switch in STIM2 (FT vs RGC)



Domain

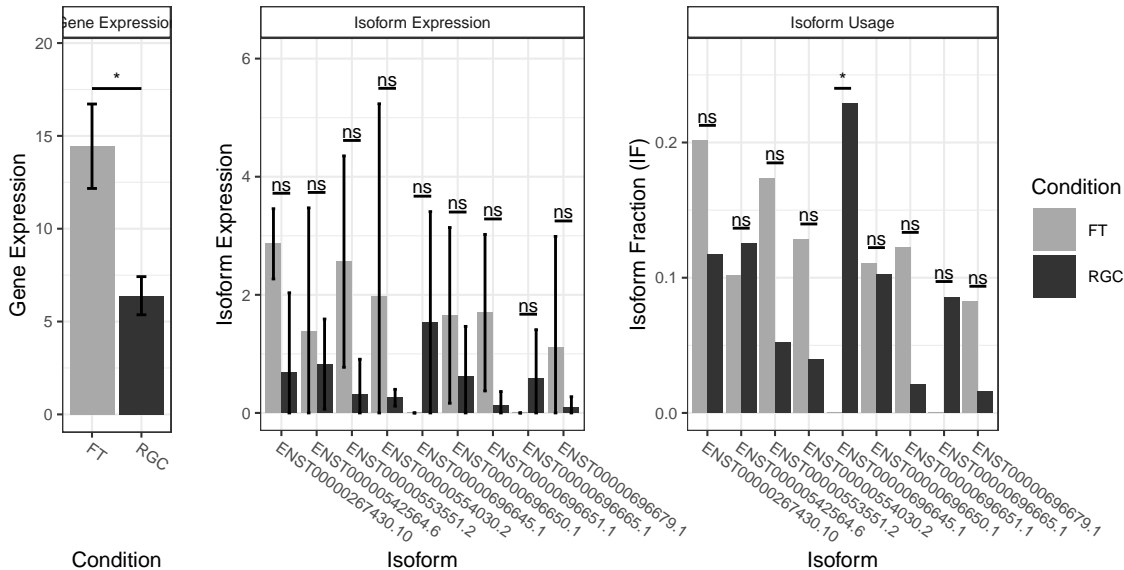
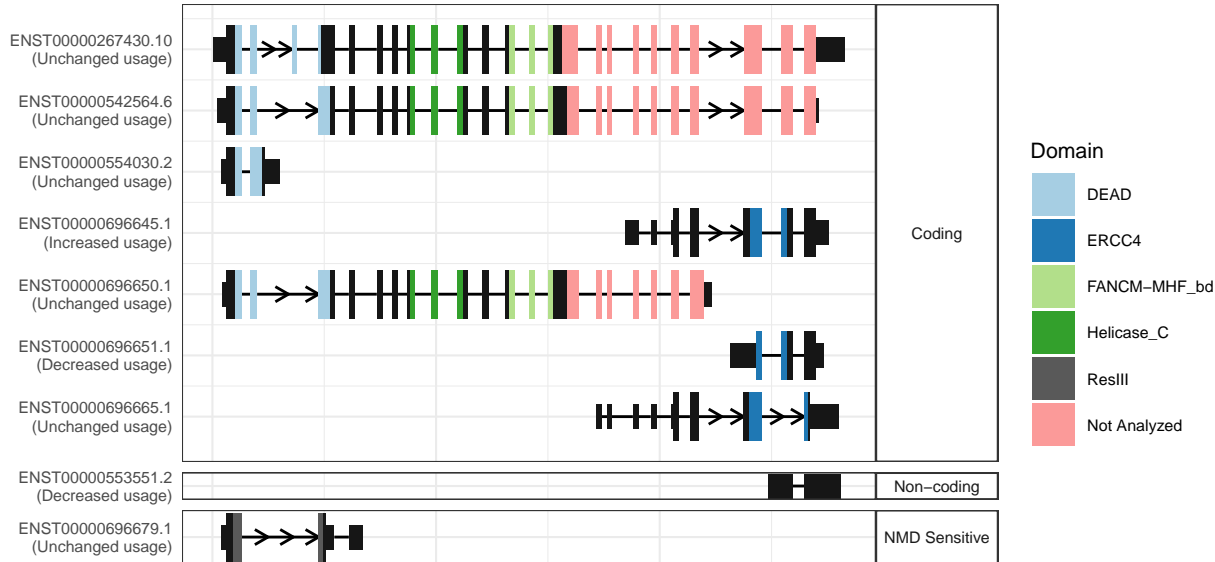


SAM_2









SOAR

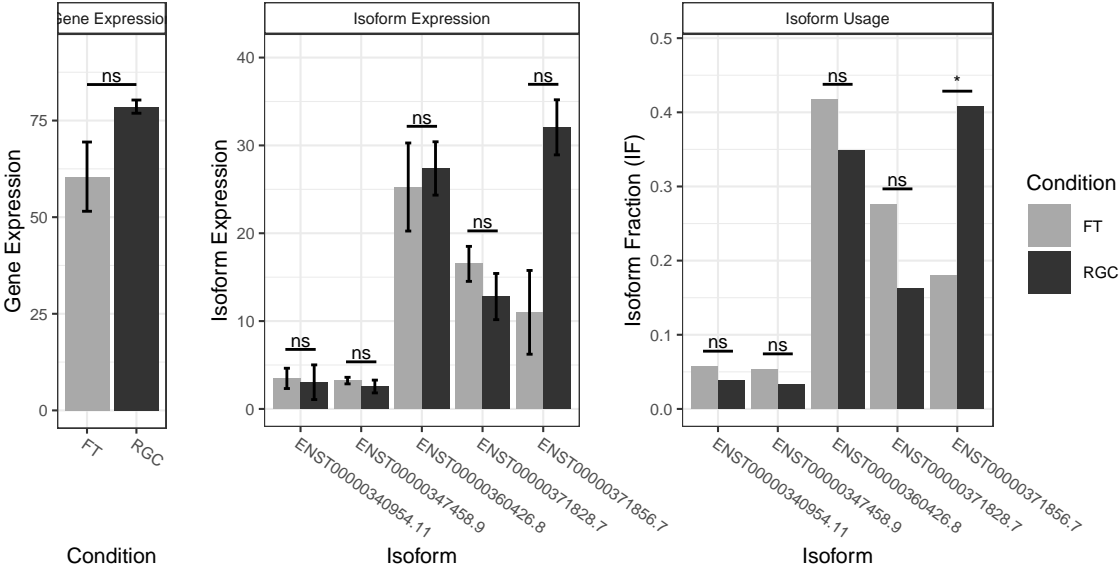
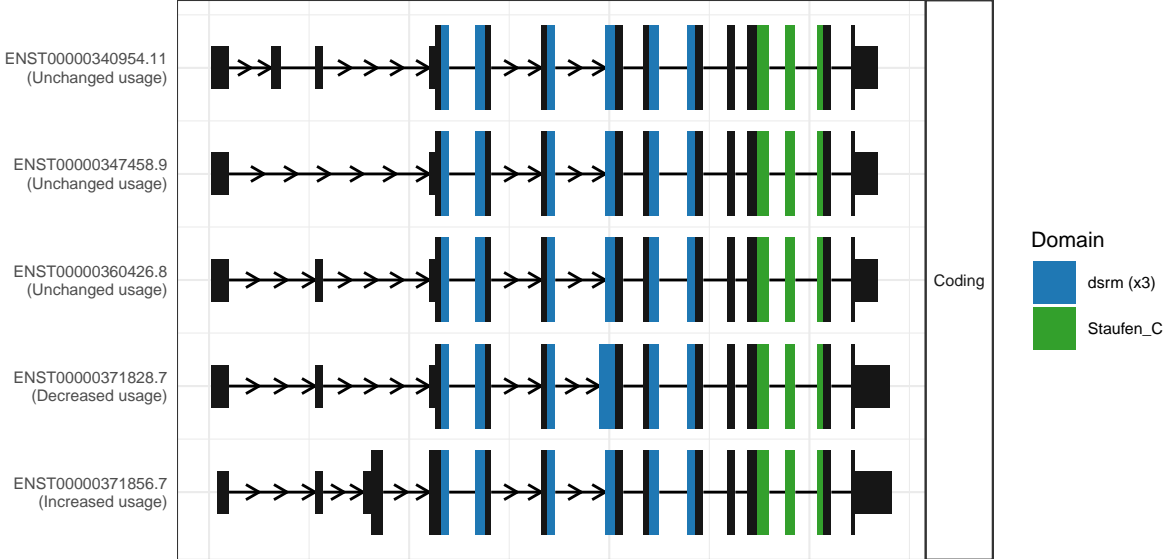
The isoform switch in FANCM (FT vs RGC)



Domain

	DEAD
	ERCC4
	FANCM-MHF_bd
	Helicase_C
	ResIII
	Not Analyzed

The isoform switch in STAU1 (FT vs RGC)



Domain

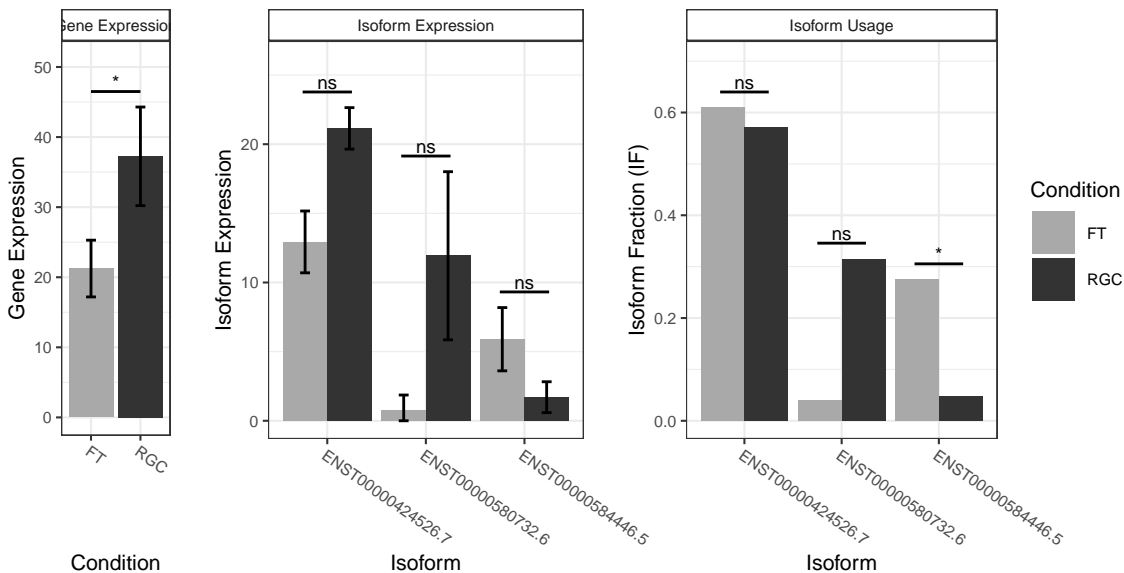
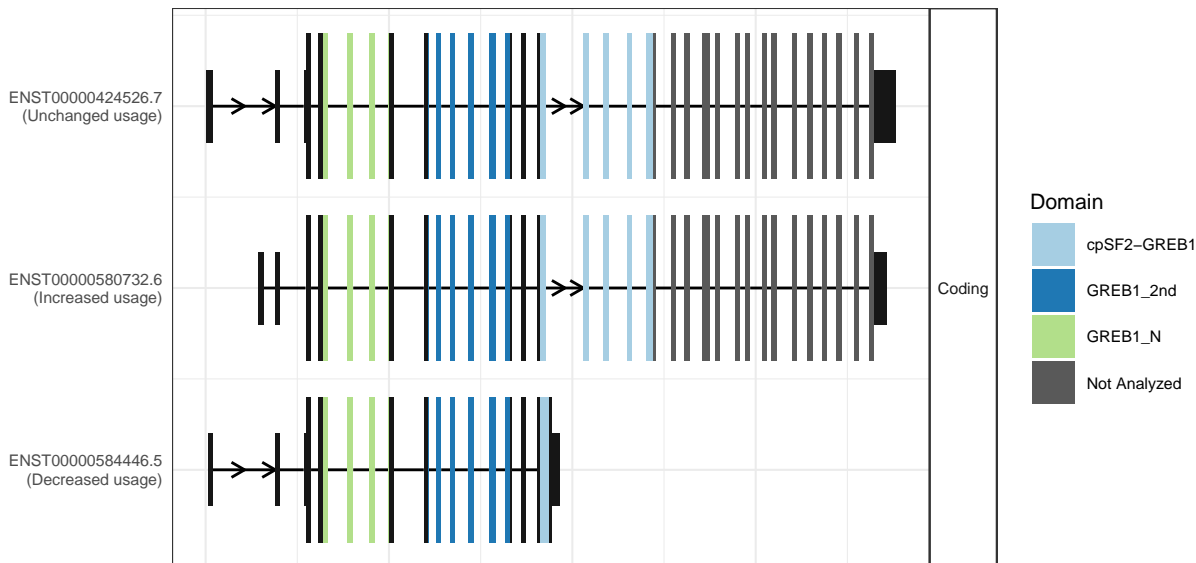


dsrm (x3)

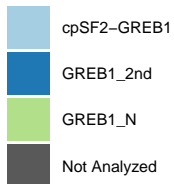


Staufen_C

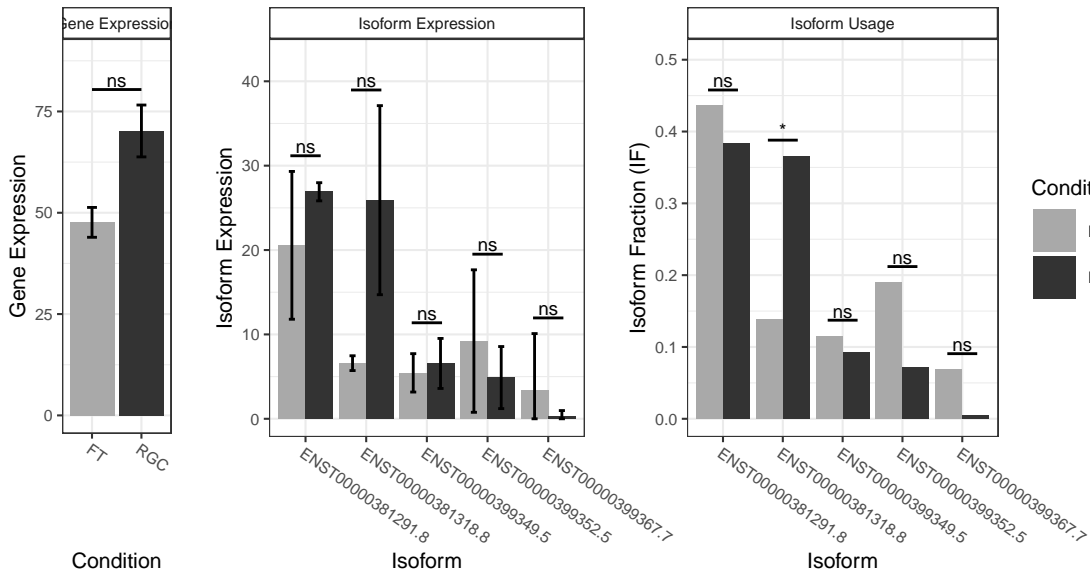
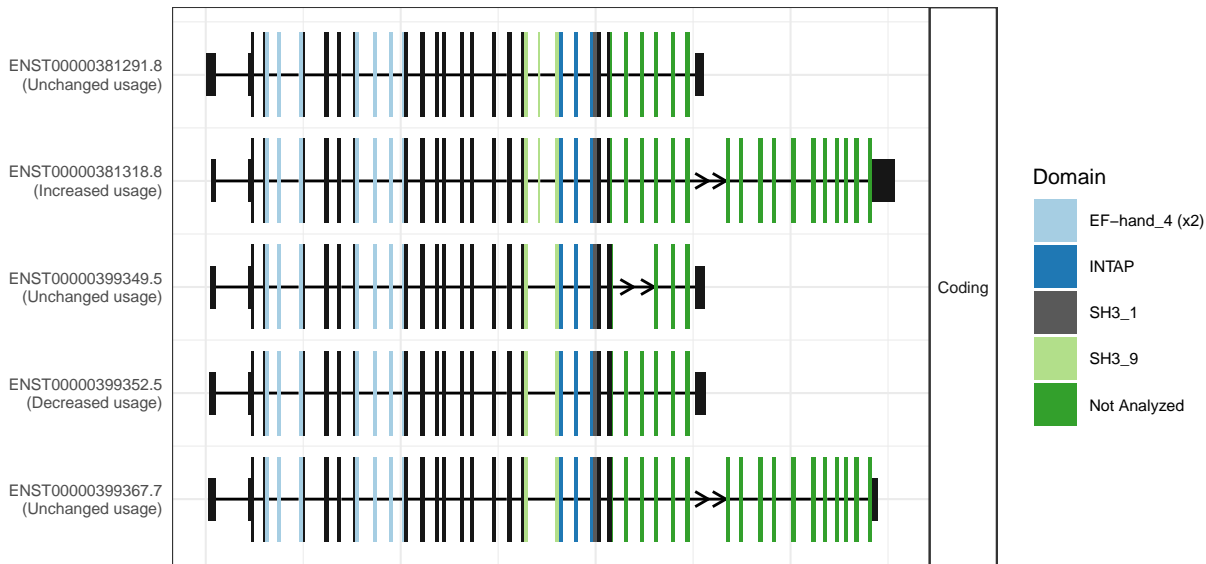
The isoform switch in GREB1L (FT vs RGC)



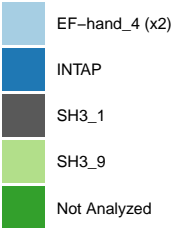
Domain



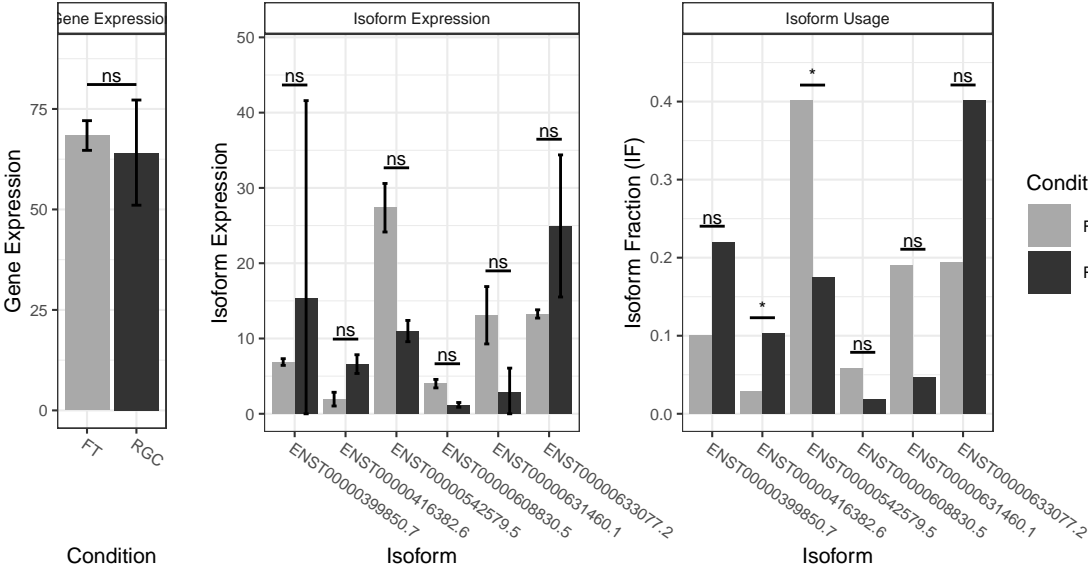
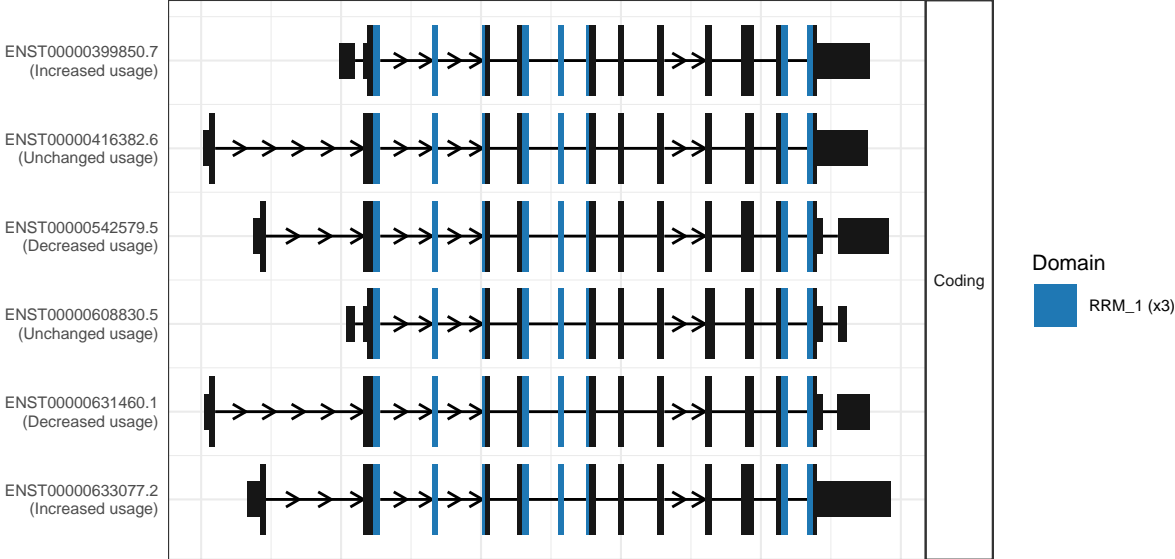
The isoform switch in ITSN1 (FT vs RGC)




Domain



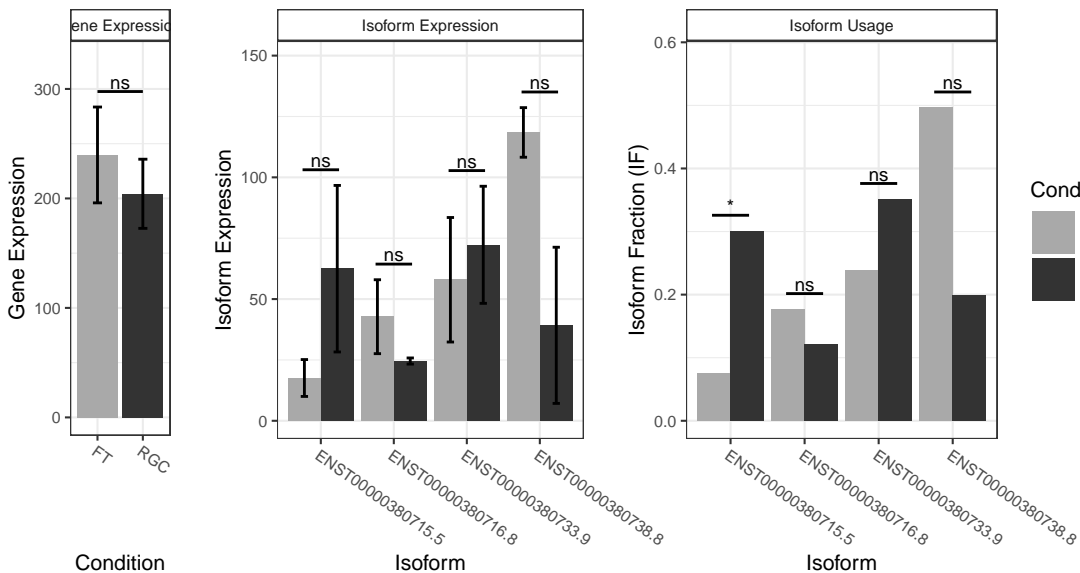
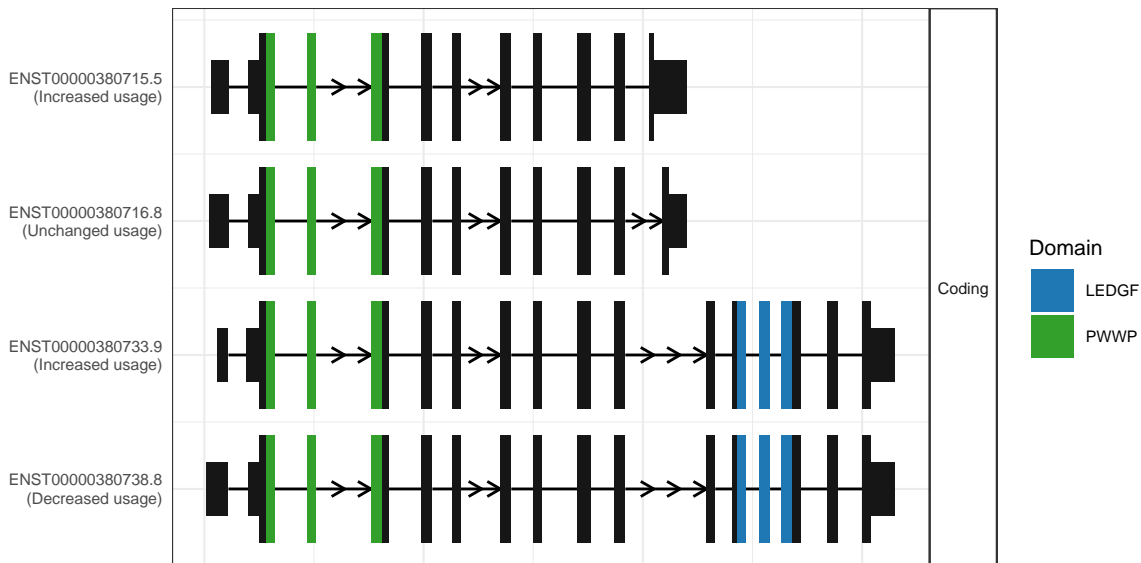
The isoform switch in CELF2 (FT vs RGC)



Domain

 RRM_1 (x3)

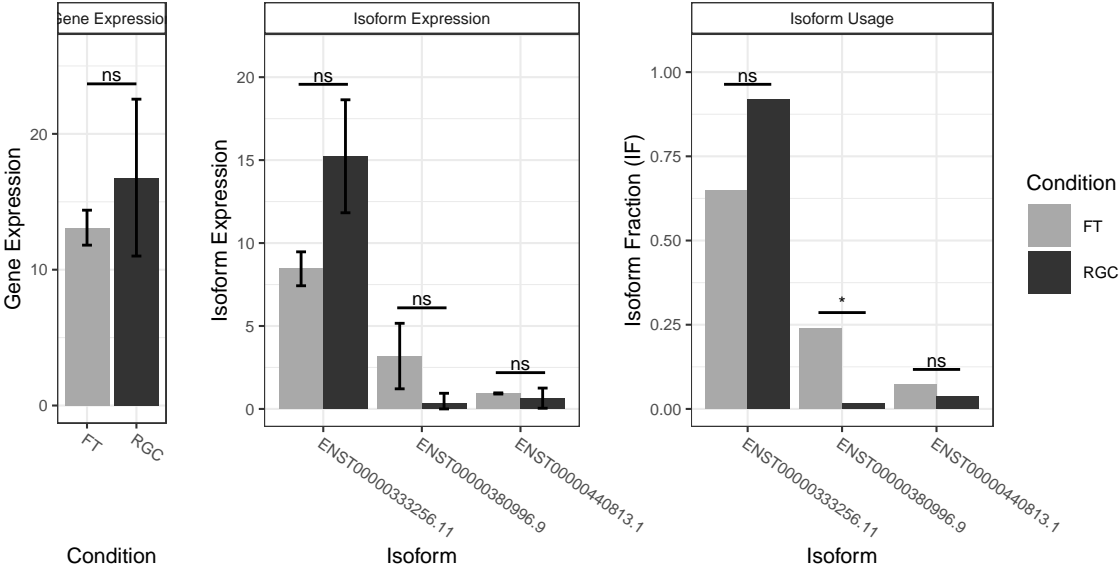
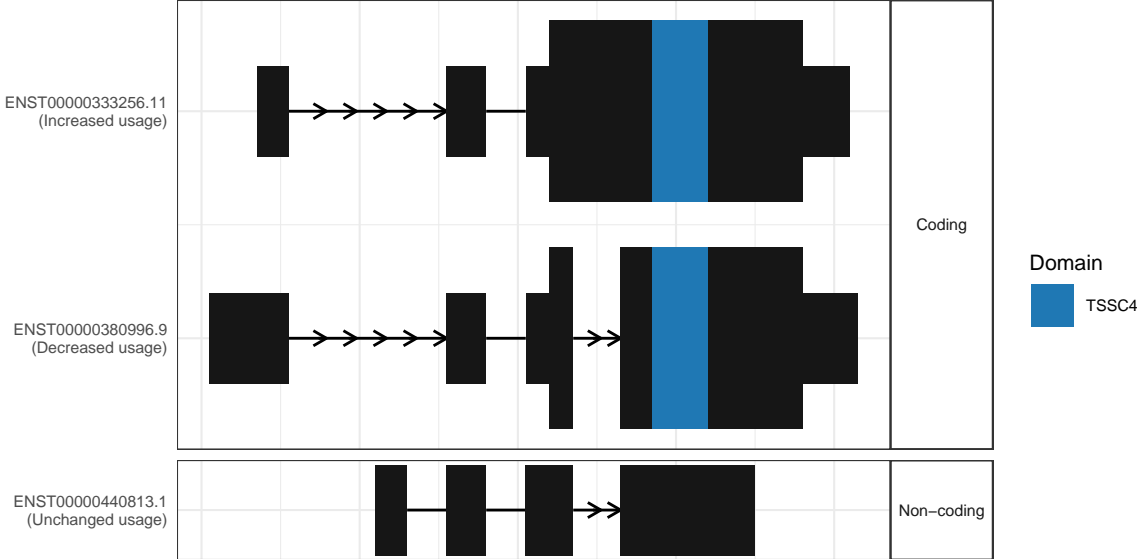
The isoform switch in PSIP1 (FT vs RGC)



Domain



The isoform switch in TSSC4 (FT vs RGC)

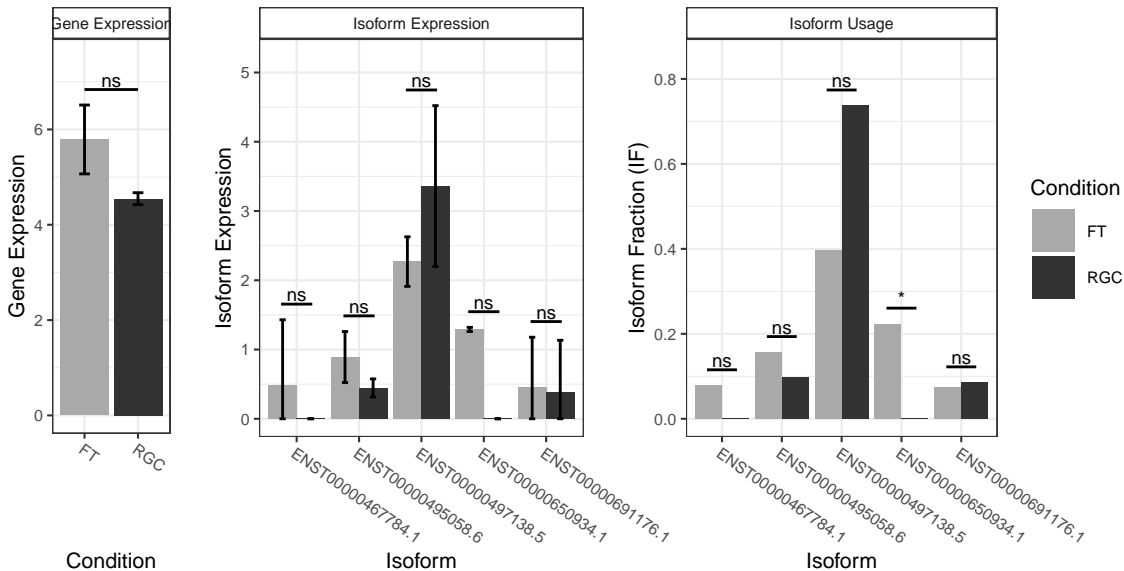
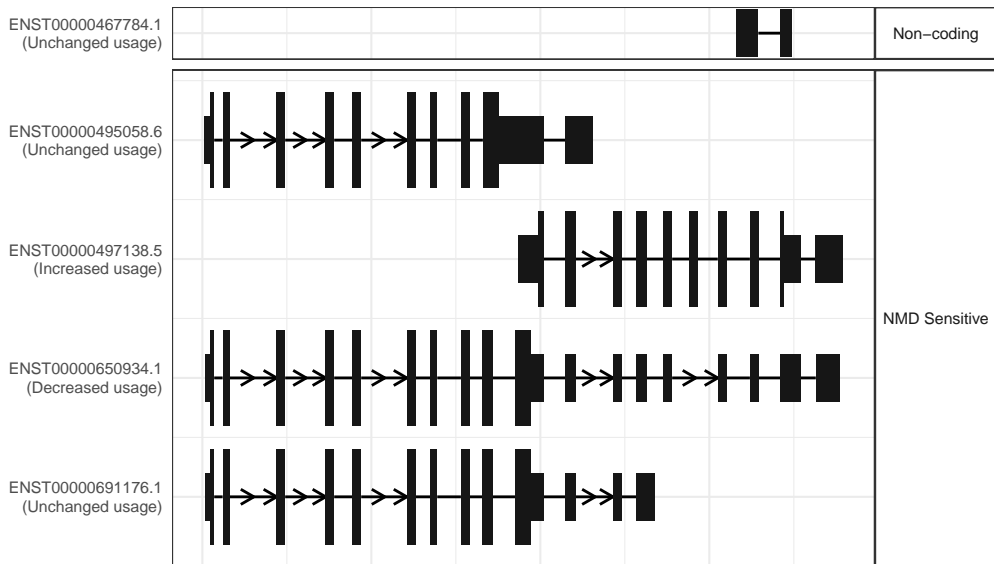


Domain

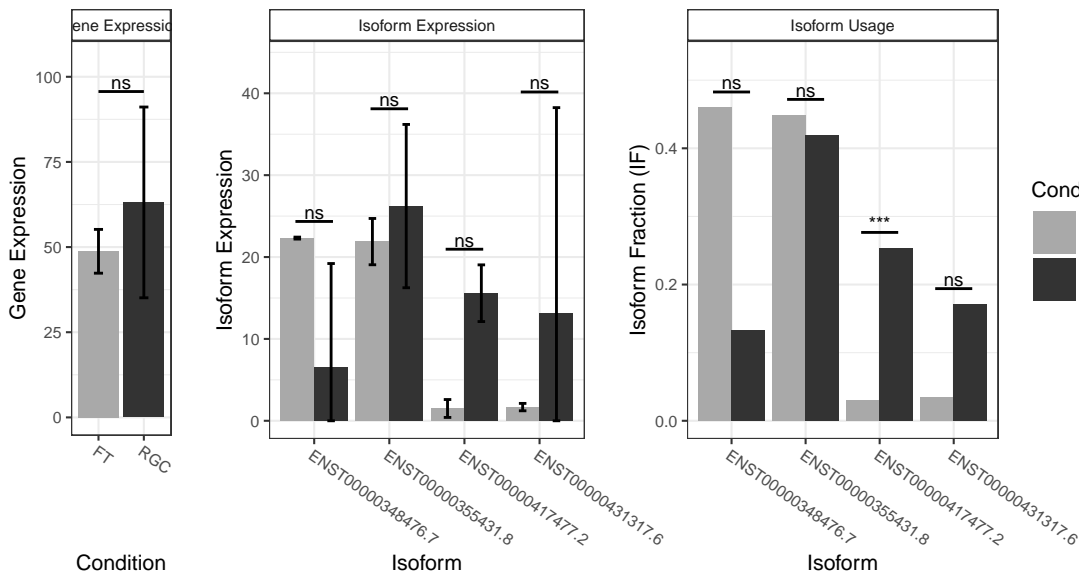
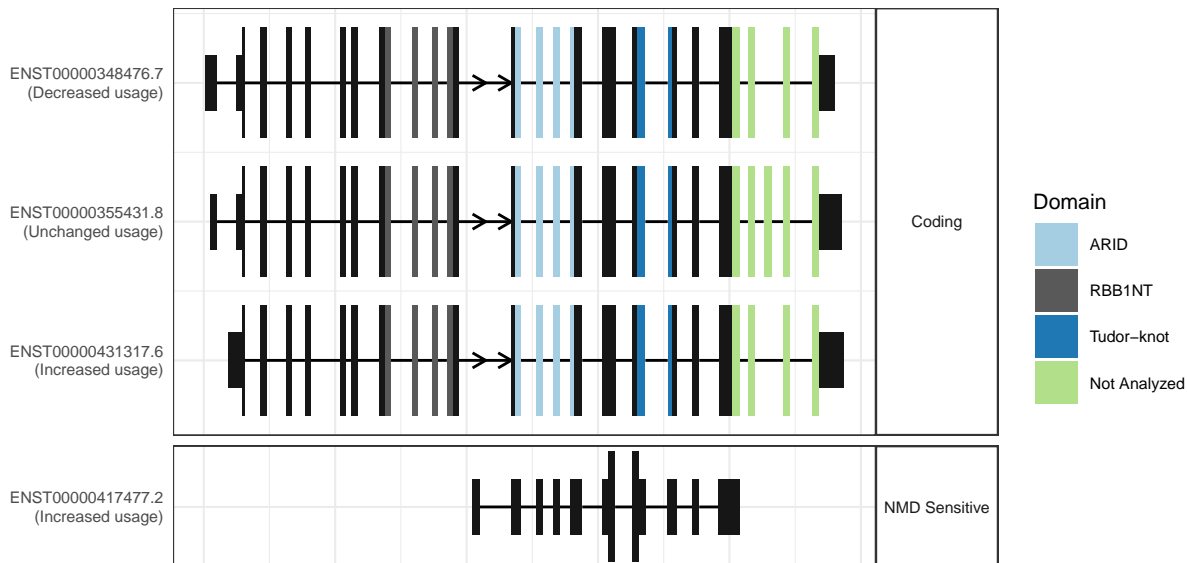


TSSC4





The isoform switch in (FT vs RGC)



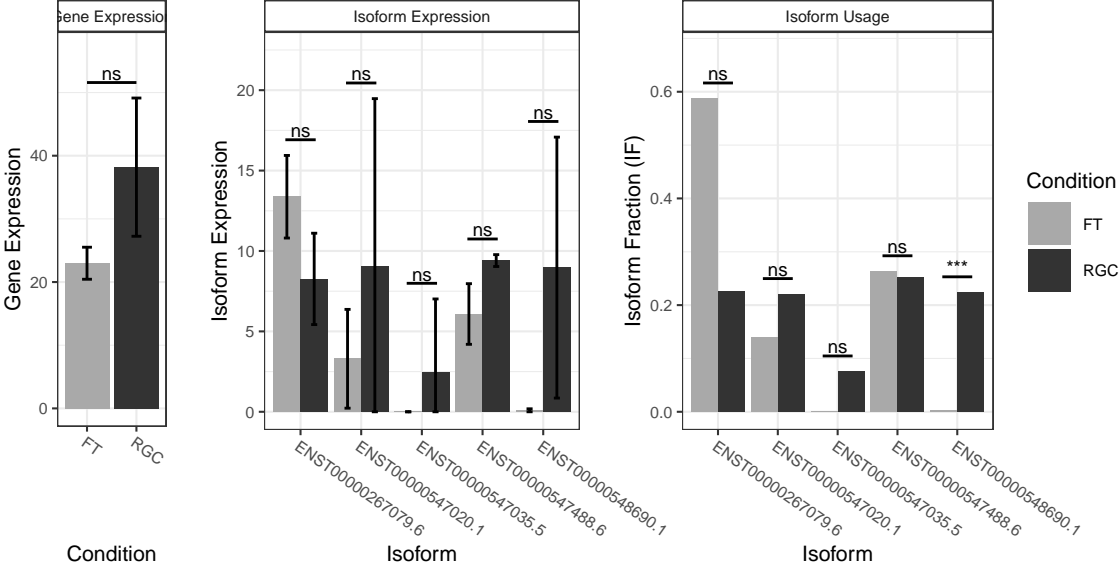
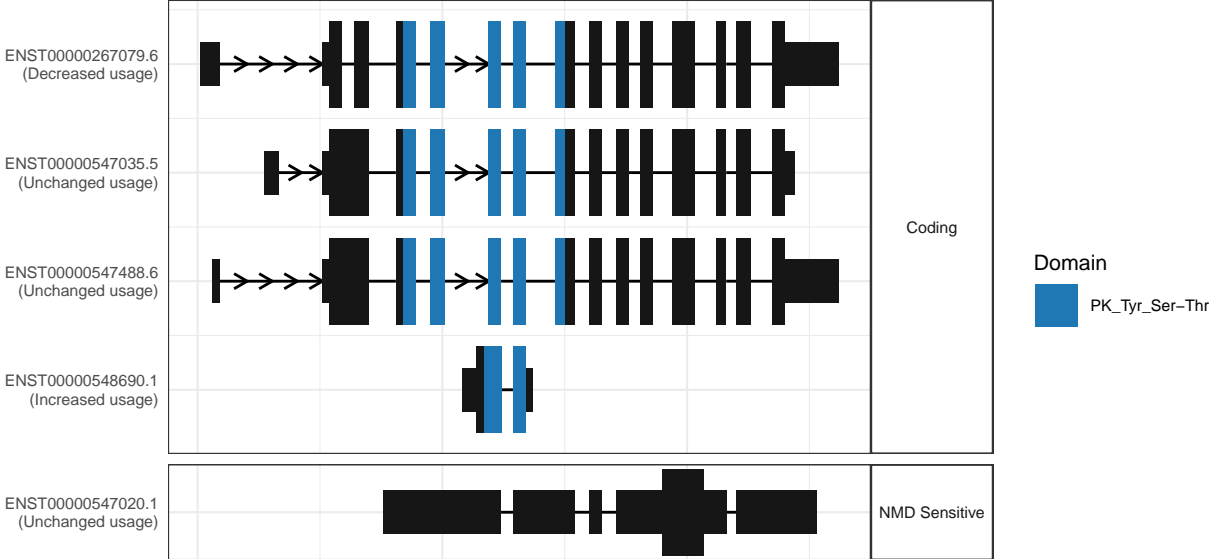
The isoform switch in ARID4A (FT vs RGC)



Domain

	ARID
	RBB1NT
	Tudor-knot
	Not Analyzed

The isoform switch in MAP3K12 (FT vs RGC)

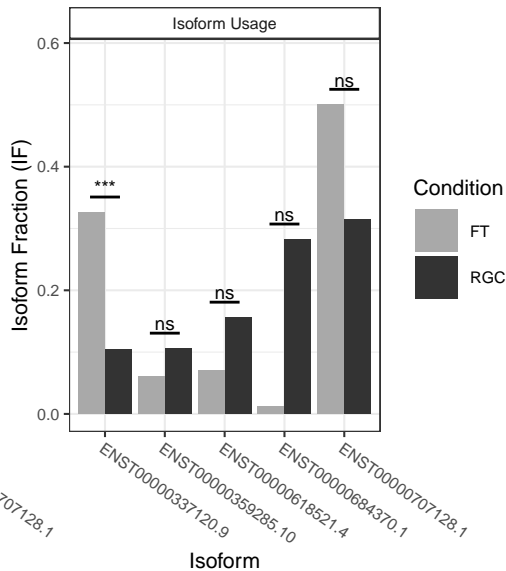
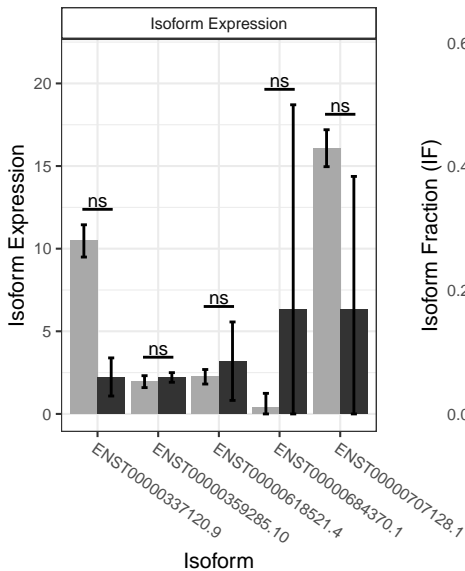
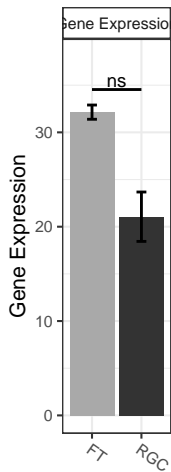
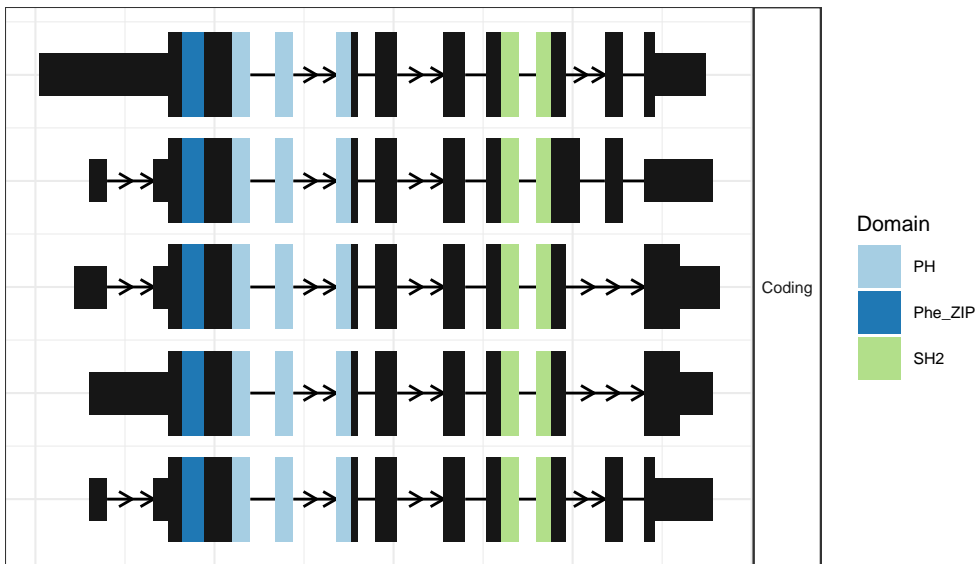


Domain



PK_Tyr_Ser-Thr

The isoform switch in SH2B1 (FT vs RGC)



Domain



PH

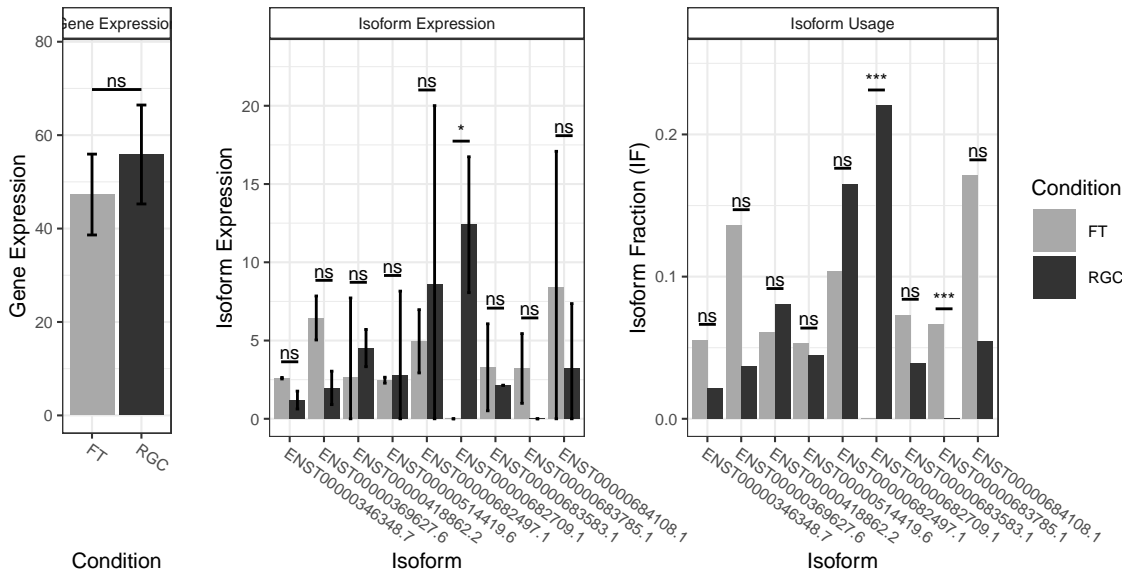
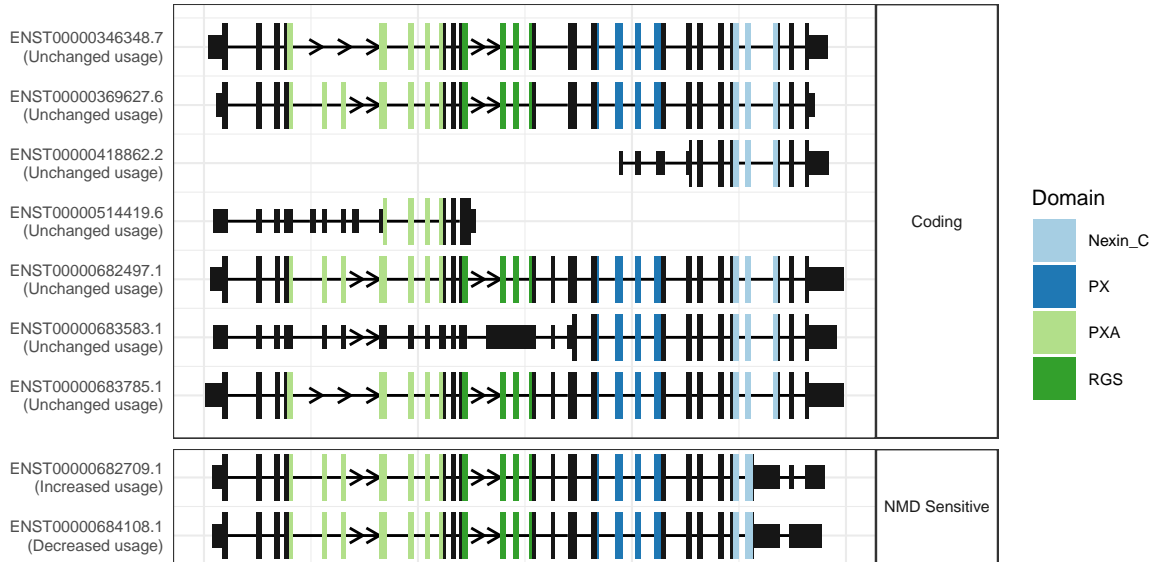


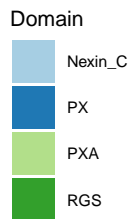
Phe_ZIP



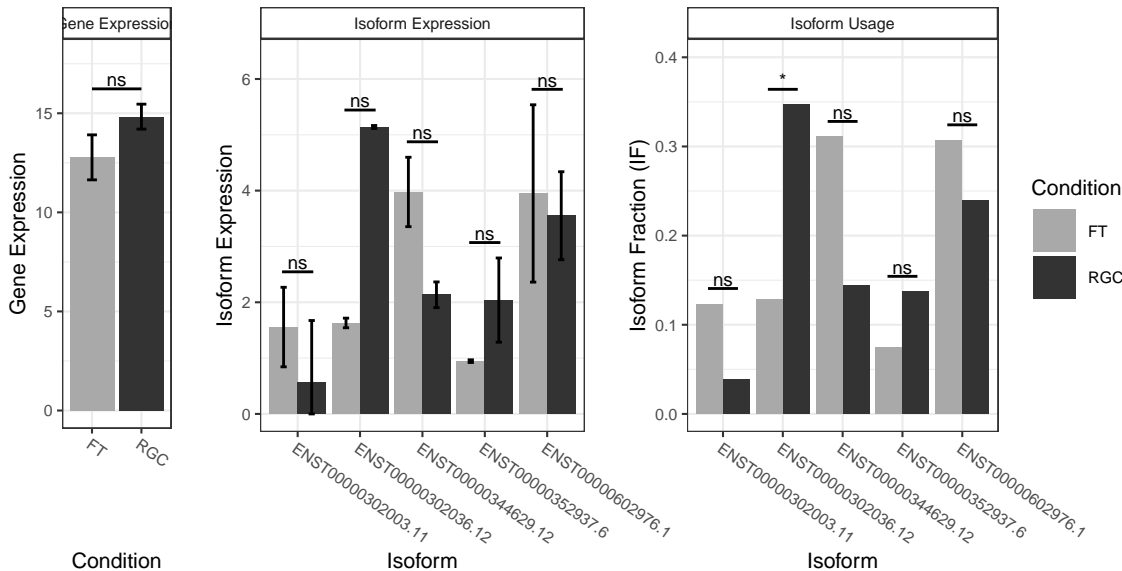
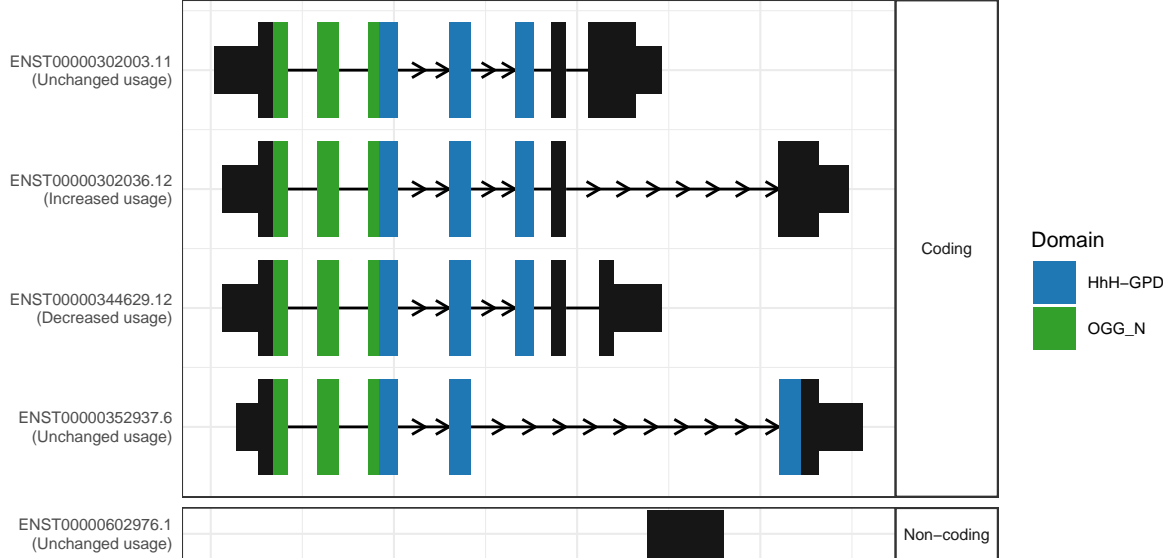
SH2

The isoform switch in SNX14 (FT vs RGC)





The isoform switch in OGG1 (FT vs RGC)



Domain

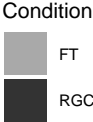
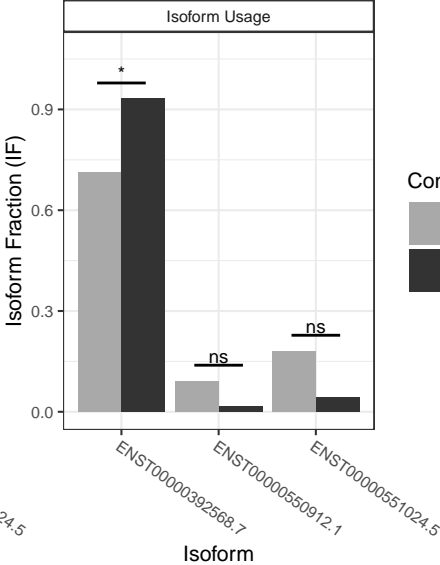
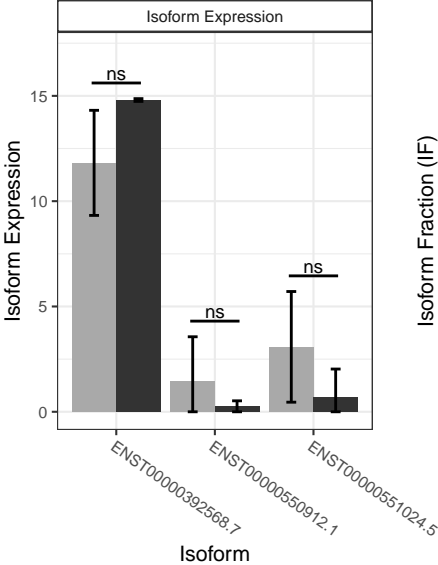
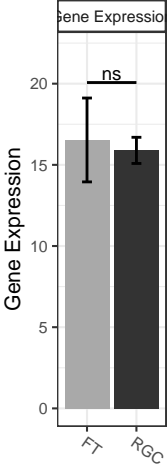
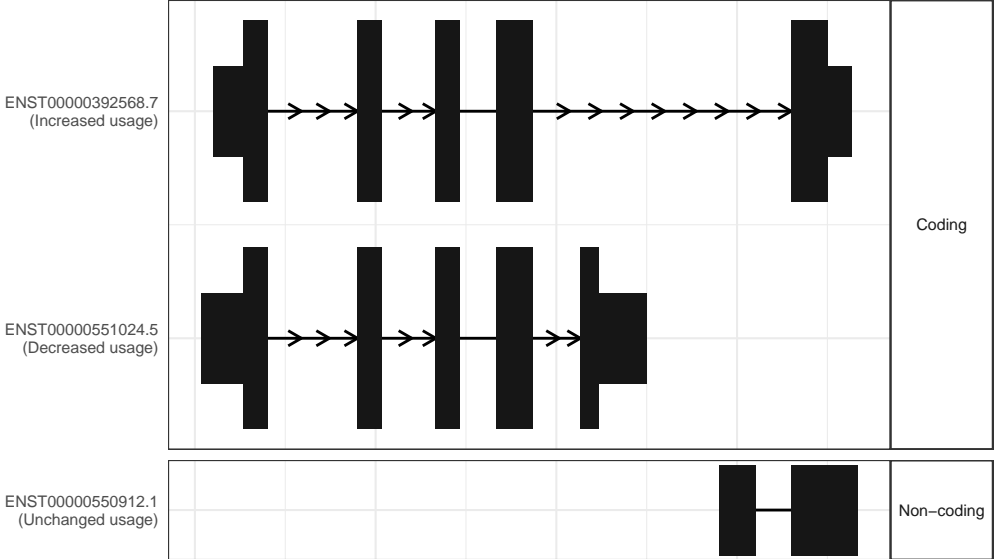


HhH-GPD

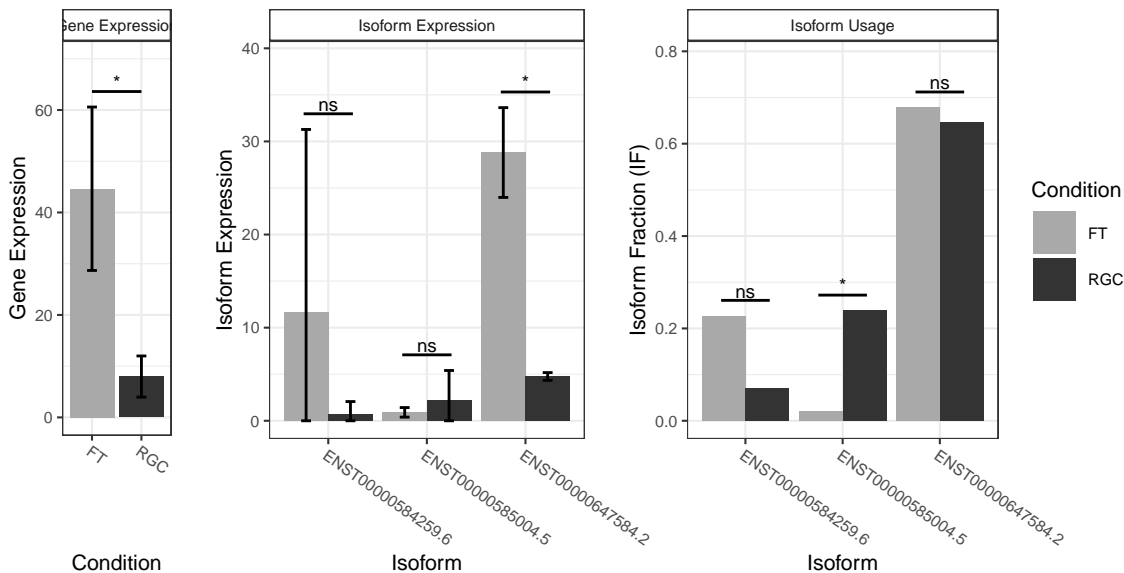
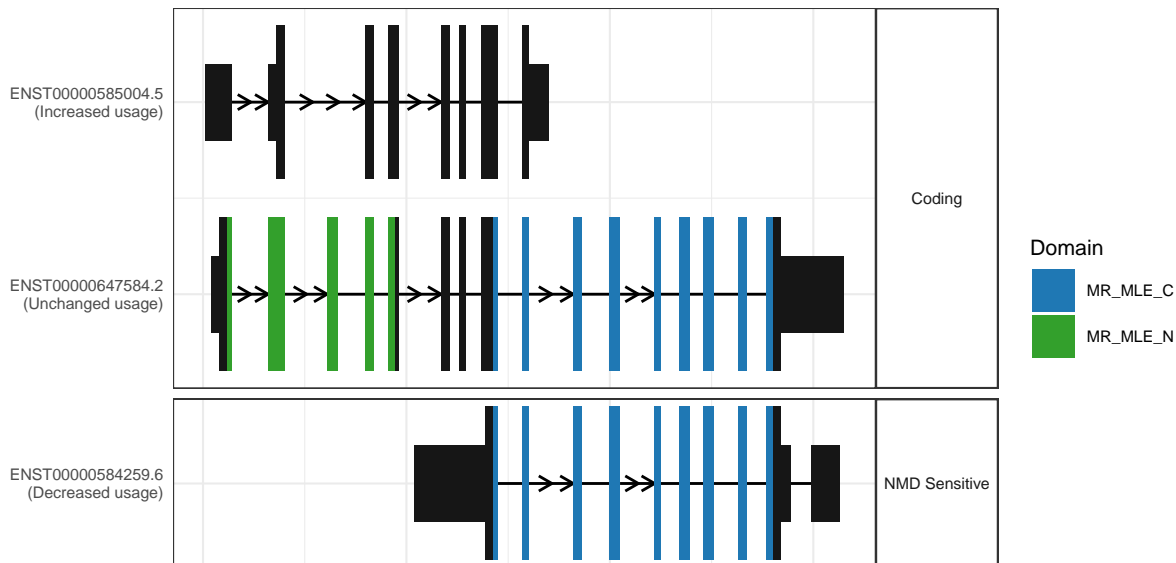


OGG_N

The isoform switch in NUDT14 (FT vs RGC)



The isoform switch in ENOSF1 (FT vs RGC)



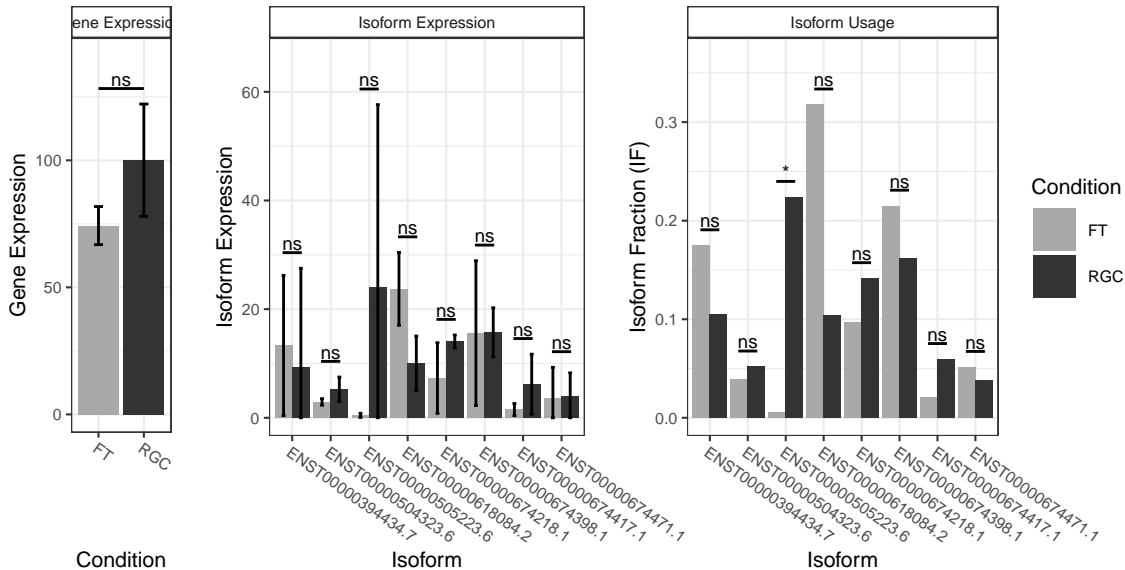
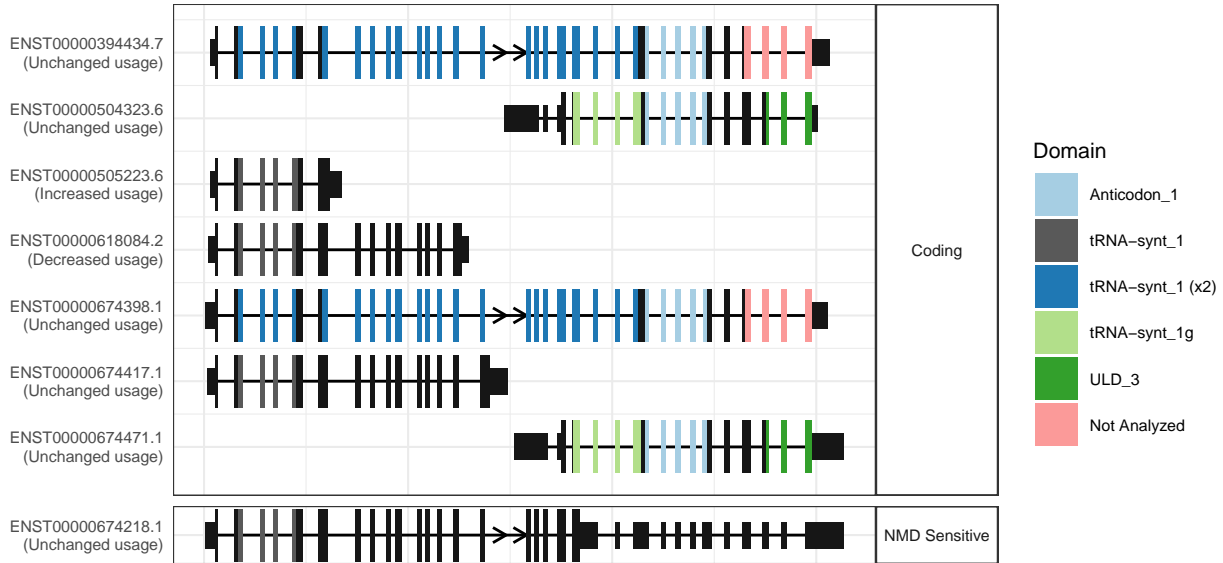
Domain









MR_MLE_C

MR_MLE_N

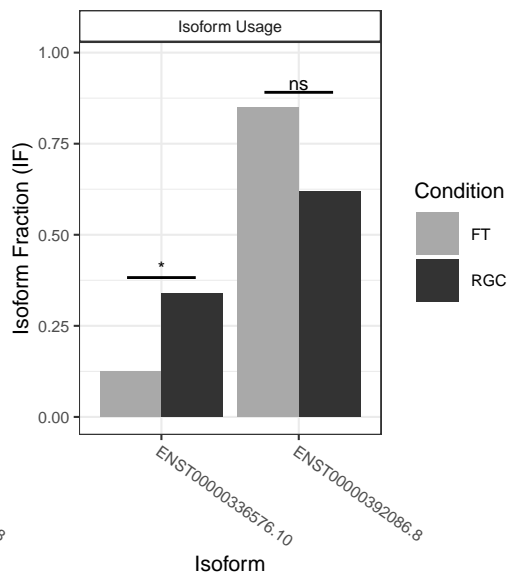
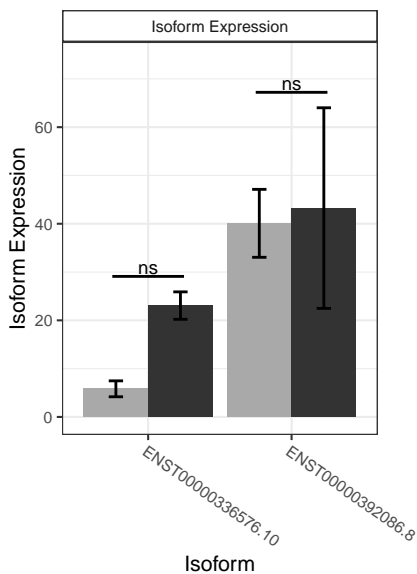
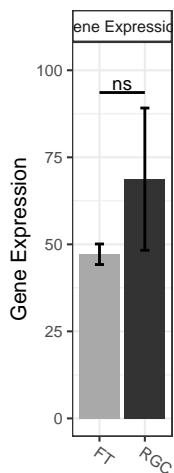
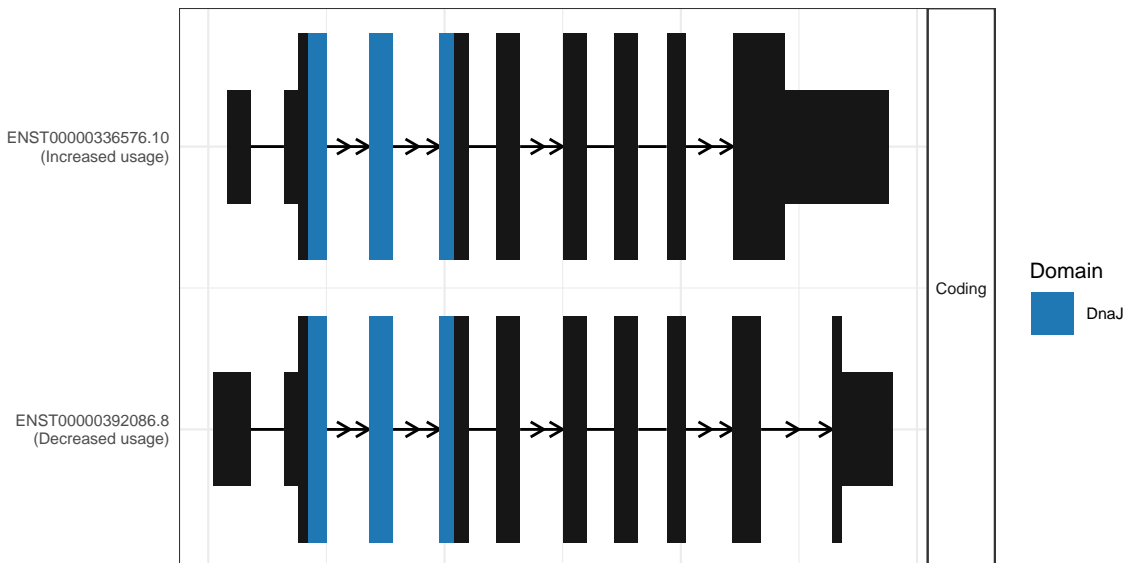
The isoform switch in LARS1 (FT vs RGC)



Domain

	Anticodon_1
	tRNA-synt_1
	tRNA-synt_1 (x2)
	tRNA-synt_1g
	ULD_3
	Not Analyzed

The isoform switch in DNAJB2 (FT vs RGC)

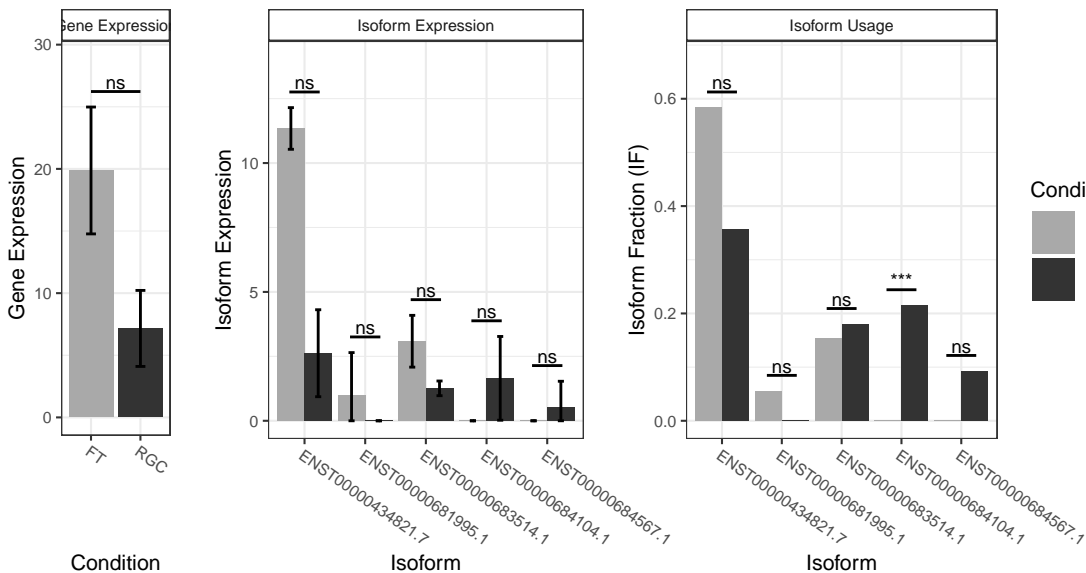
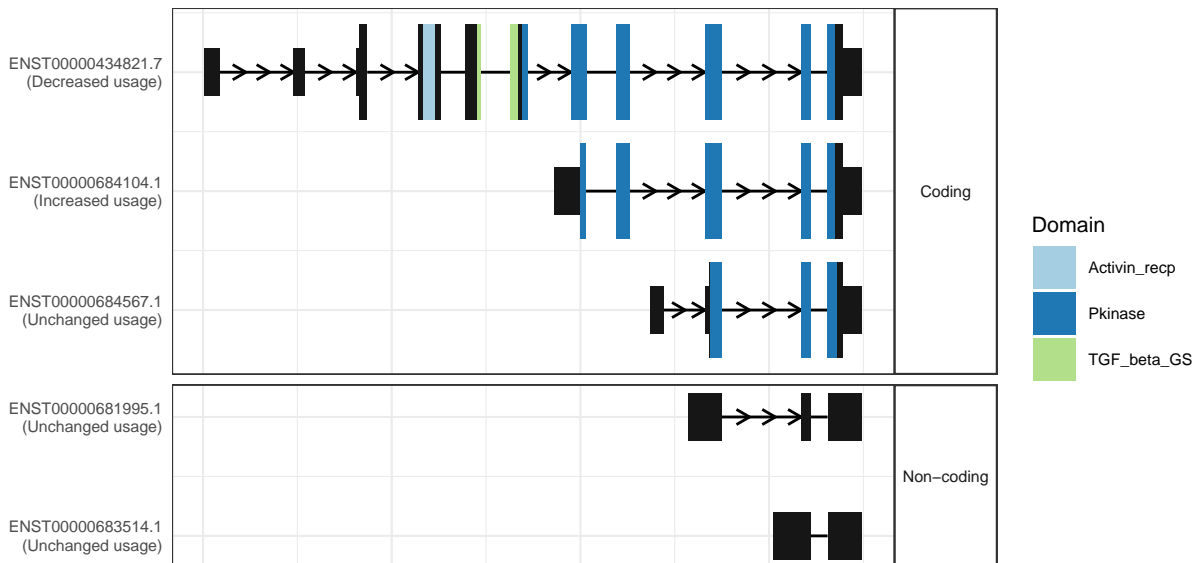


Domain



DnaJ

The isoform switch in ACVR1 (FT vs RGC)



Domain



Activin_recp

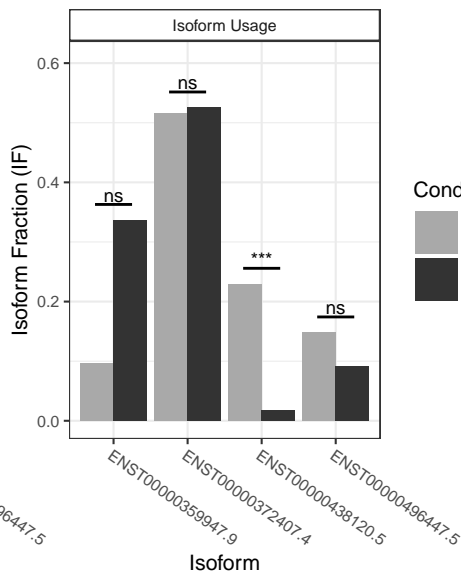
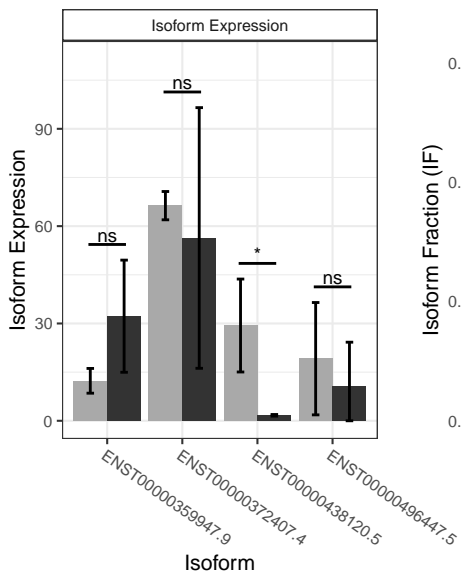
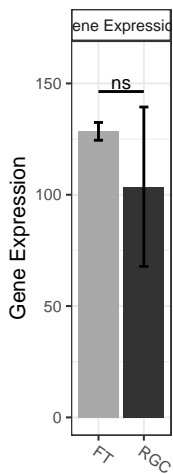
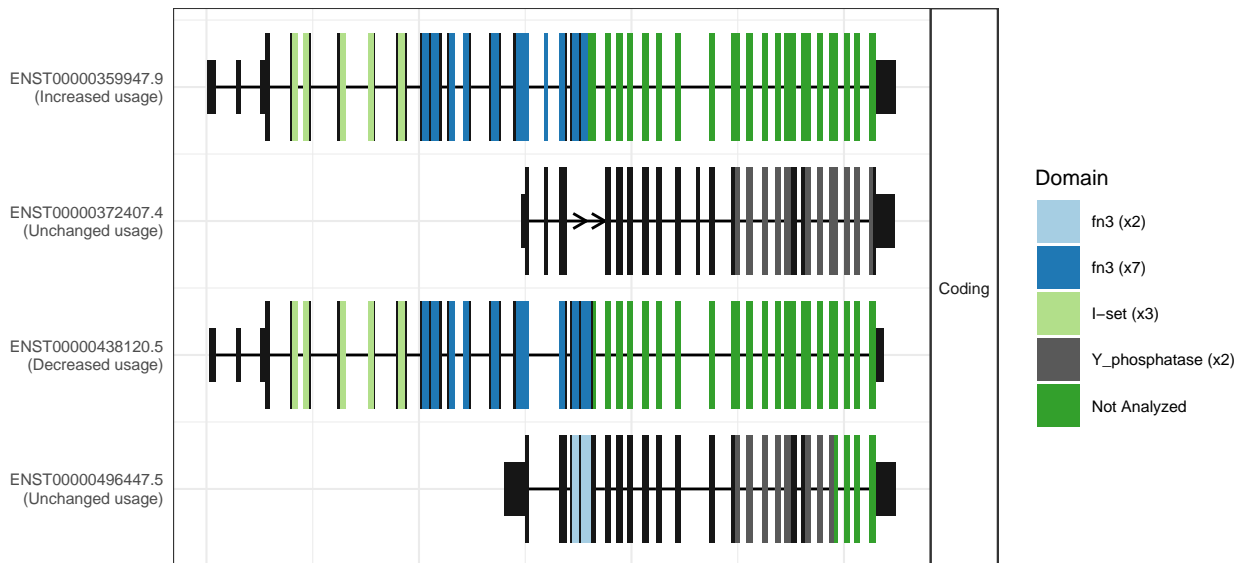


Pkinase



TGF_beta_GS

The isoform switch in PTPRF (FT vs RGC)



Condition






Isoform

Isoform

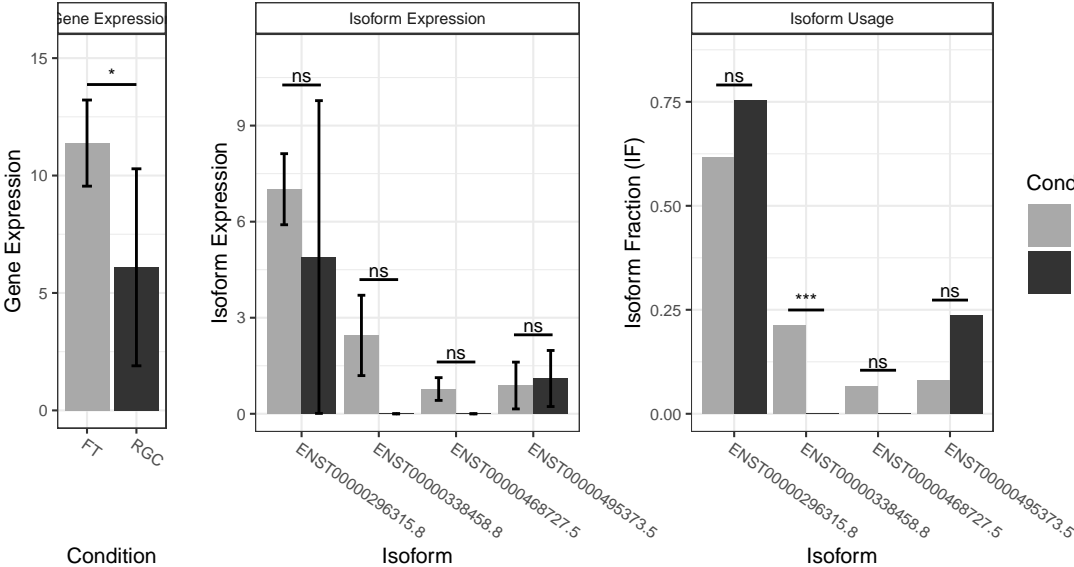
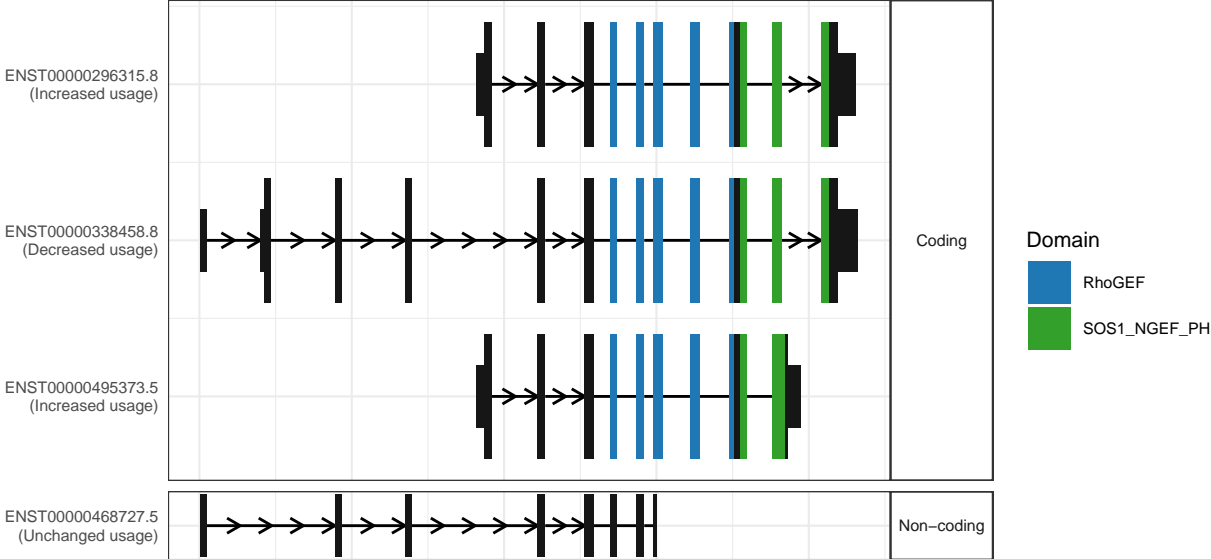
Condition



Domain

	fn3 (x2)
	fn3 (x7)
	l-set (x3)
	Y_phosphatase (x2)
	Not Analyzed

The isoform switch in ARHGEF3 (FT vs RGC)



Domain

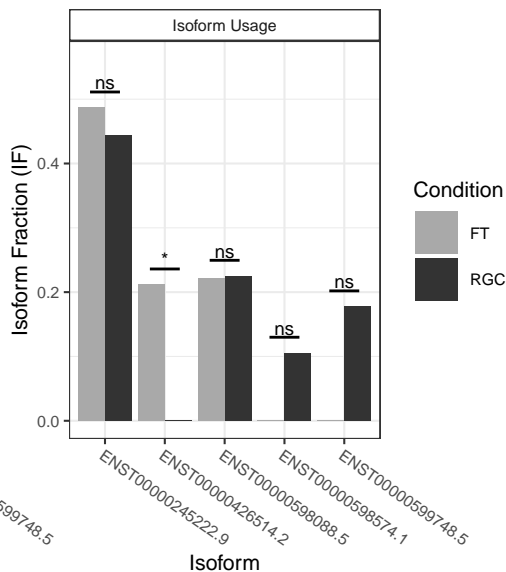
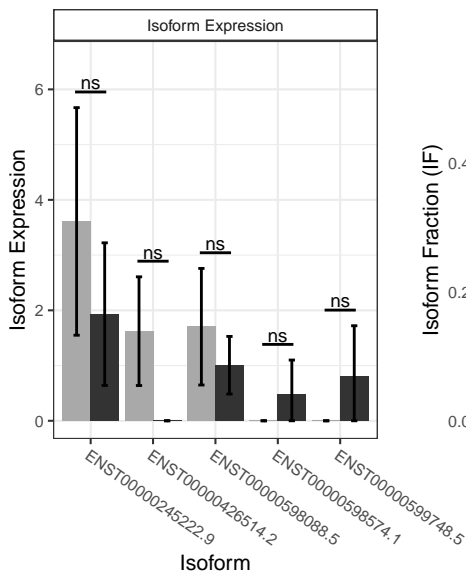
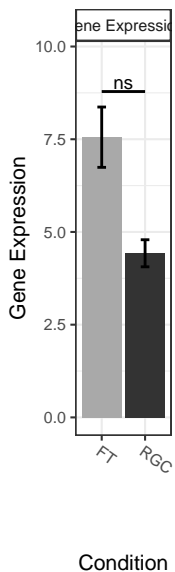
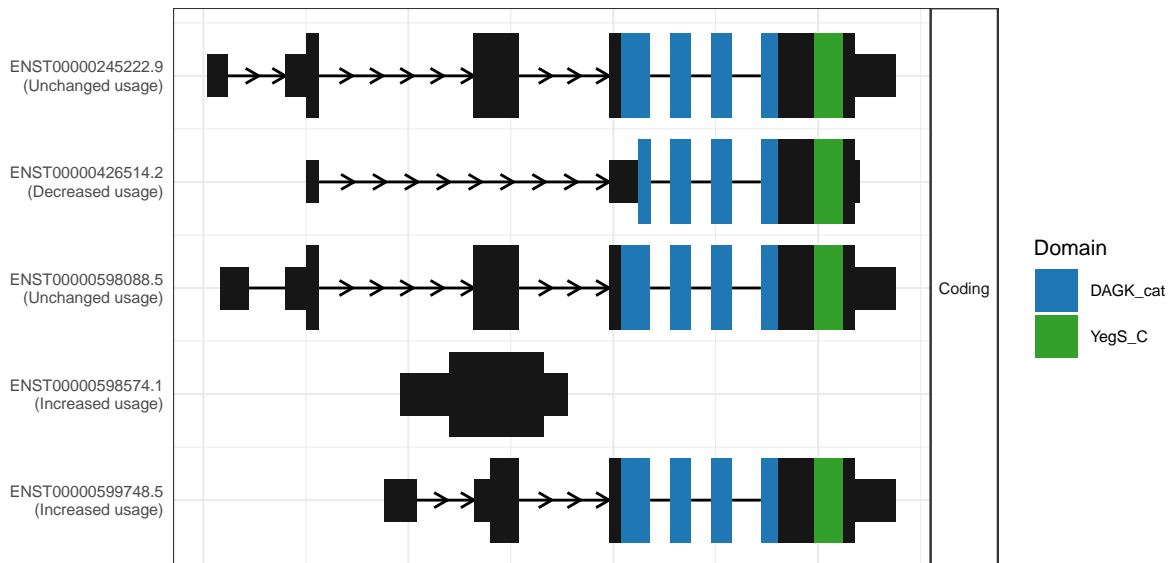


RhoGEF

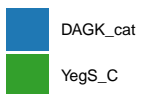


SOS1_NGEF_PH

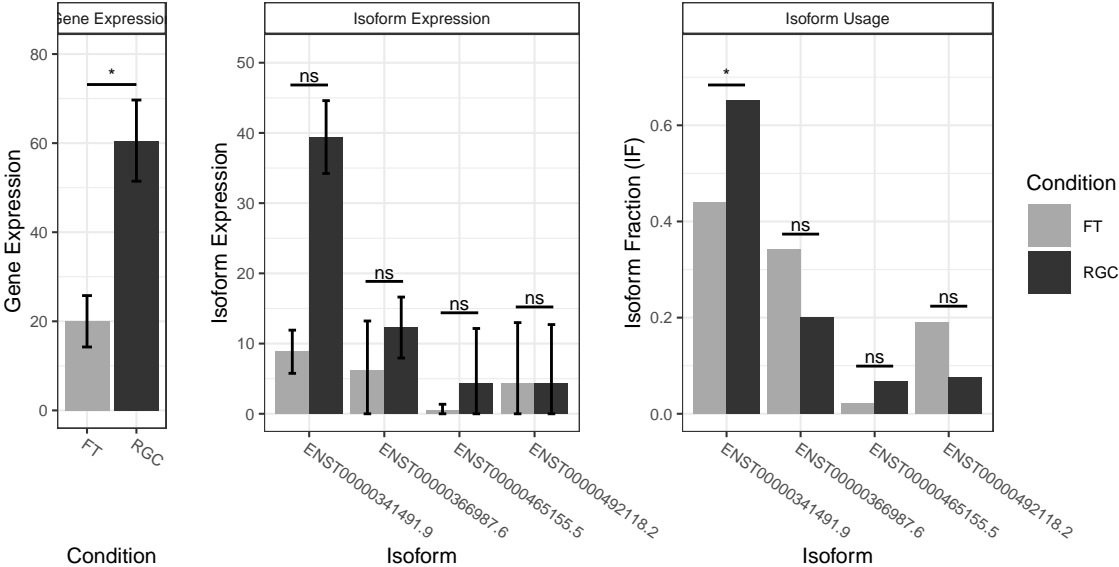
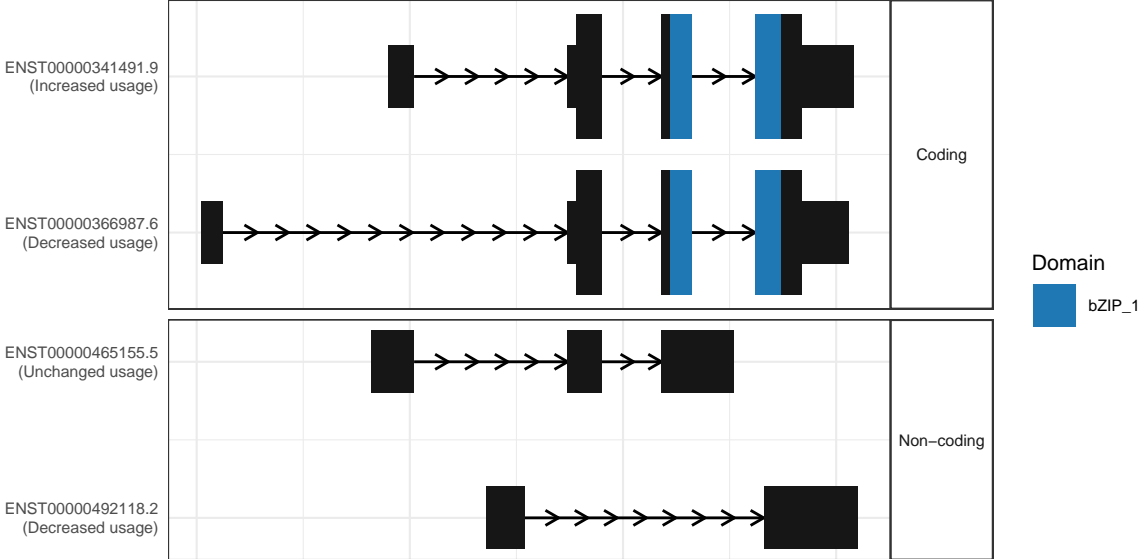
The isoform switch in SPHK2 (FT vs RGC)



Domain



The isoform switch in ATF3 (FT vs RGC)

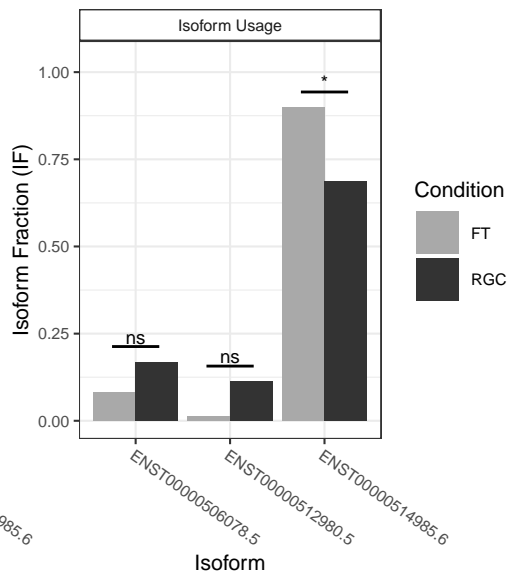
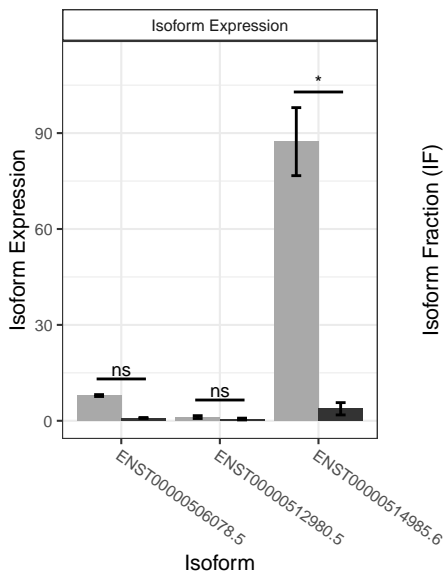
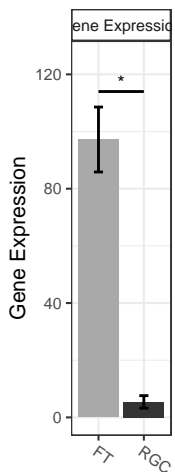
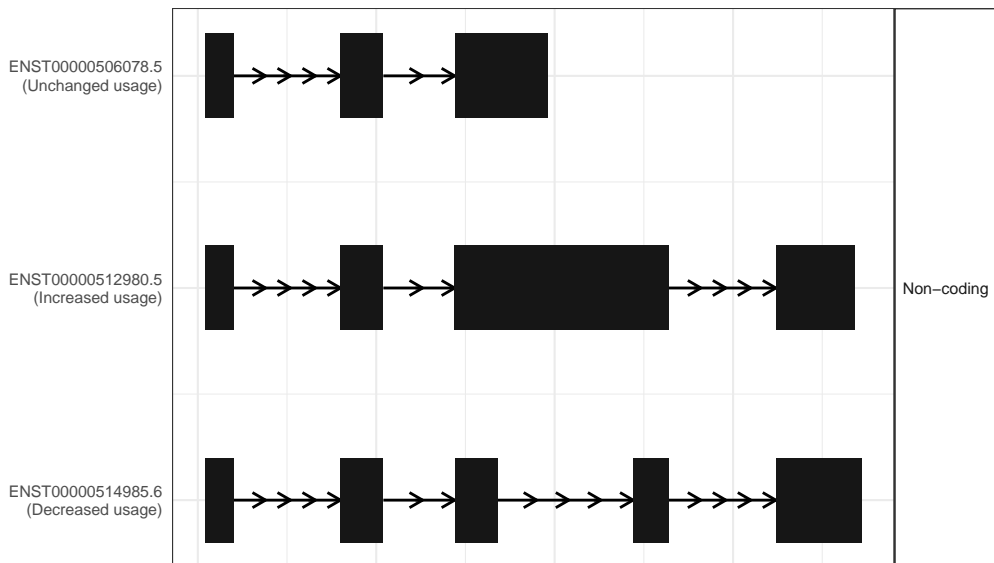


Domain

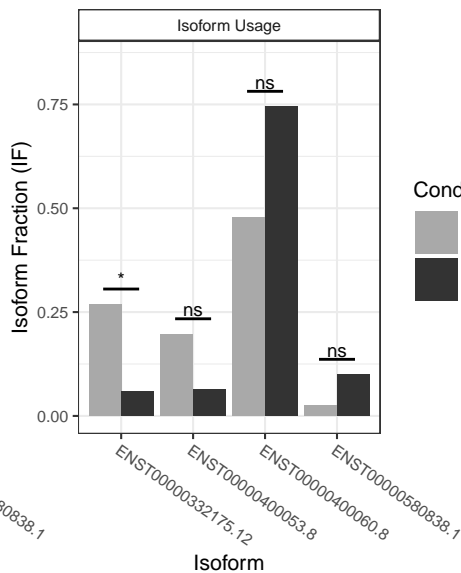
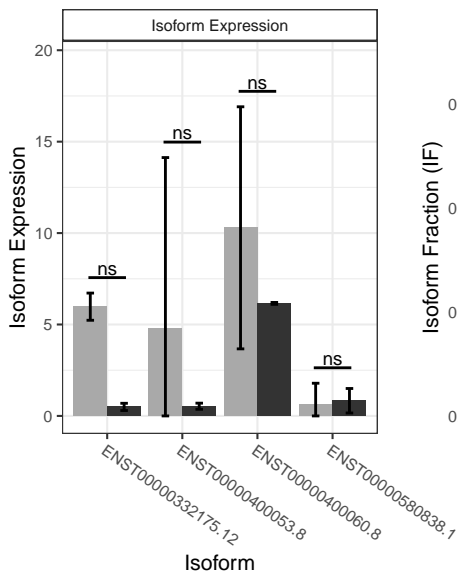
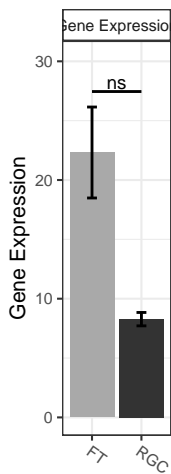
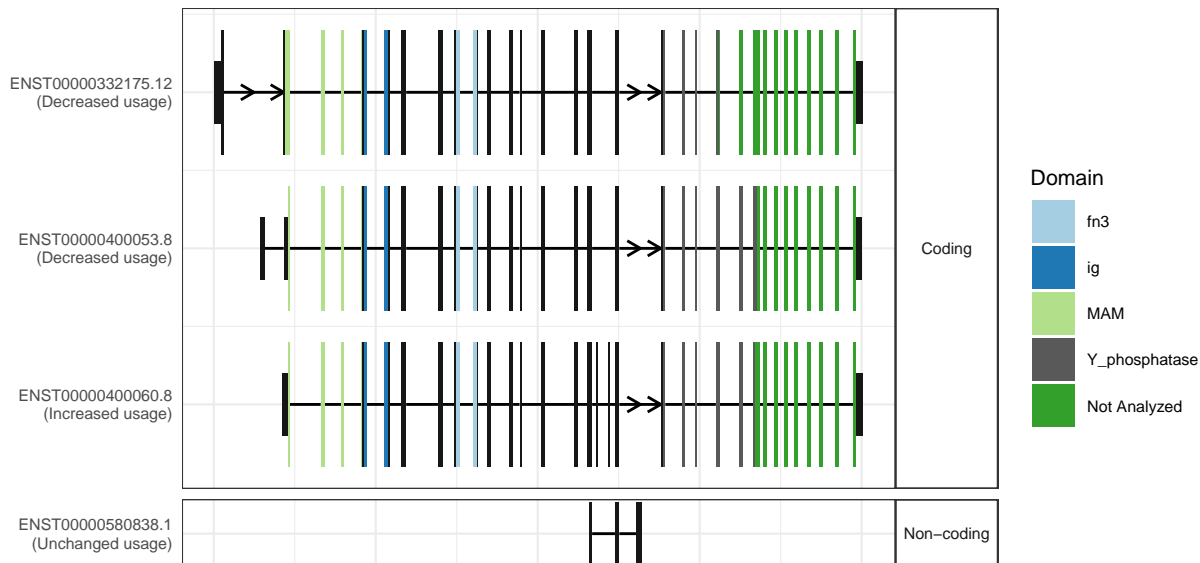


bZIP_1

The isoform switch in SELENOP (FT vs RGC)



The isoform switch in PTPRM (FT vs RGC)



Domain



fn3



ig



MAM

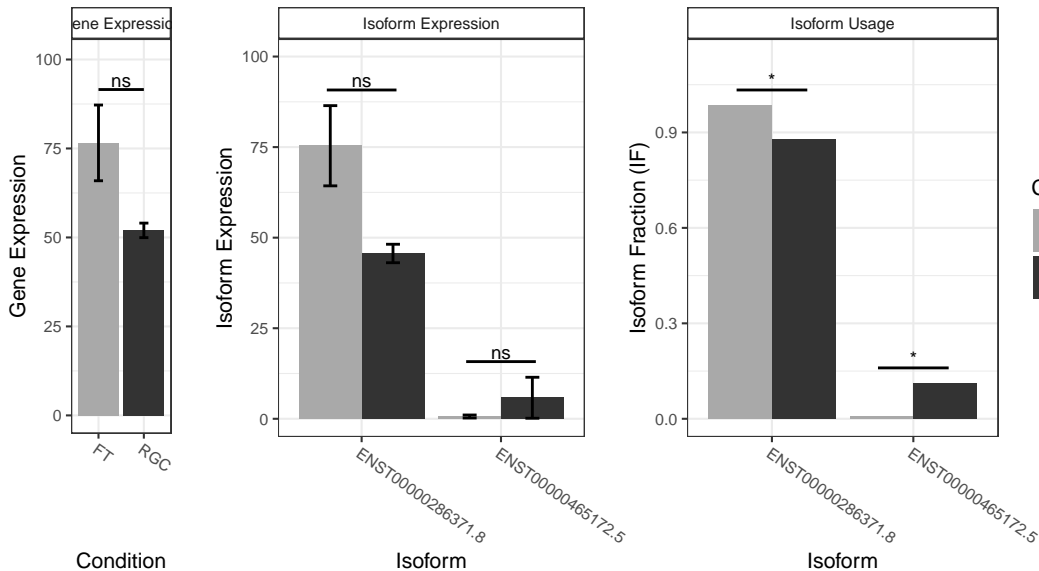
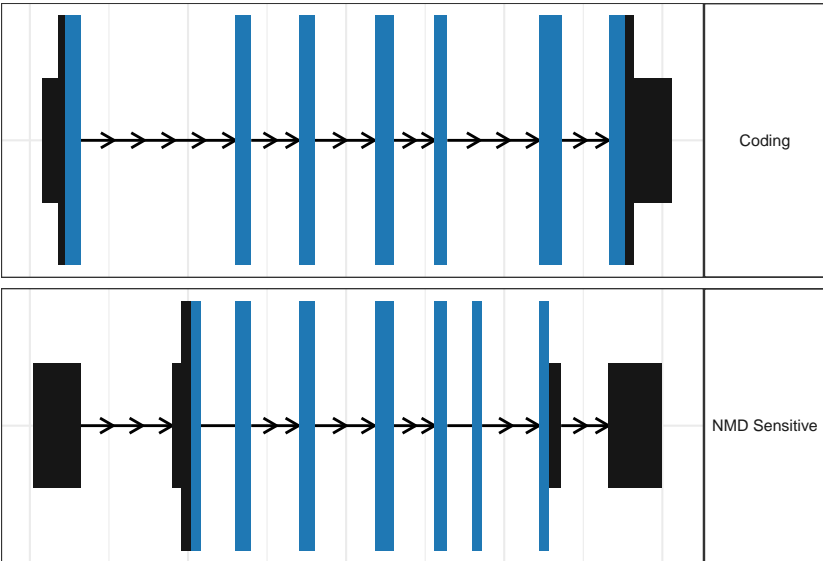


Y_phosphatase



Not Analyzed

The isoform switch in ATP1B3 (FT vs RGC)

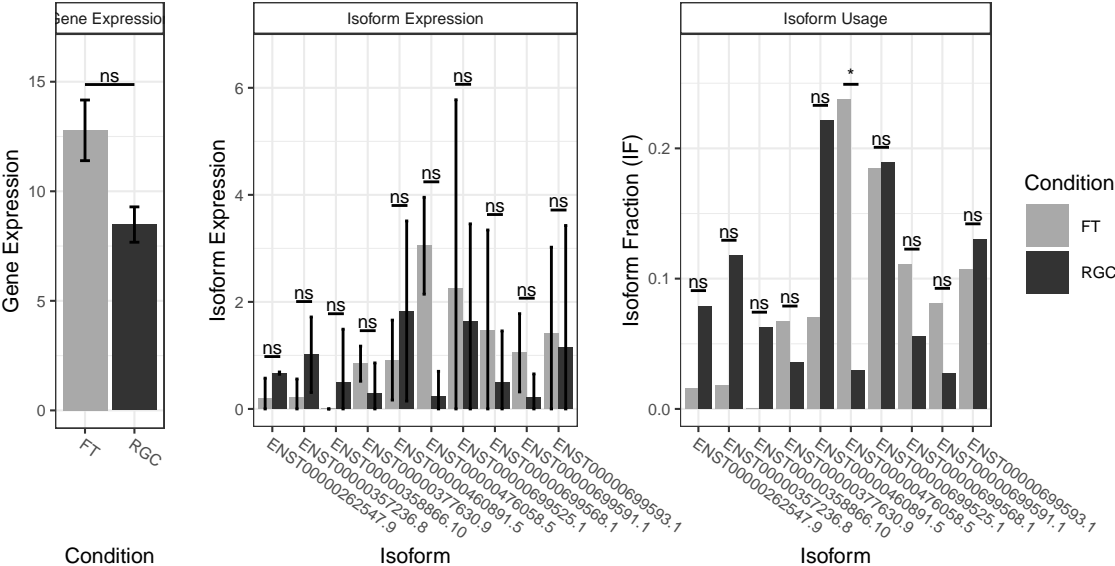
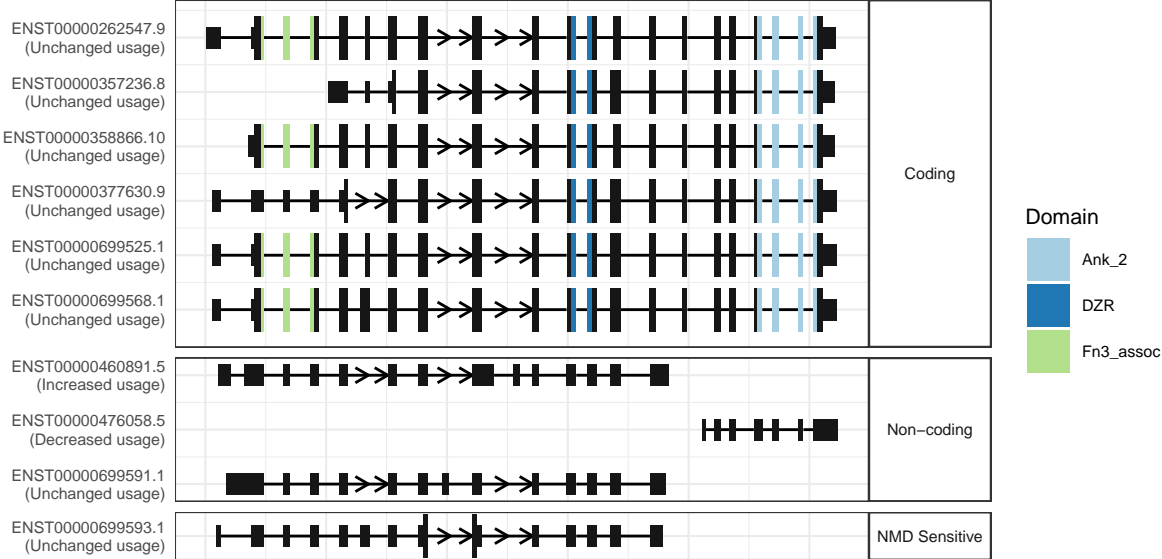


Domain



Na_K-ATPase

The isoform switch in DZANK1 (FT vs RGC)



Domain



Ank_2

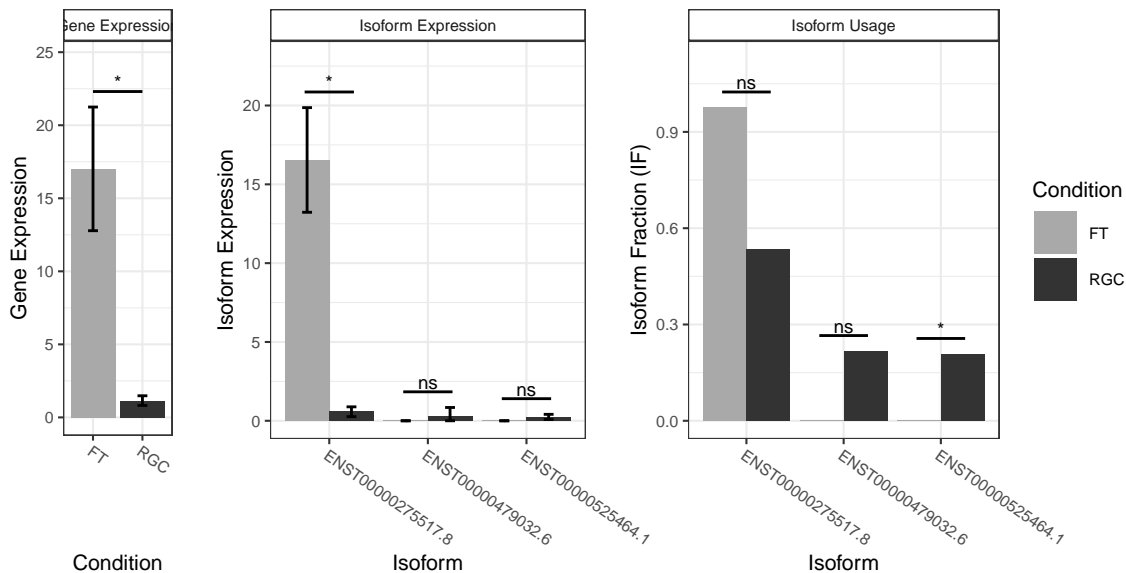
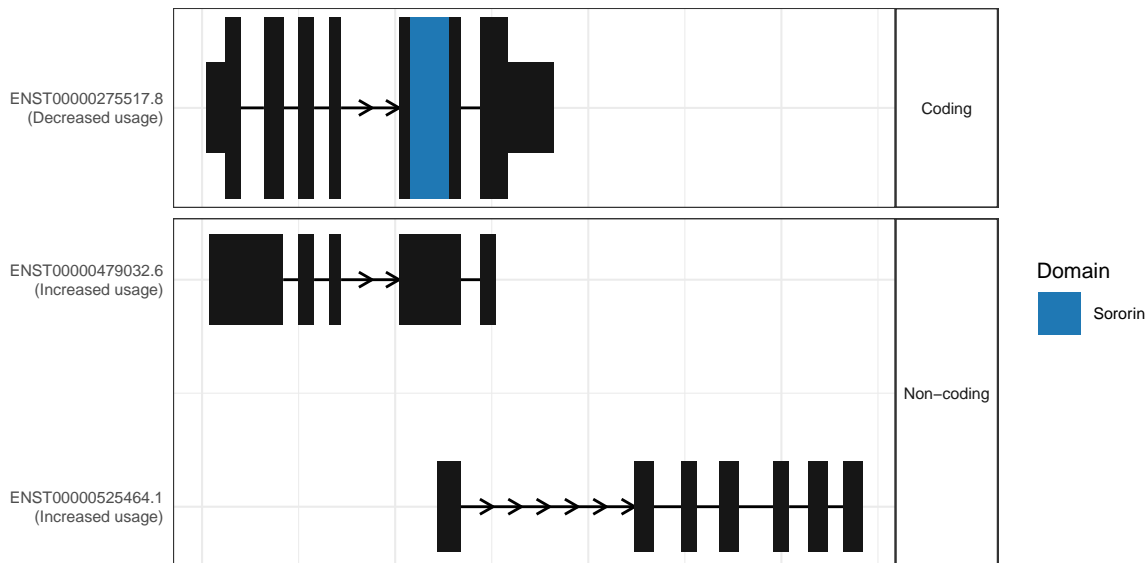


DZR



Fn3_assoc

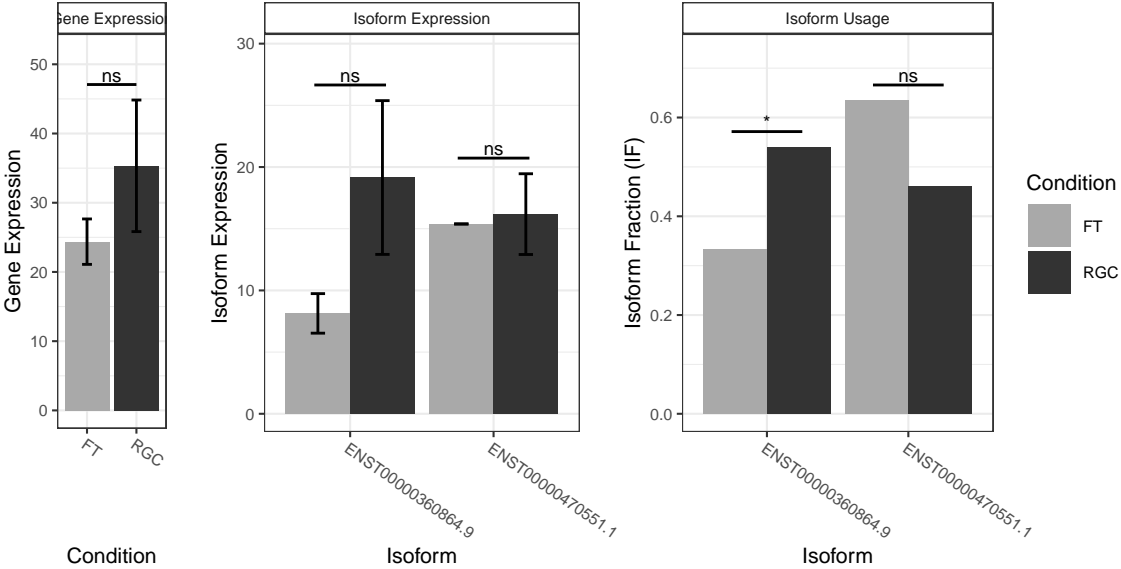
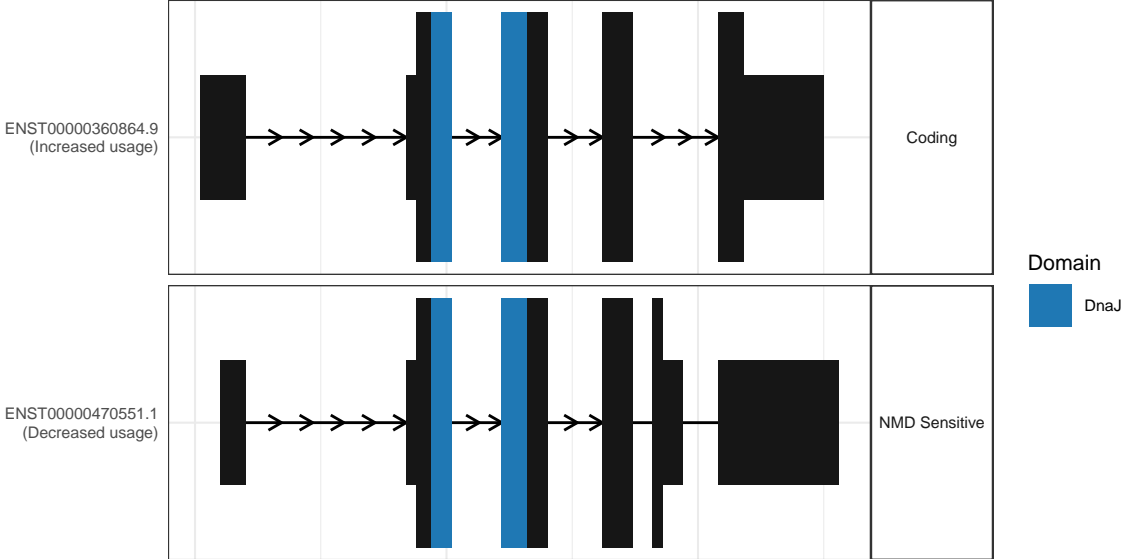
The isoform switch in CDCA5 (FT vs RGC)



Domain

 Sororin

The isoform switch in DNAJC5 (FT vs RGC)

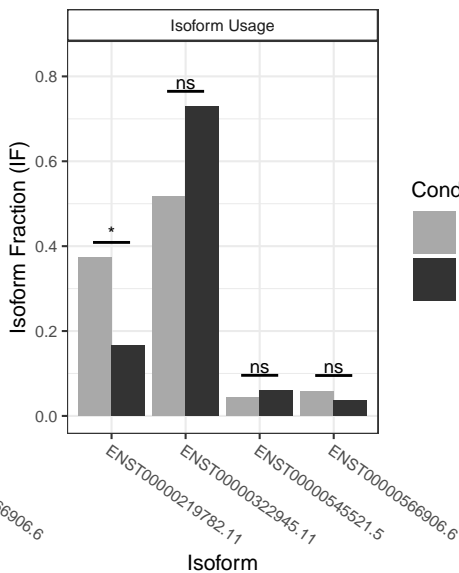
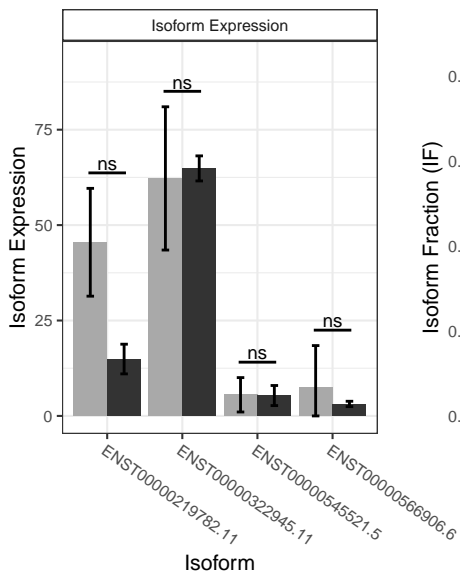
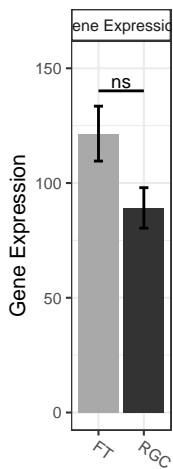
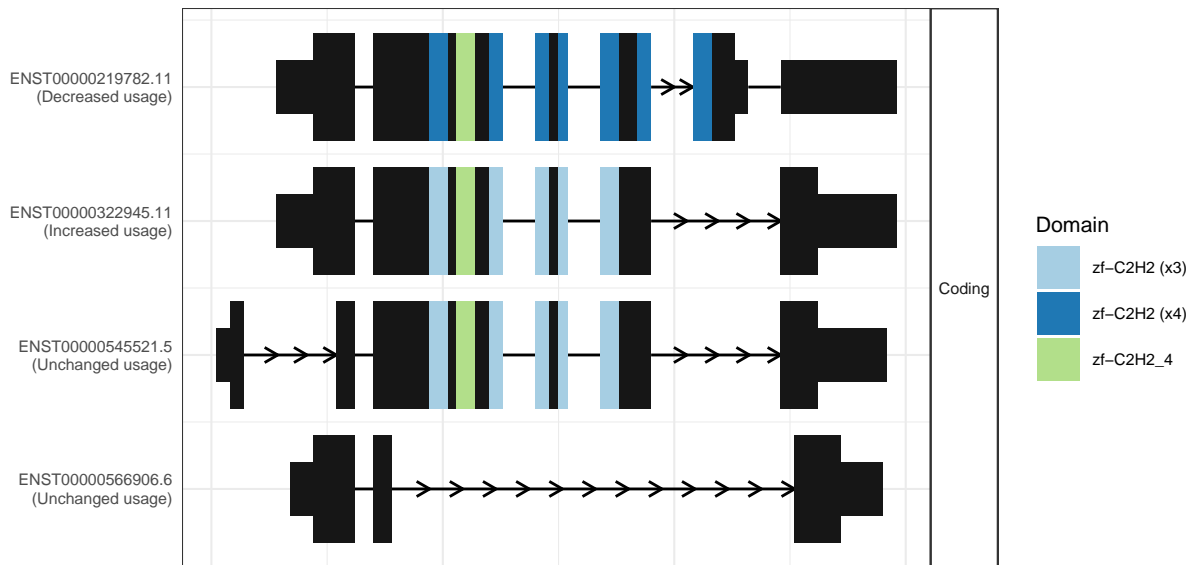


Domain



DnaJ

The isoform switch in MAZ (FT vs RGC)



Domain



zf-C2H2 (x3)

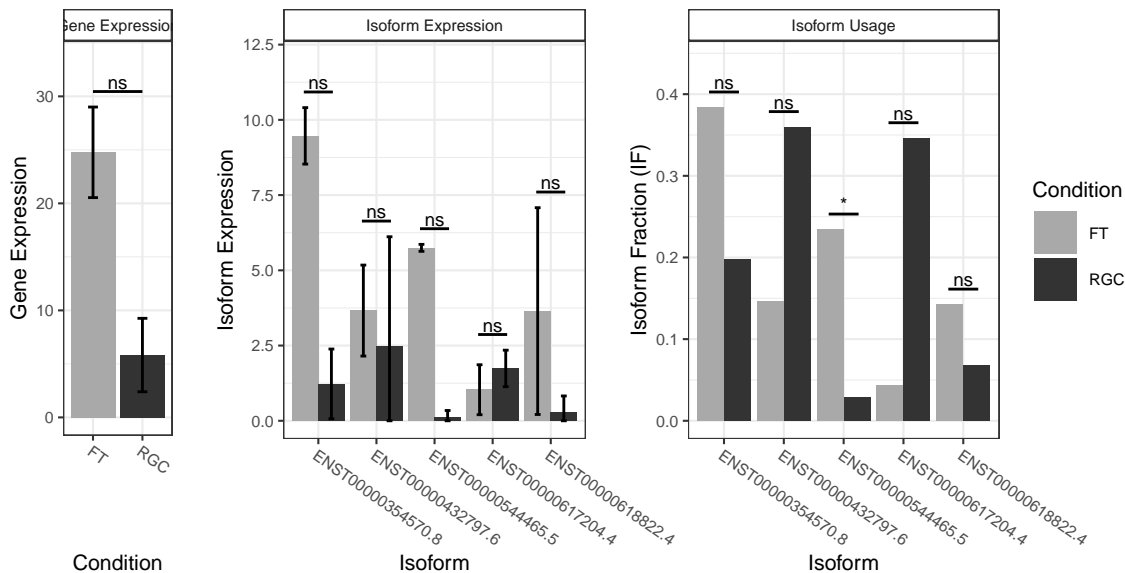
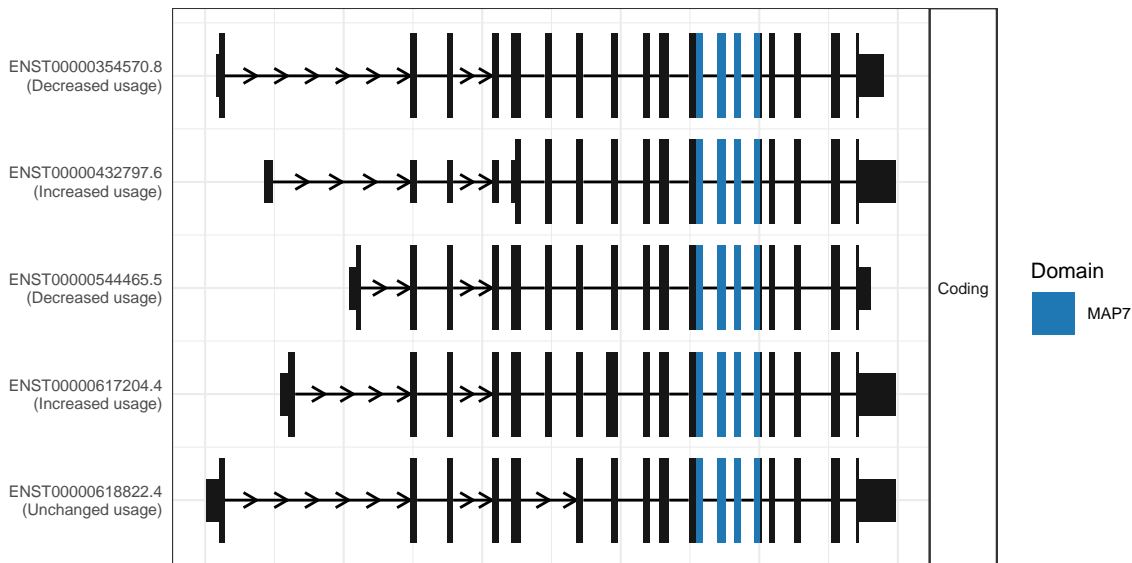


zf-C2H2 (x4)



zf-C2H2_4

The isoform switch in MAP7 (FT vs RGC)

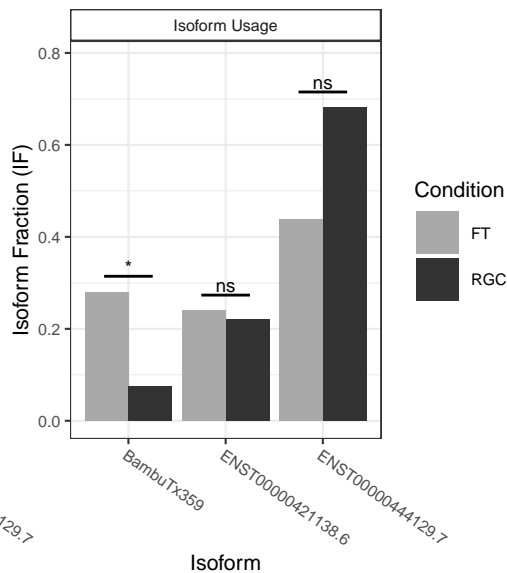
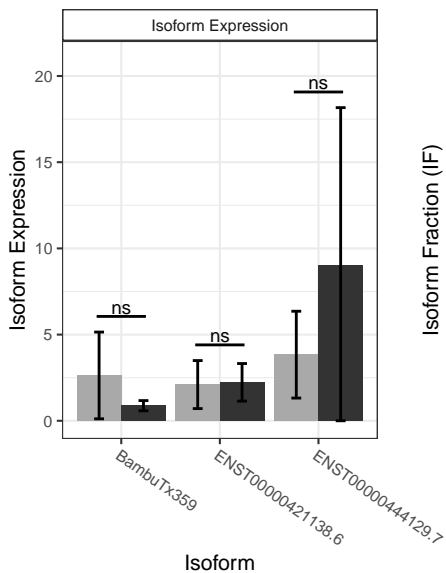
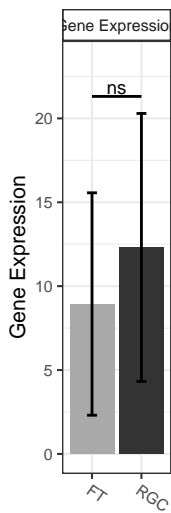
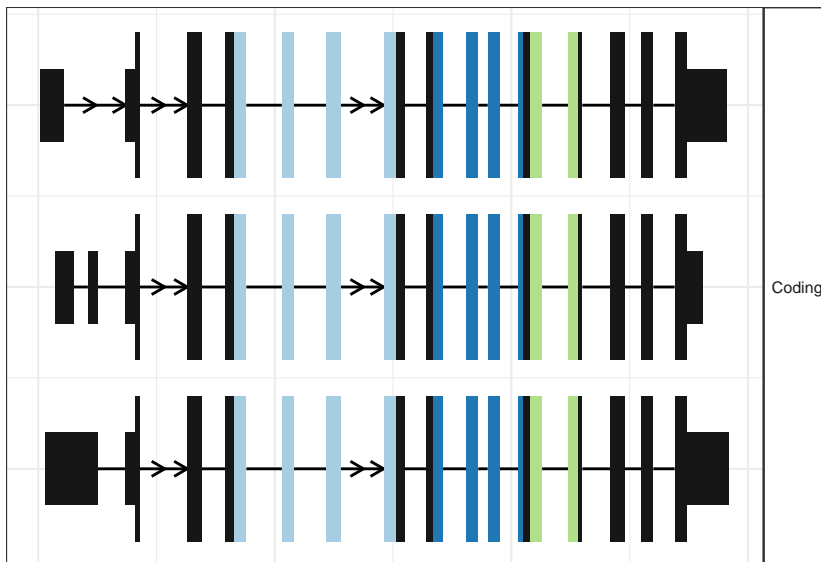


Domain



MAP7

The isoform switch in RECQL (FT vs RGC)



Domain



DEAD

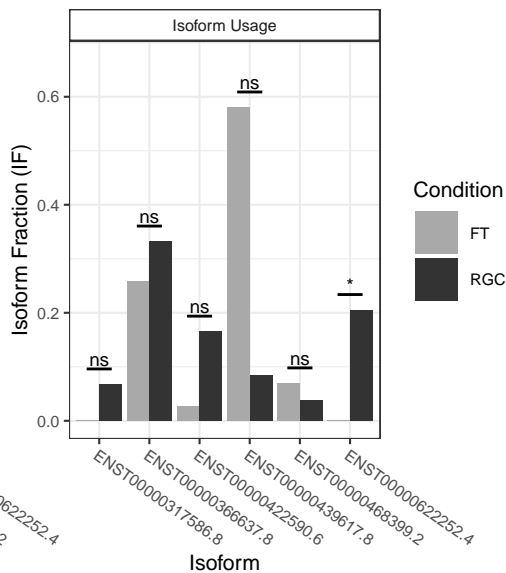
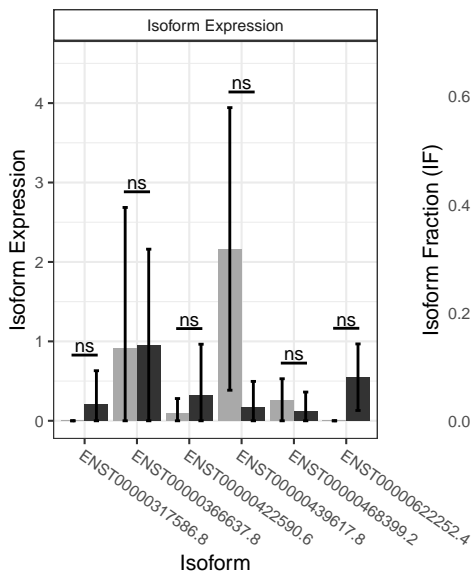
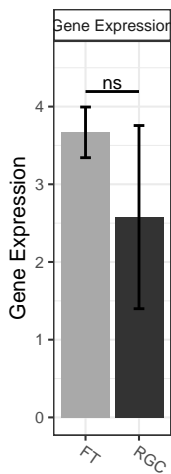
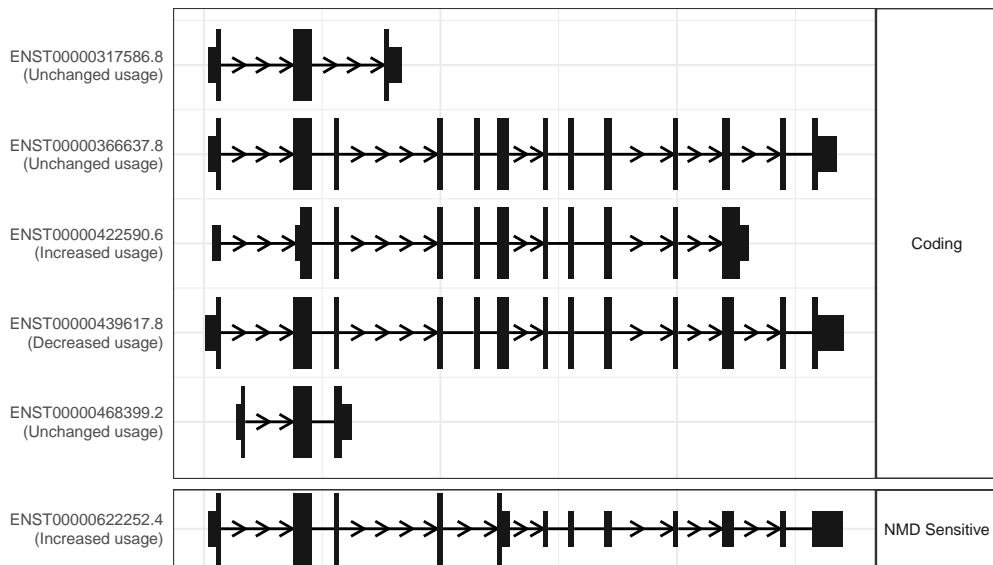


Helicase_C

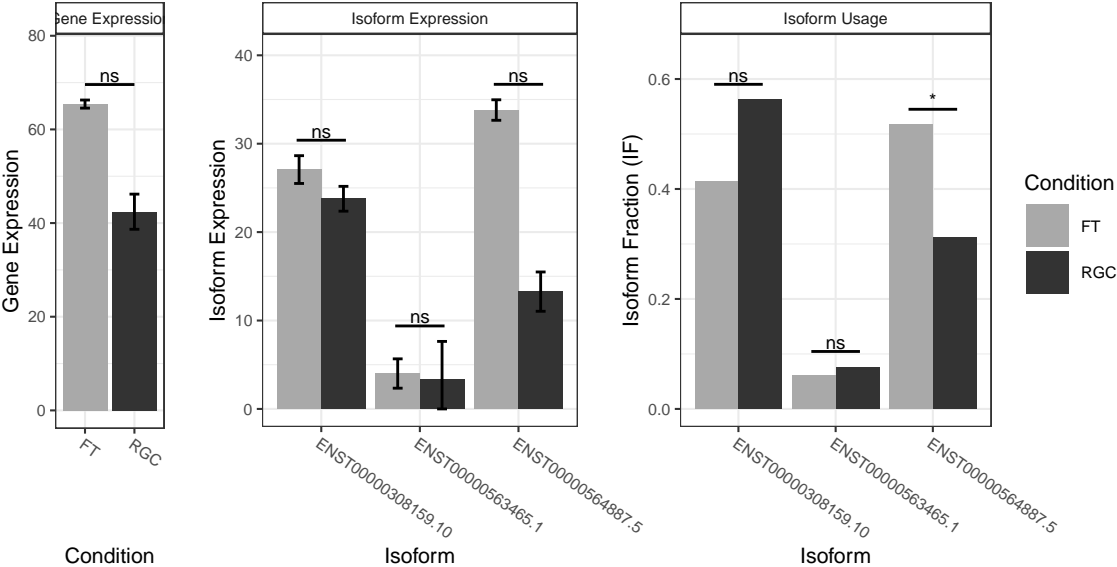
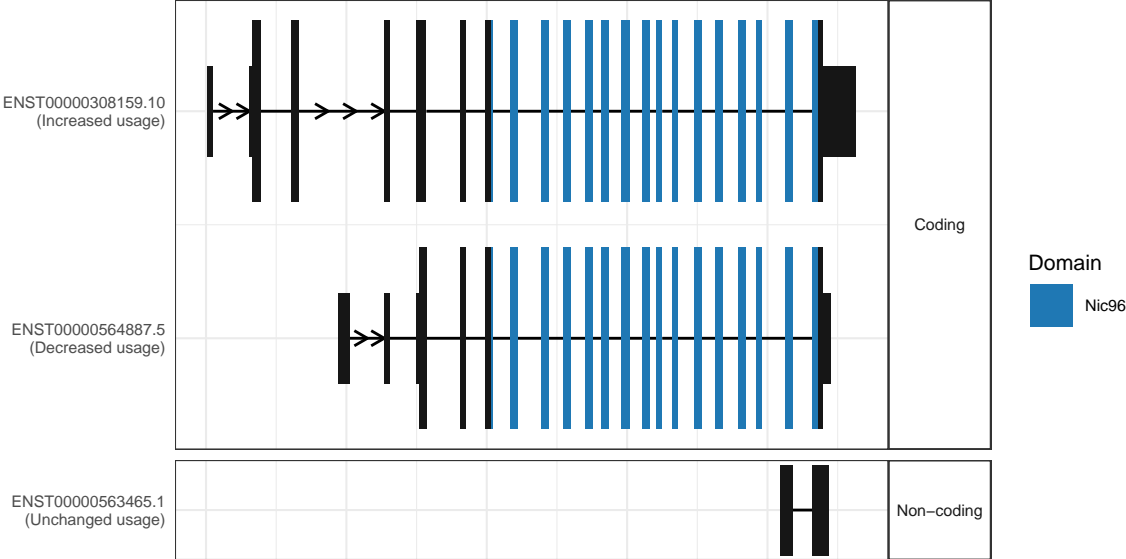


RecQ_Zn_bind

The isoform switch in DISC1 (FT vs RGC)



The isoform switch in NUP93 (FT vs RGC)

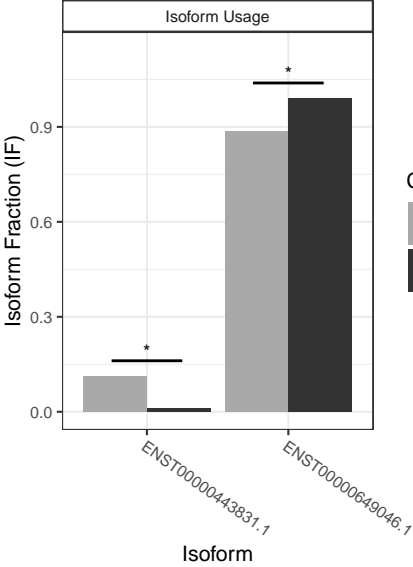
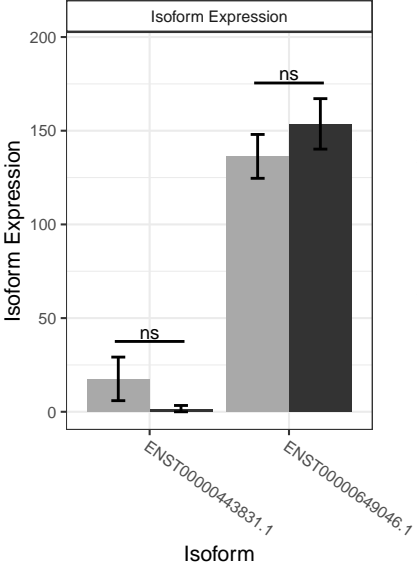
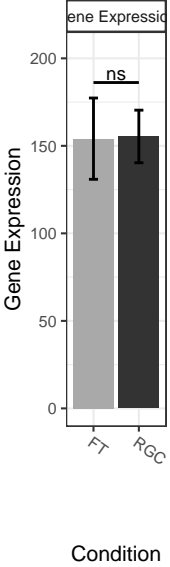
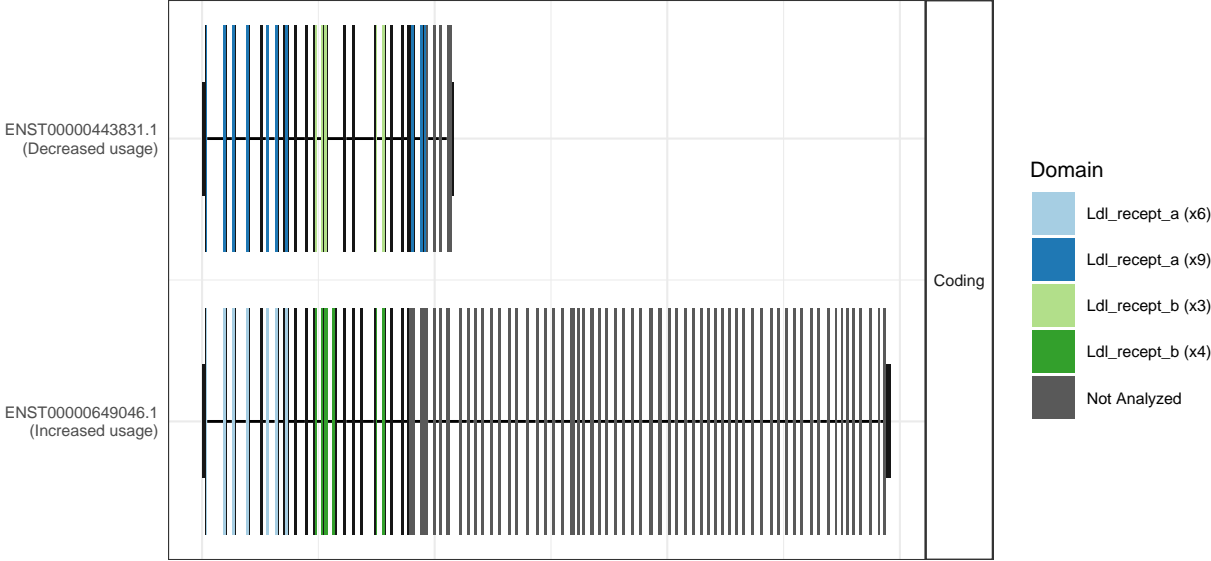


Domain



Nic96

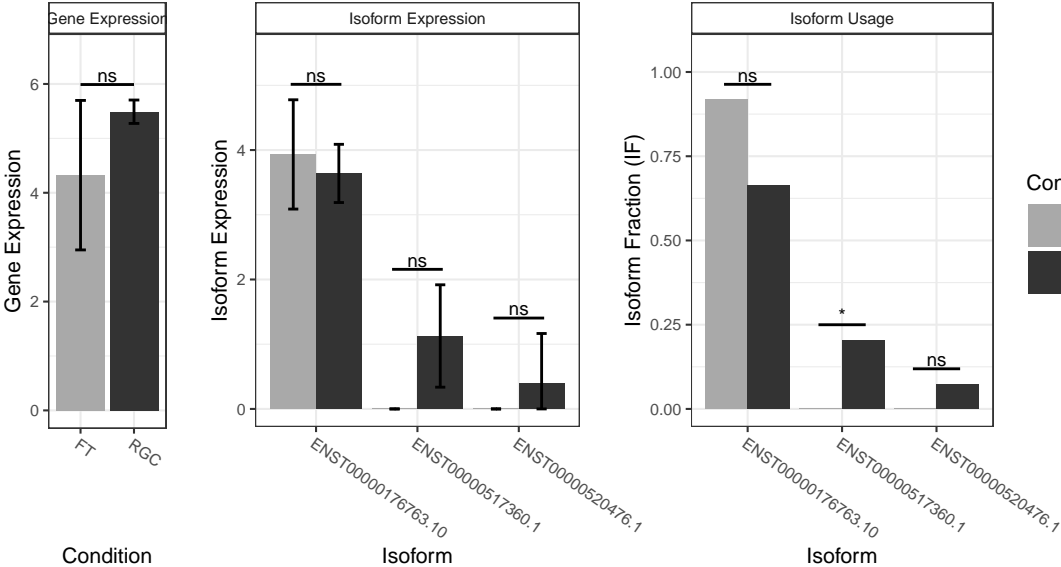
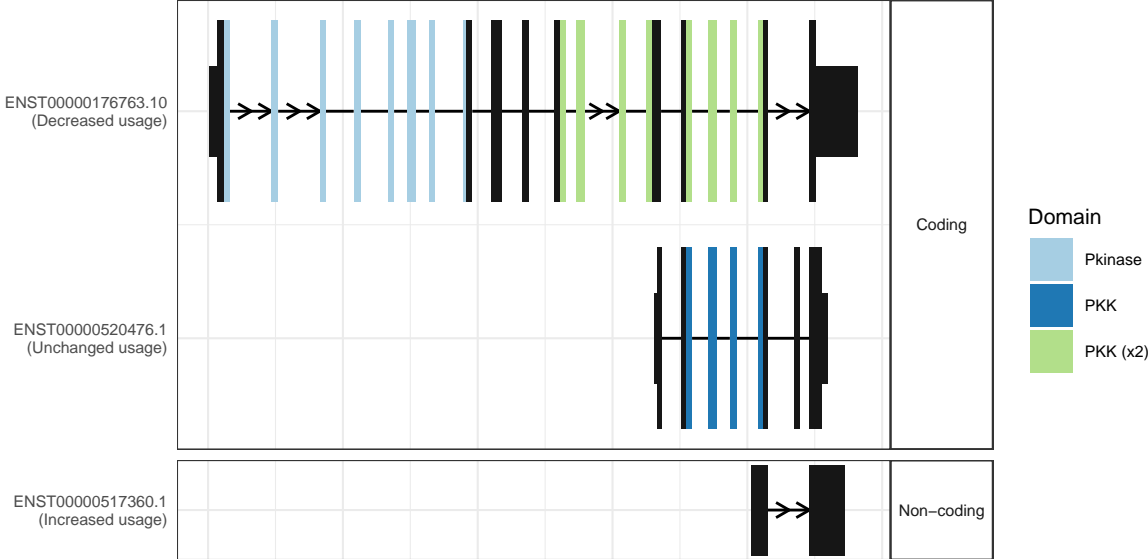
The isoform switch in LRP2 (FT vs RGC)



Domain



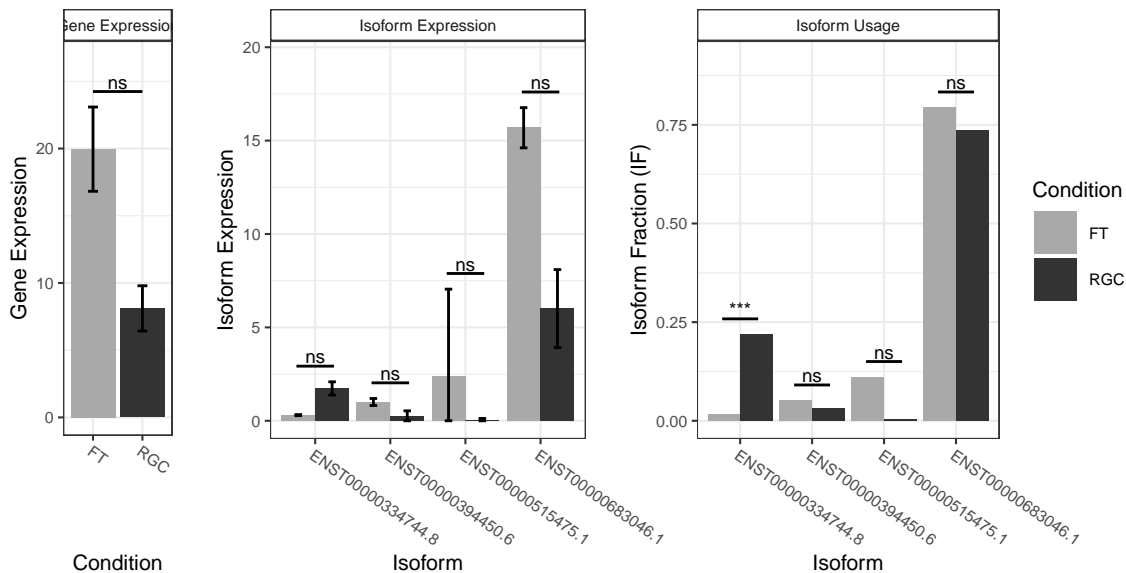
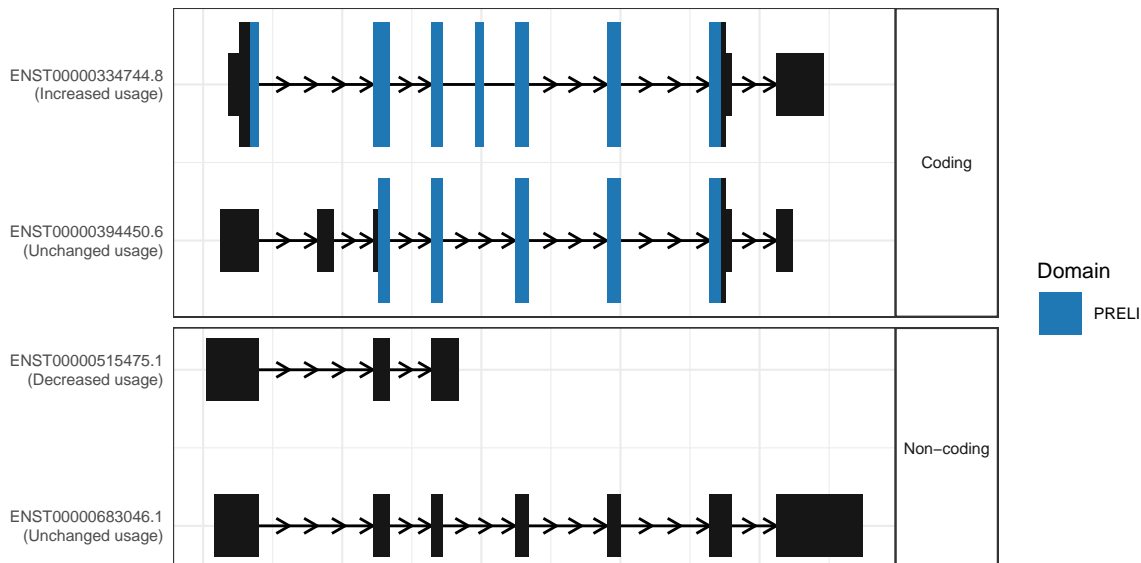
The isoform switch in STK10 (FT vs RGC)



Domain

	Pkinase
	PKK
	PKK (x2)








The isoform switch in PRELID2 (FT vs RGC)



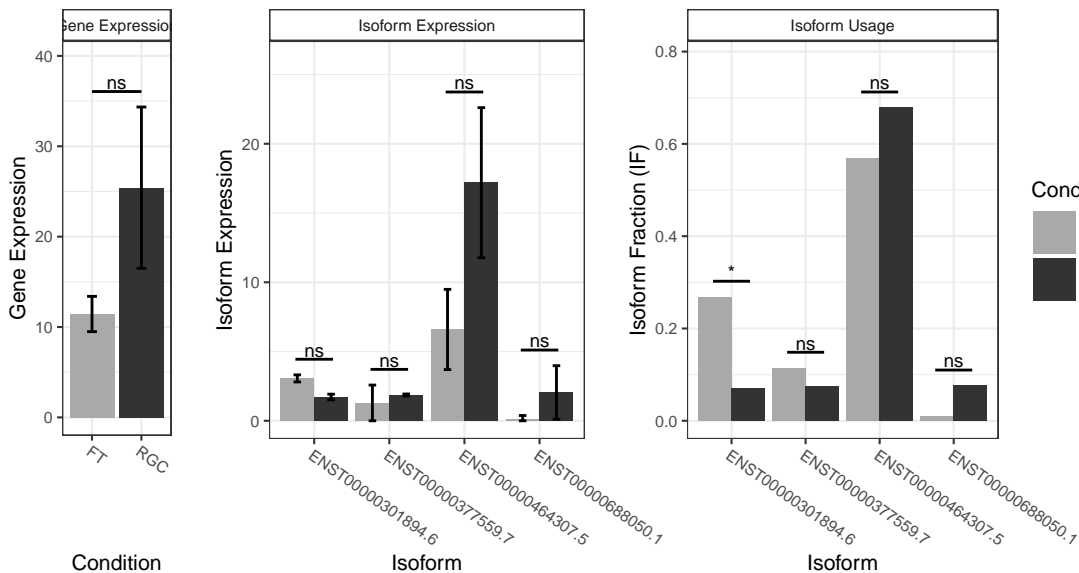
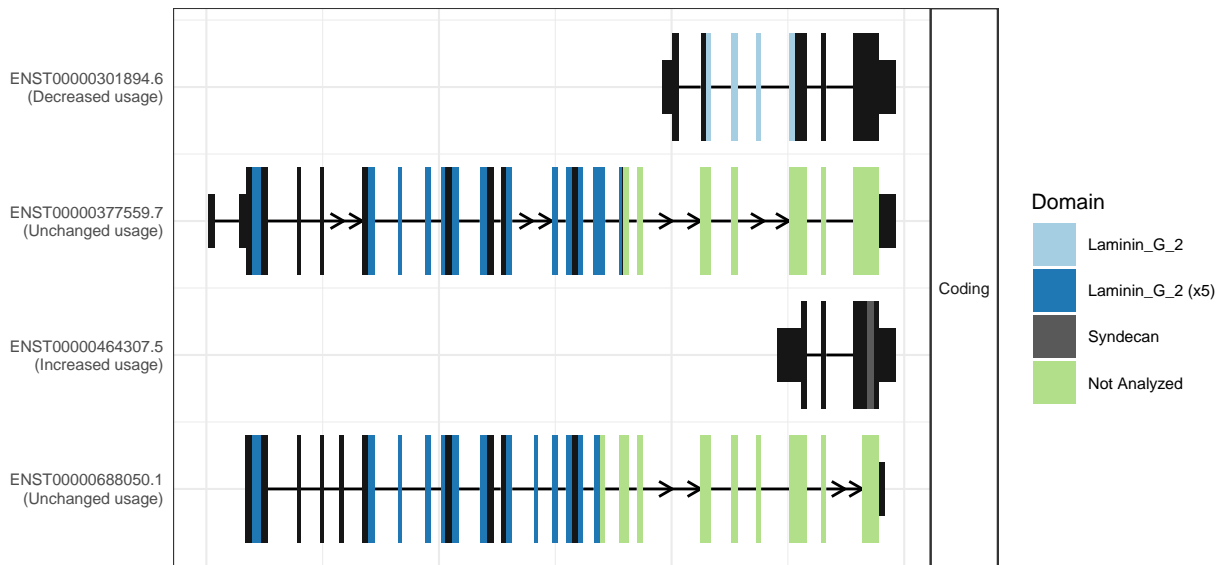
Domain

 PRELI





Domain

	betaPIX_CC
	CH
	PH
	RhoGEF
	RhoGEF67_u1
	RhoGEF67_u2
	SH3_2

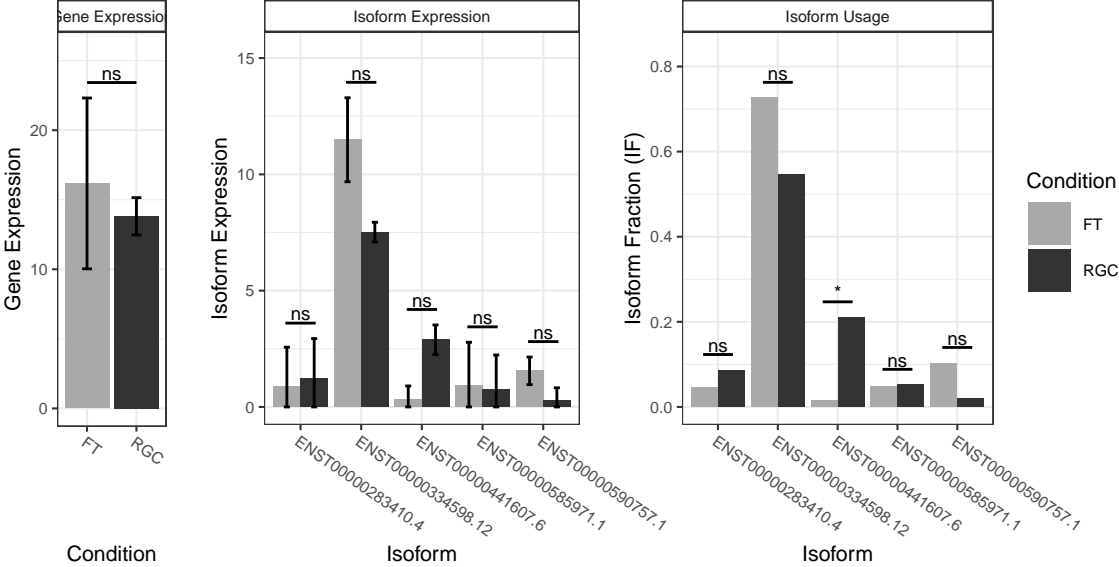
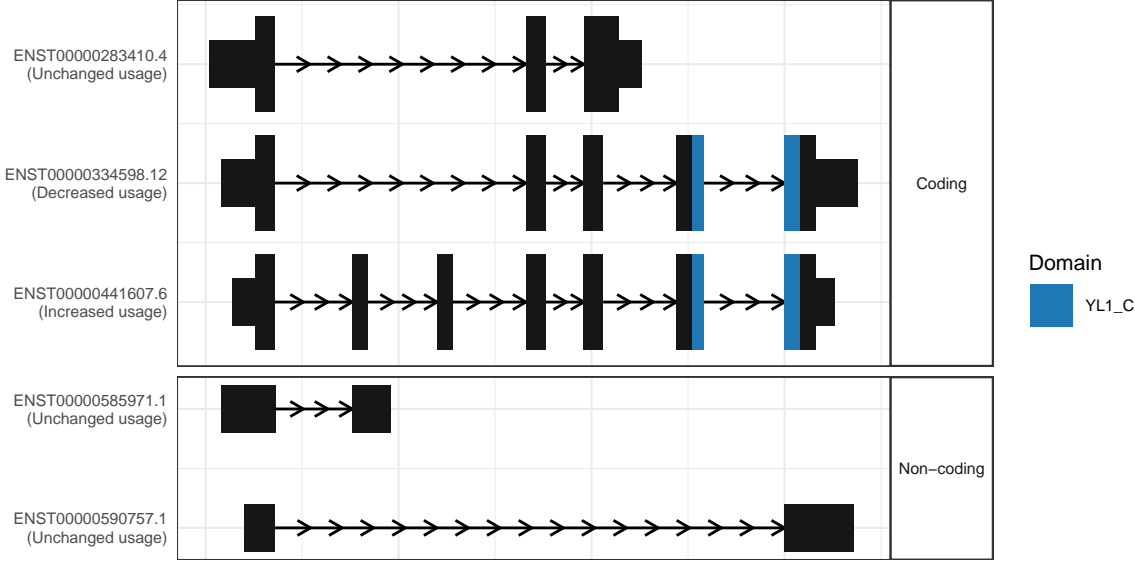
The isoform switch in NRXN2 (FT vs RGC)



Domain

	Laminin_G_2
	Laminin_G_2 (x5)
	Syndecan
	Not Analyzed

The isoform switch in INO80C (FT vs RGC)

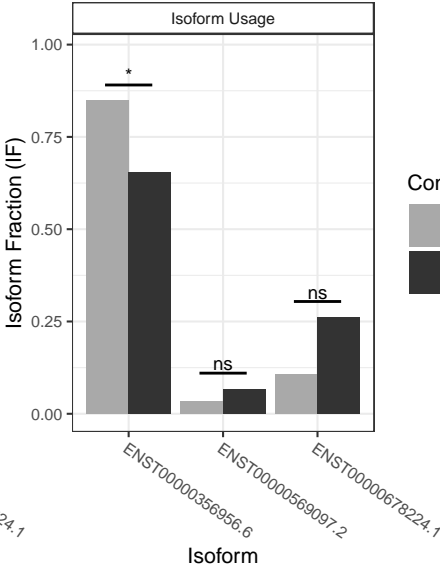
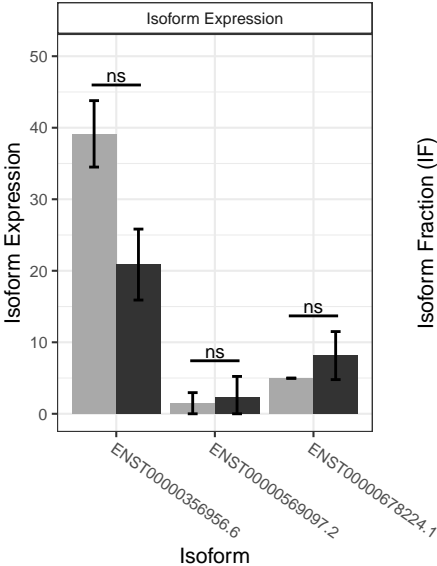
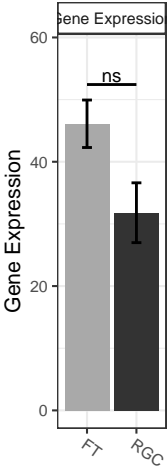
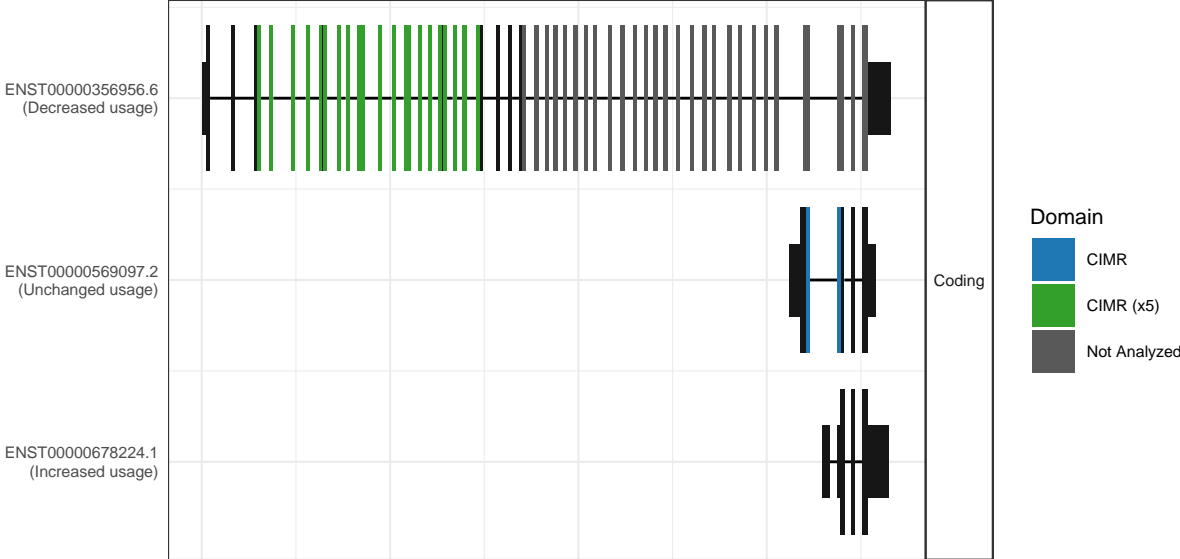


Domain

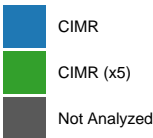


YL1_C

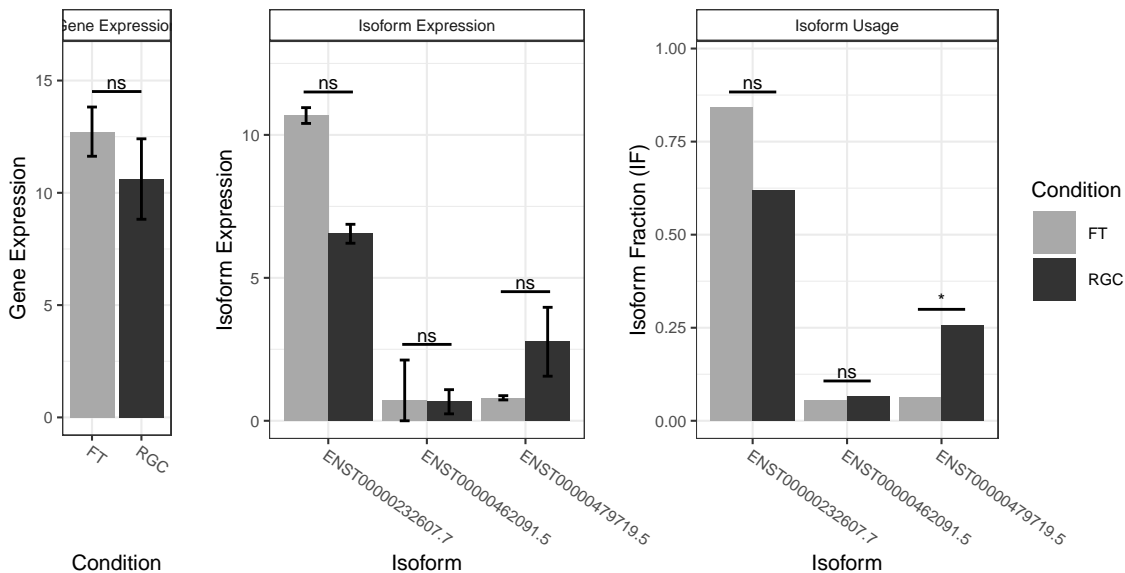
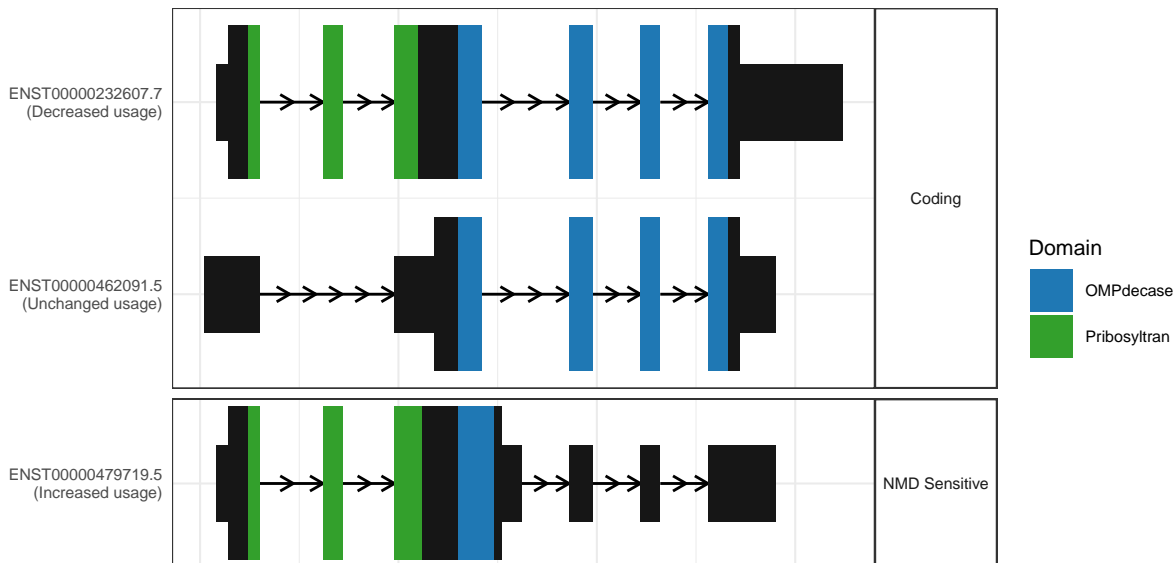
The isoform switch in IGF2R (FT vs RGC)



Domain



The isoform switch in UMPS (FT vs RGC)



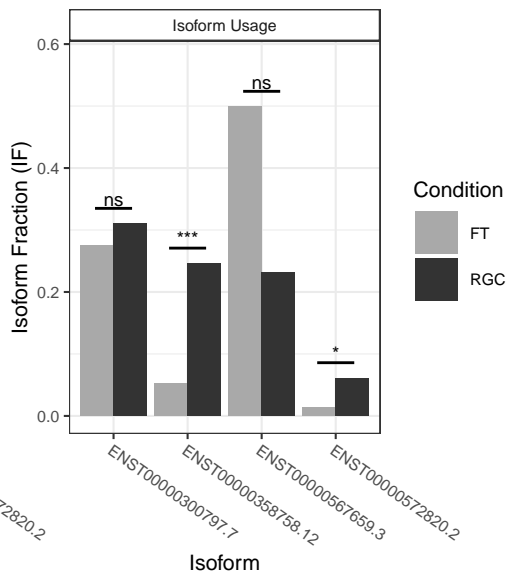
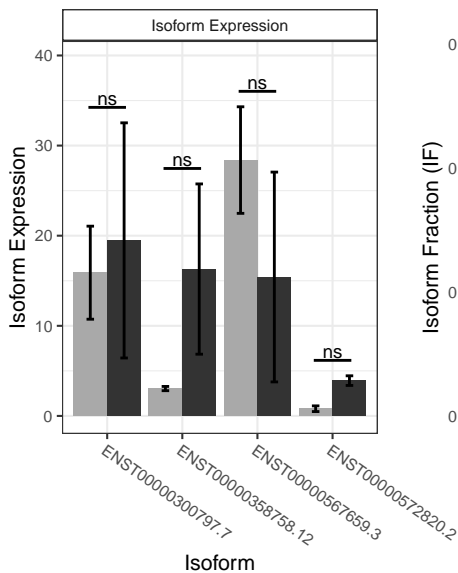
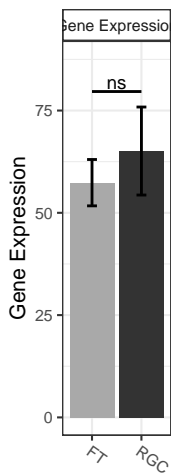
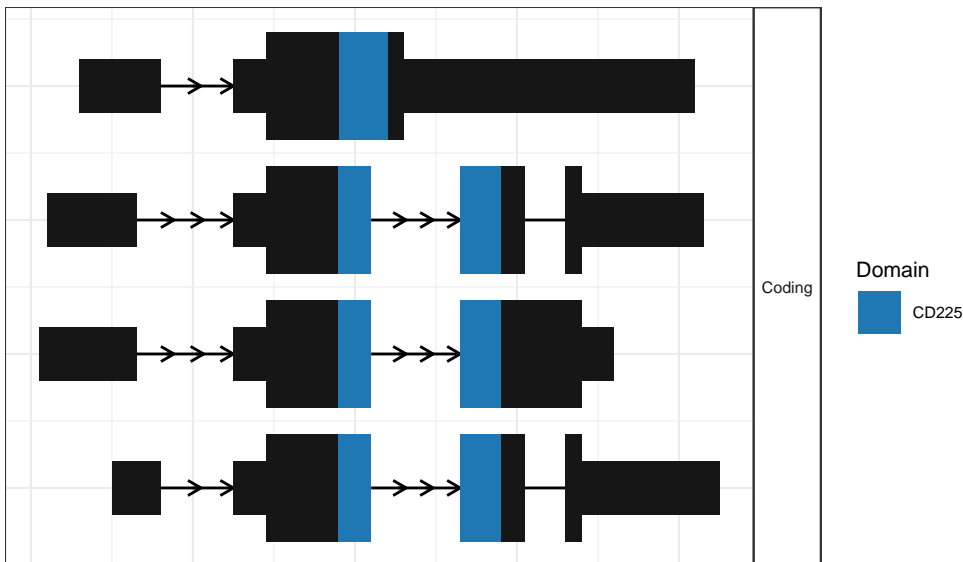
Domain



OMPdecase

Pribosyltran

The isoform switch in PRRT2 (FT vs RGC)



Condition

Isoform

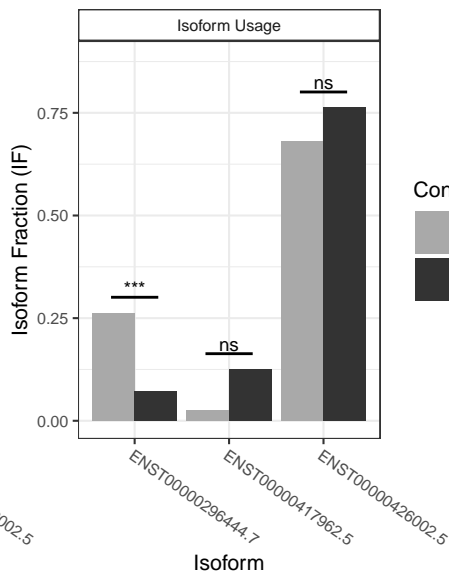
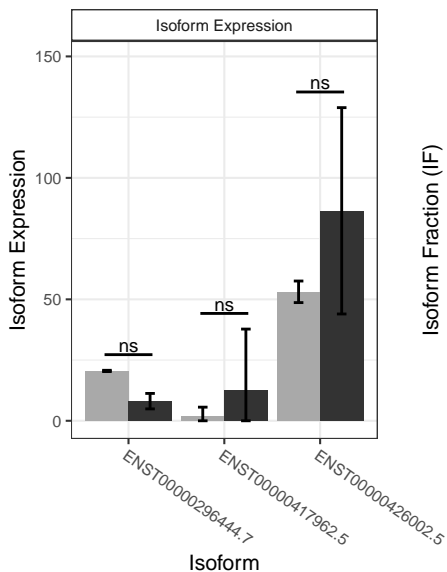
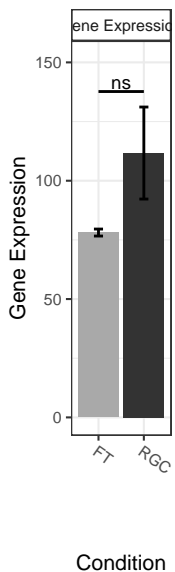
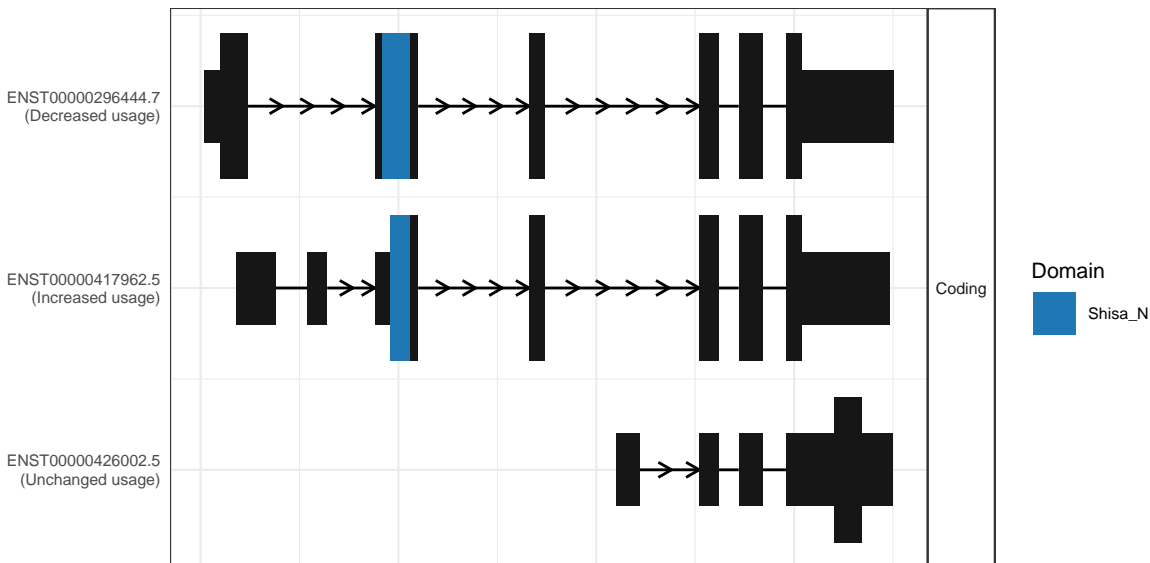
Isoform

Domain



CD225

The isoform switch in SHISA5 (FT vs RGC)

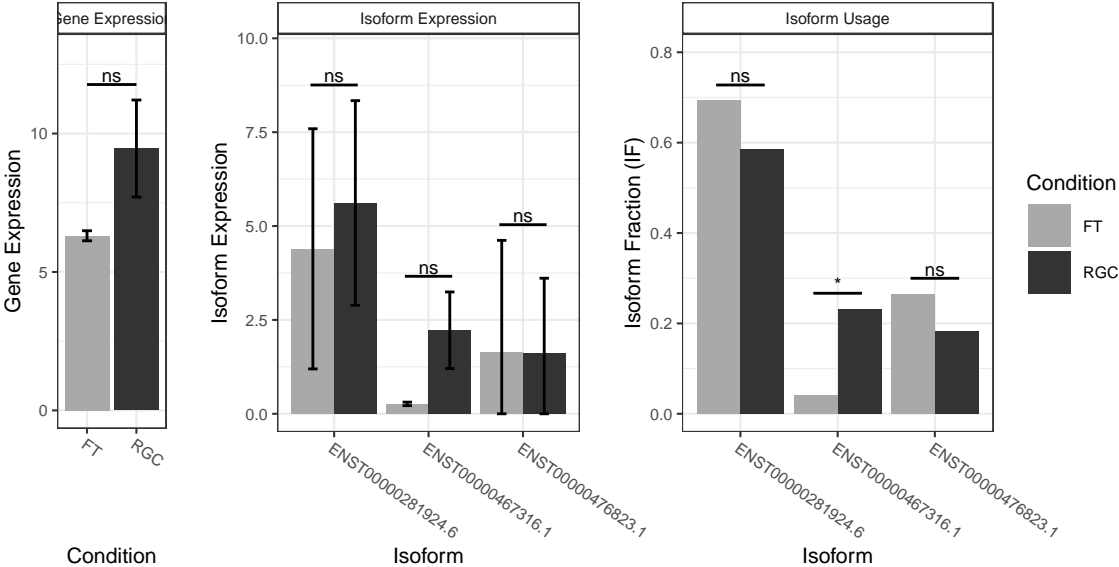
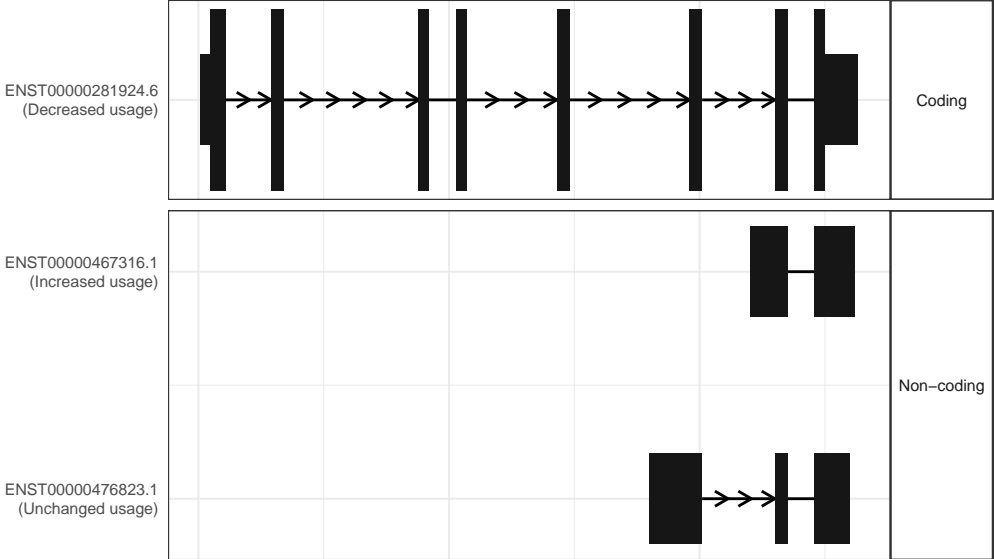


Domain

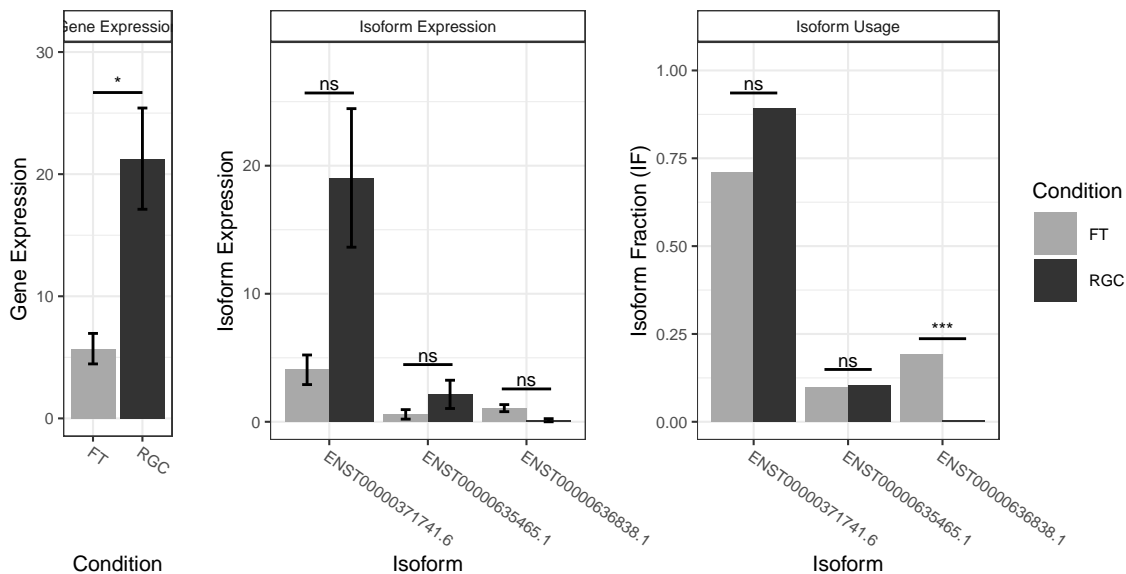
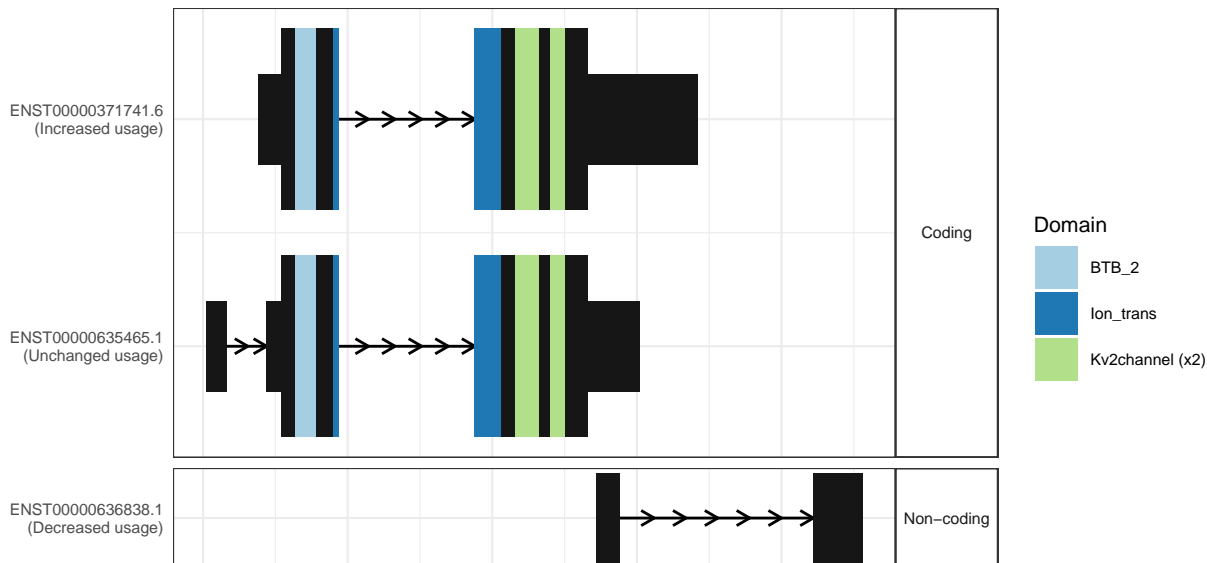


Shisa_N

The isoform switch in TMEM163 (FT vs RGC)



The isoform switch in KCNB1 (FT vs RGC)



Domain



BTB_2

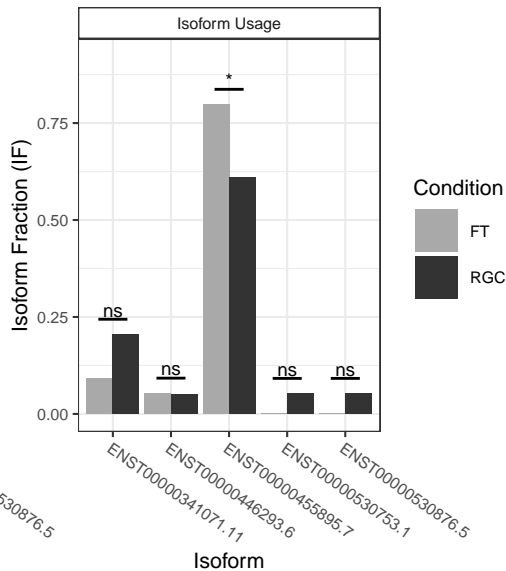
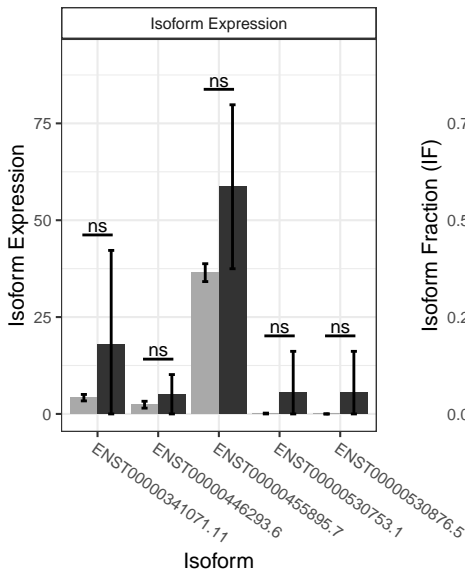
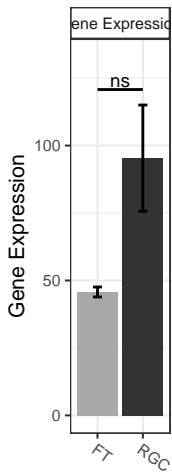
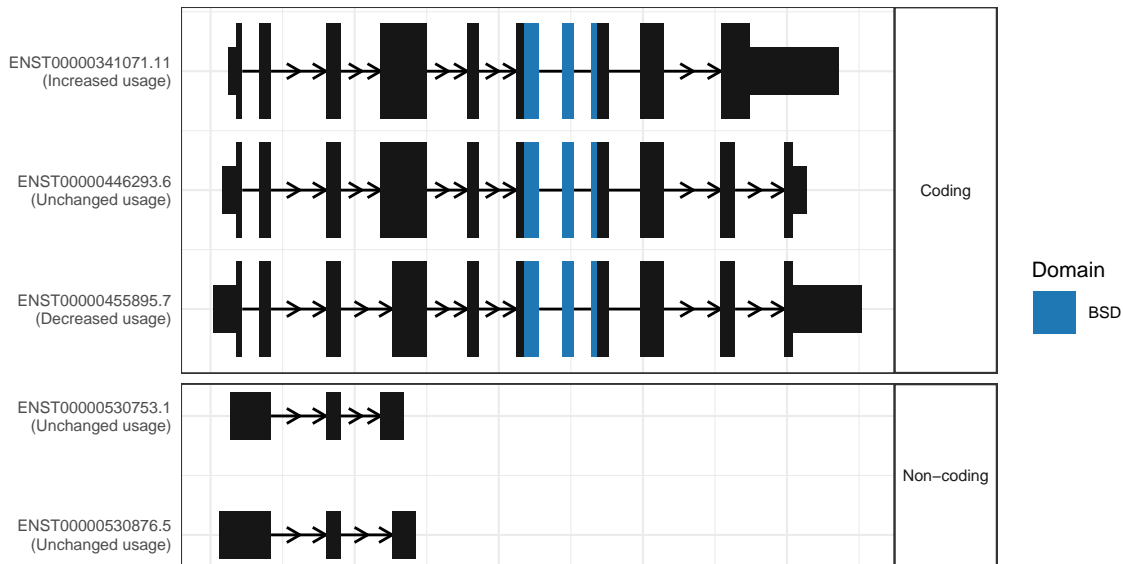


lon_trans



Kv2channel (x2)

The isoform switch in BSDC1 (FT vs RGC)

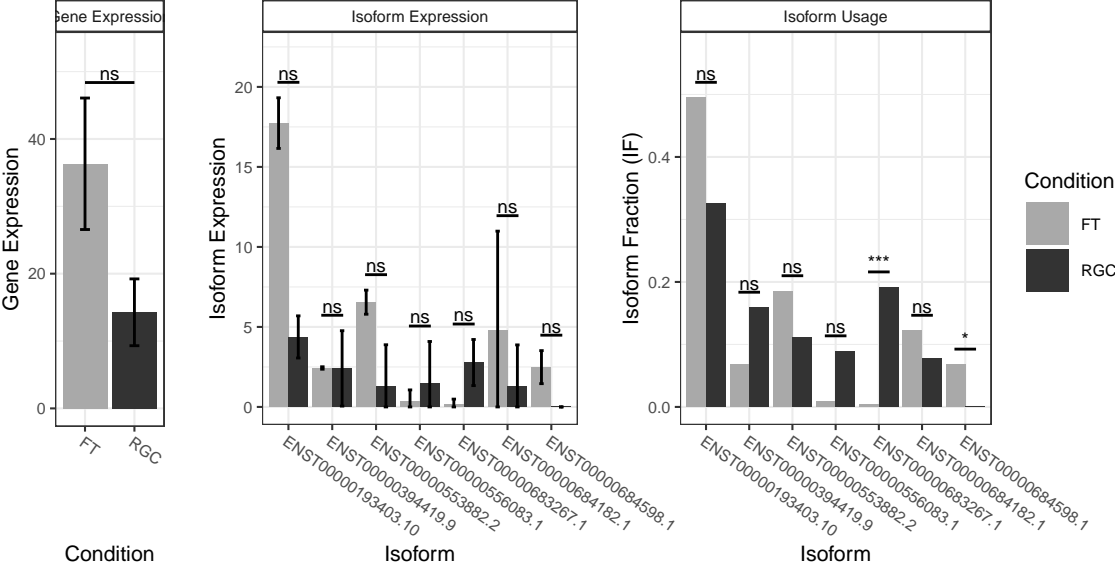
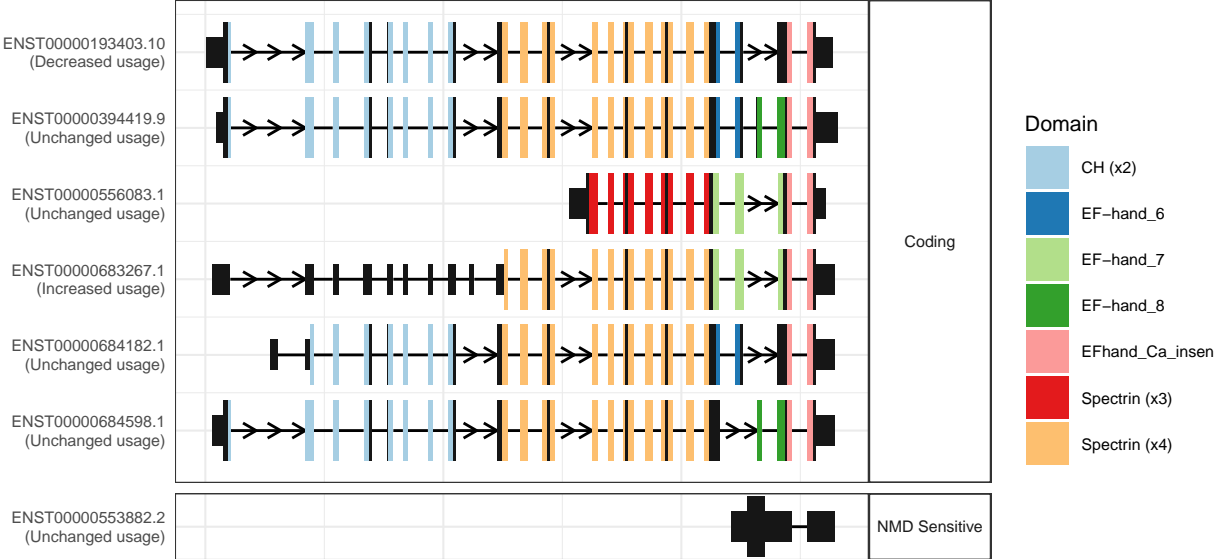


Domain










BSD

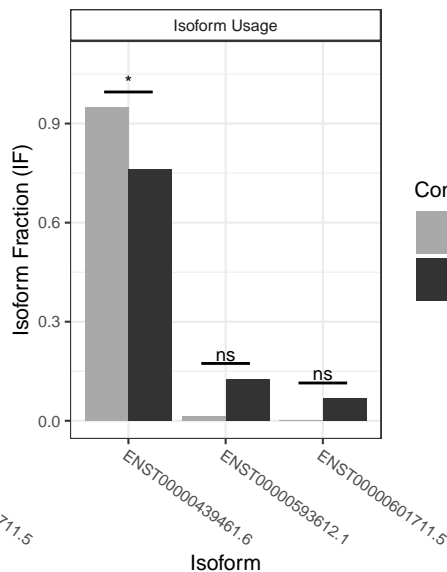
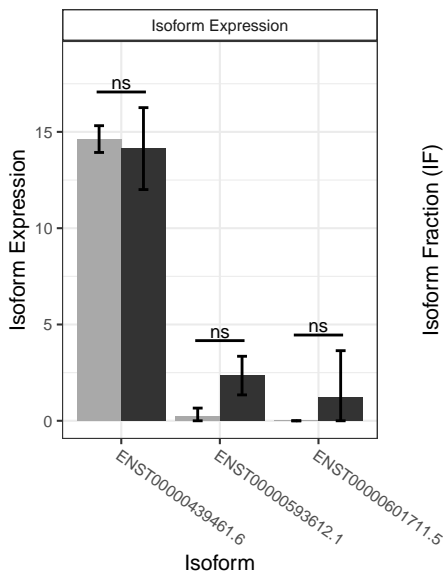
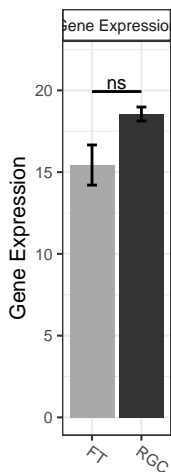
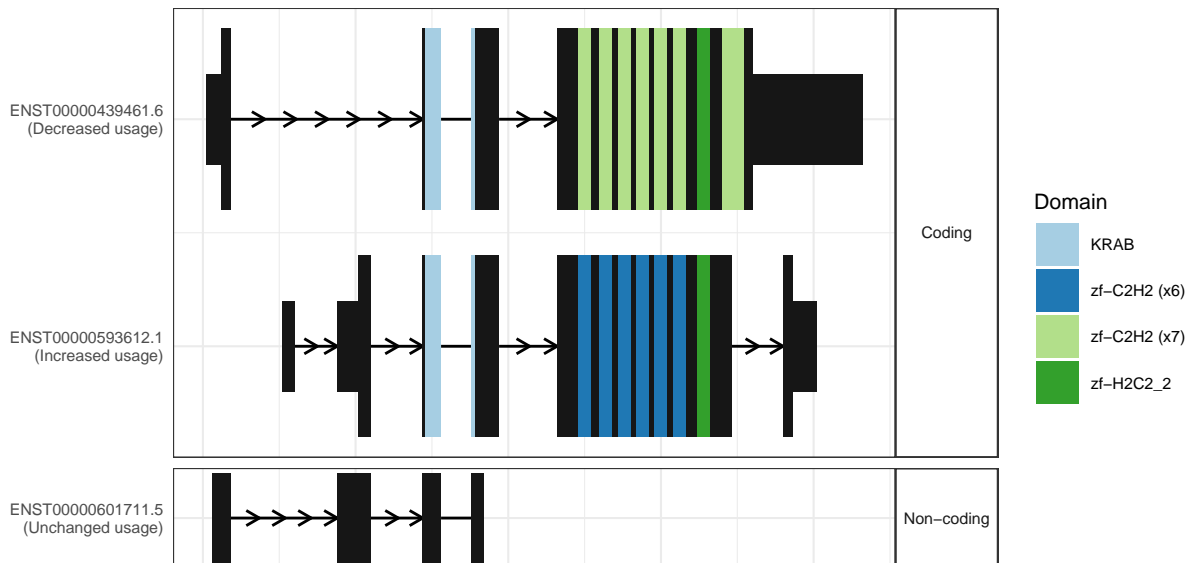
The isoform switch in ACTN1 (FT vs RGC)



Domain

	CH (x2)
	EF-hand_6
	EF-hand_7
	EF-hand_8
	EFhand_Ca_insen
	Spectrin (x3)
	Spectrin (x4)

The isoform switch in ZNF766 (FT vs RGC)



Condition

Isoform

Isoform

Domain



KRAB



zf-C2H2 (x6)

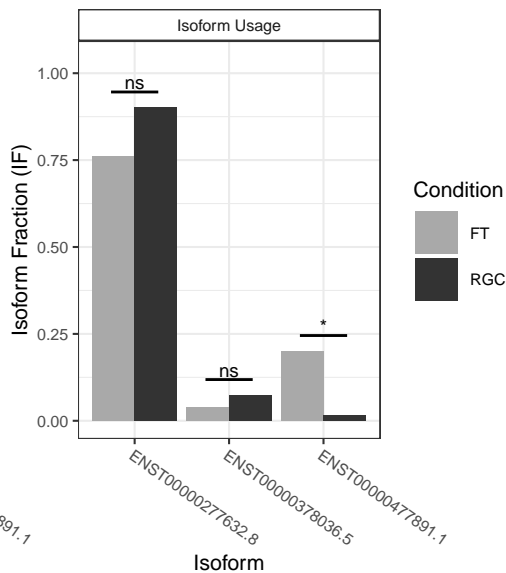
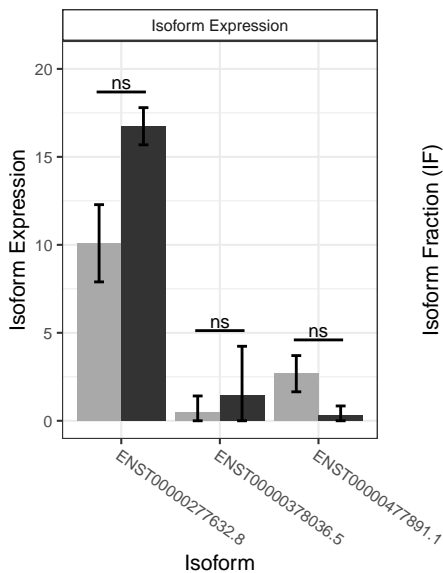
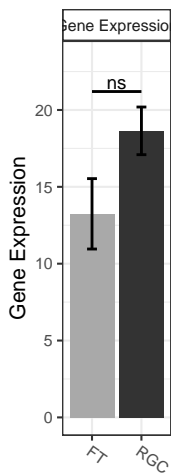
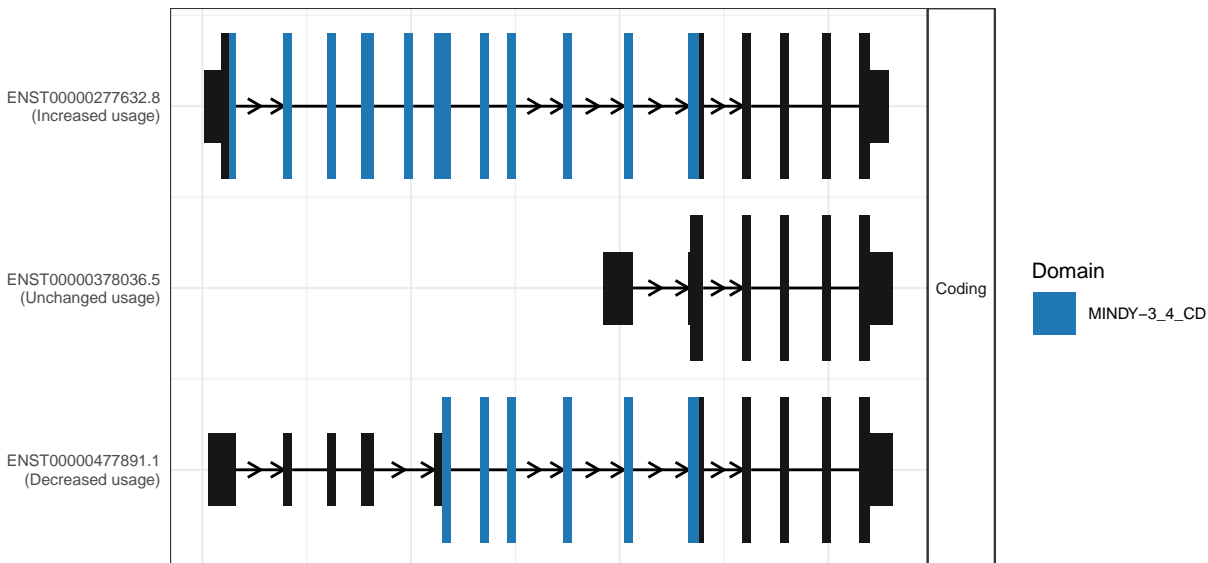


zf-C2H2 (x7)



zf-H2C2_2

The isoform switch in MINDY3 (FT vs RGC)

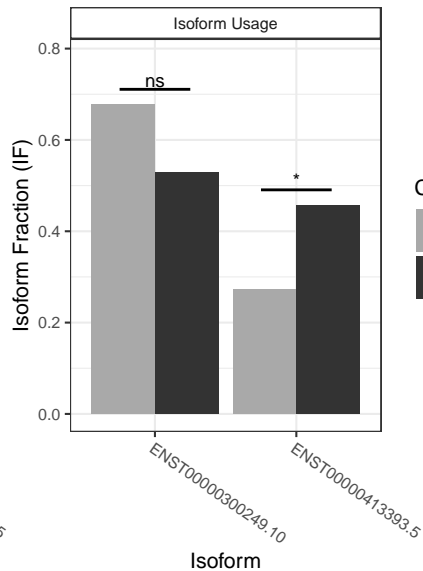
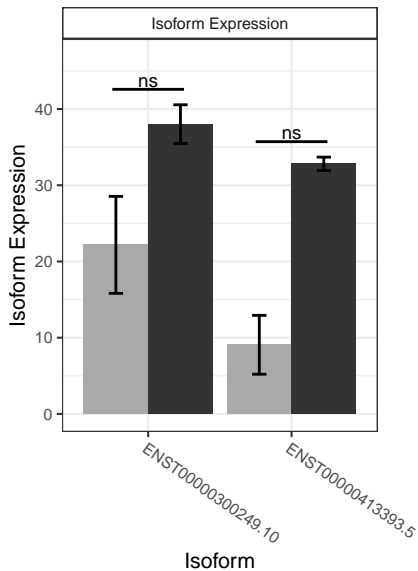
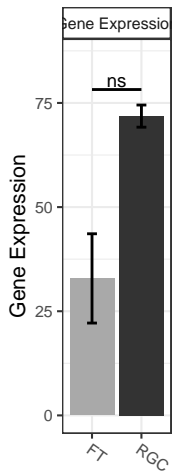
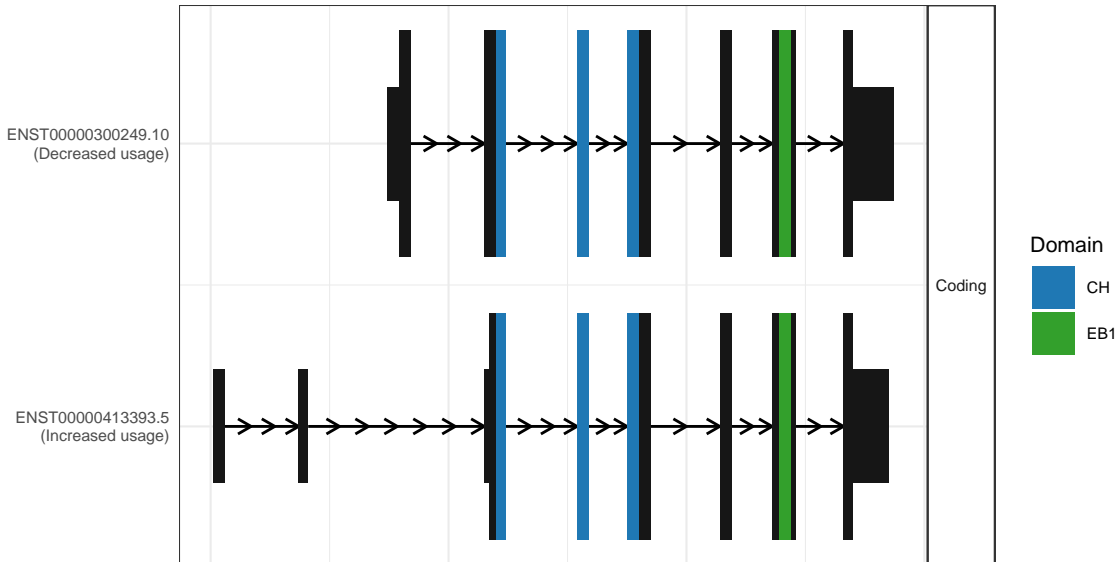


Domain



MINDY-3_4_CD

The isoform switch in MAPRE2 (FT vs RGC)



Domain

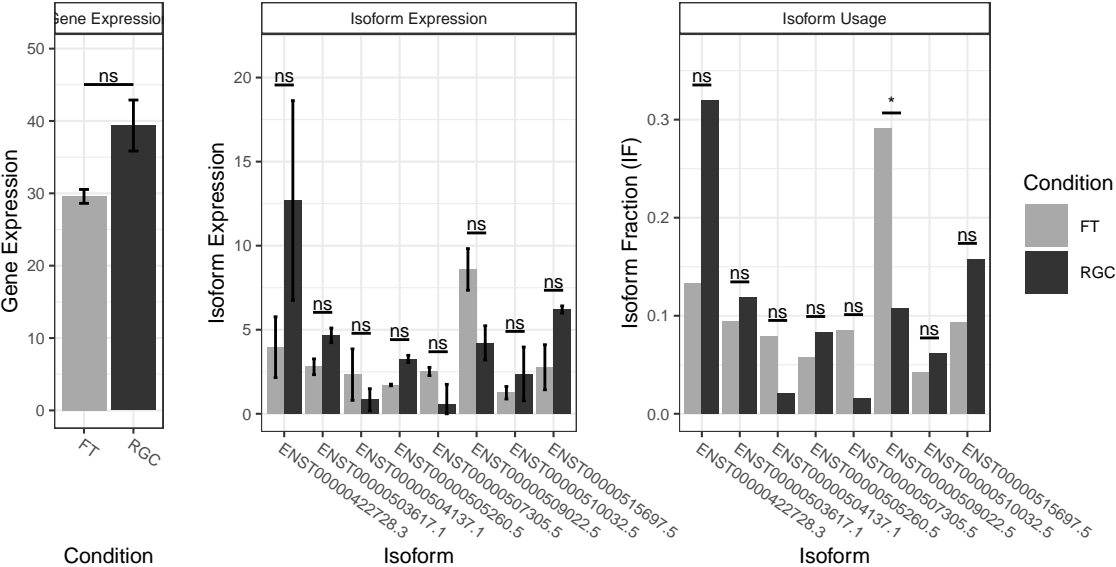
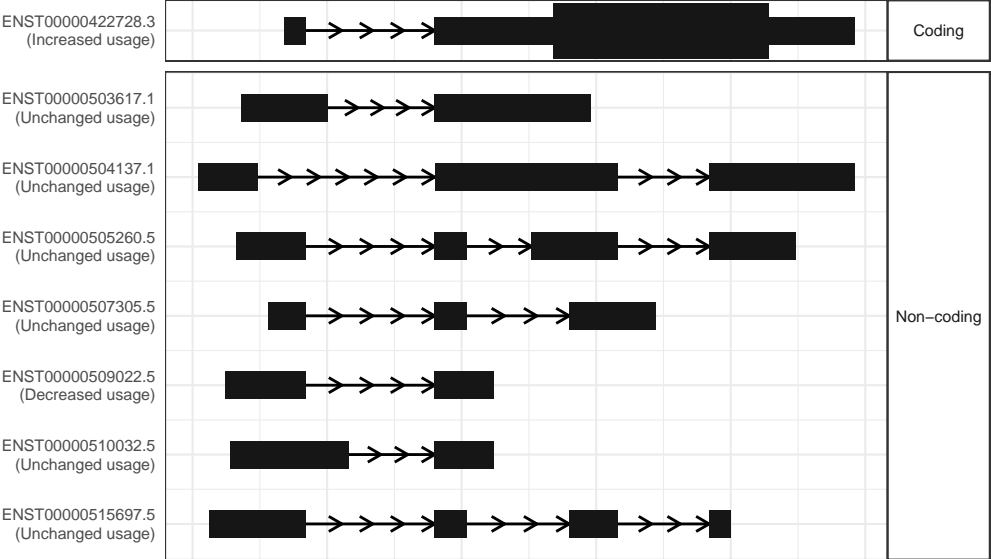


CH

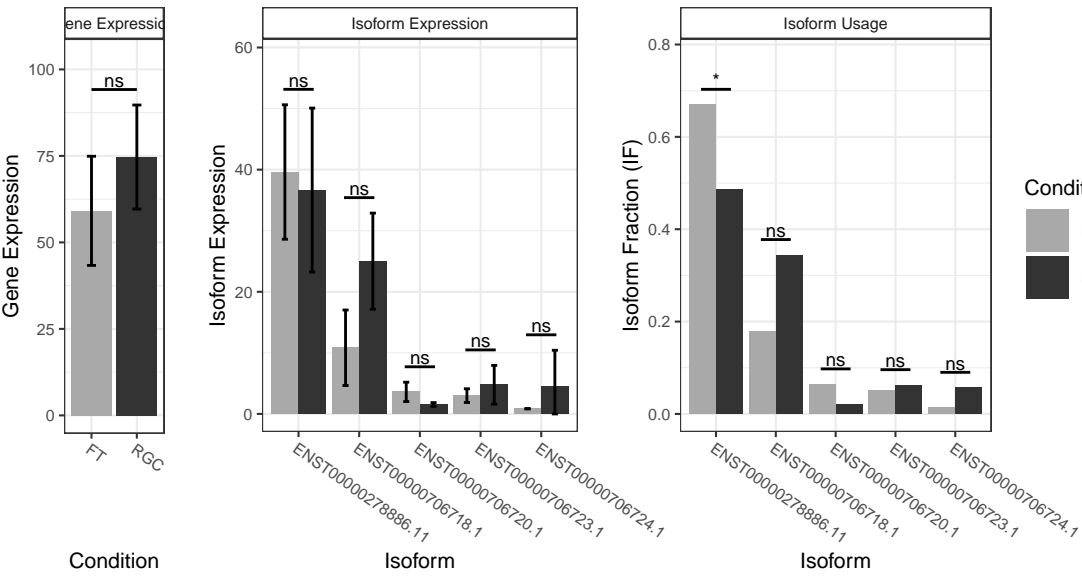
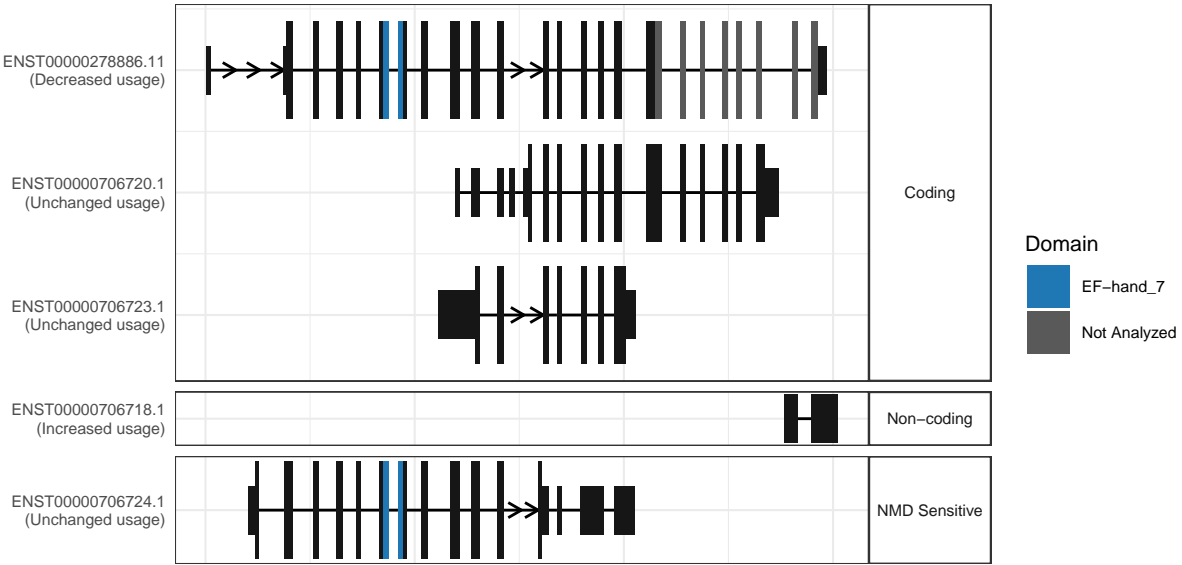


EB1

The isoform switch in FAM200B (FT vs RGC)



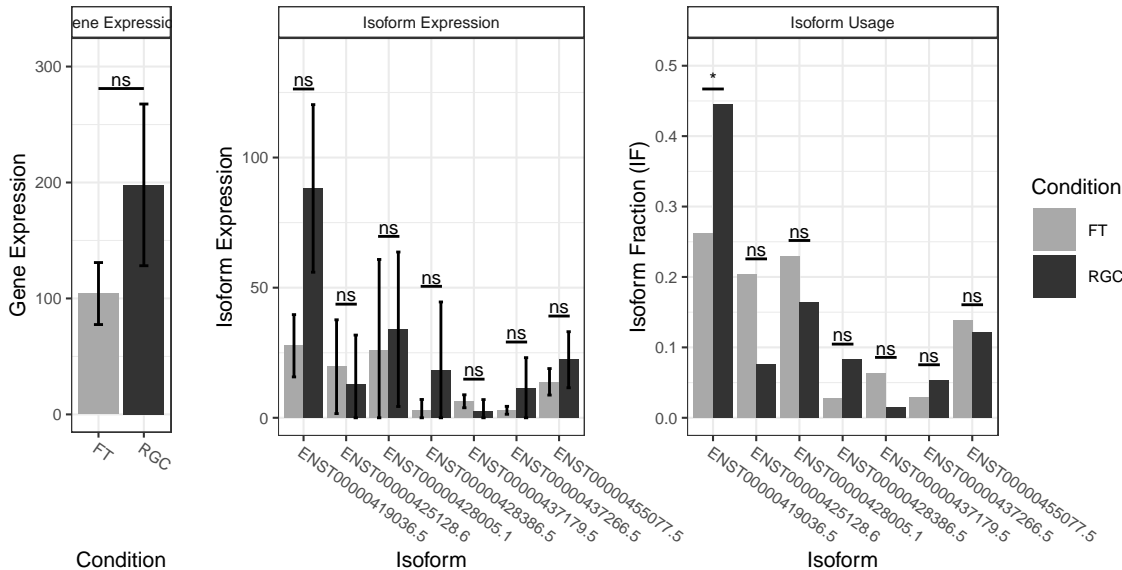
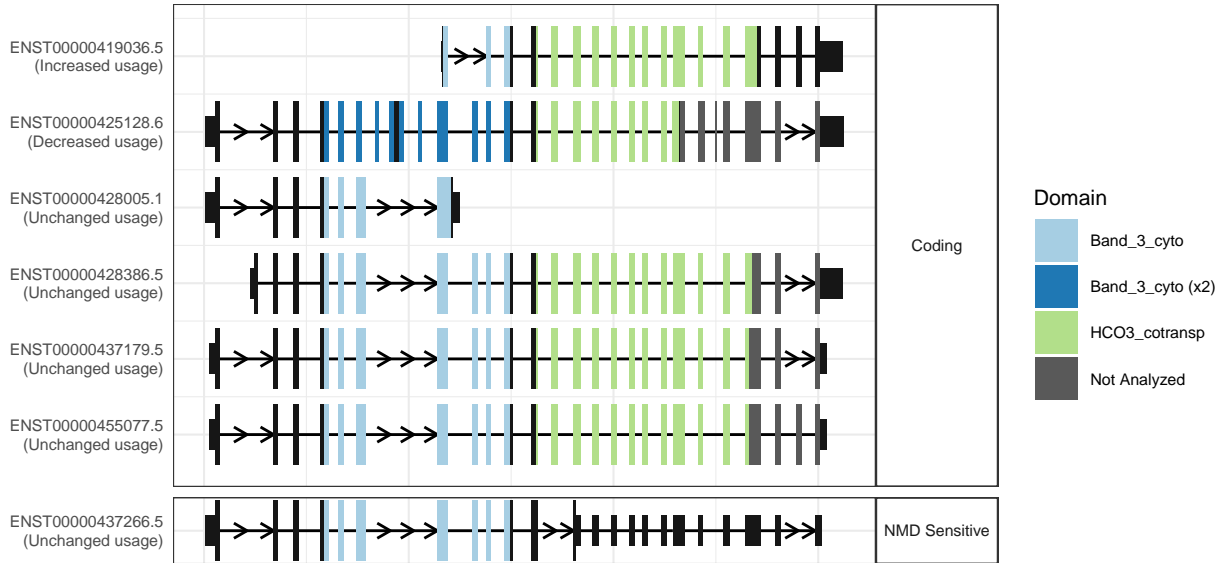
The isoform switch in NINL (FT vs RGC)



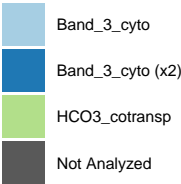
Domain

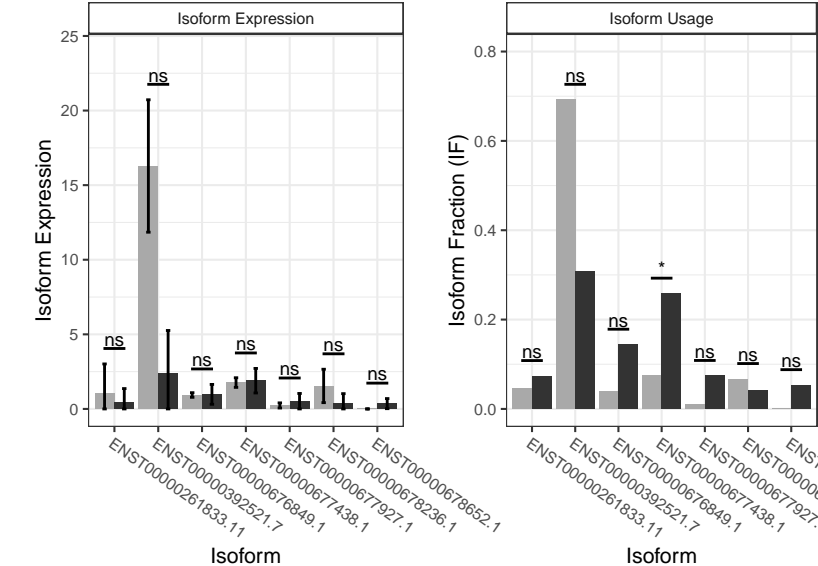


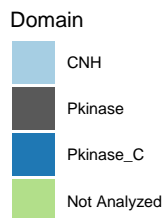
The isoform switch in SLC4A7 (FT vs RGC)



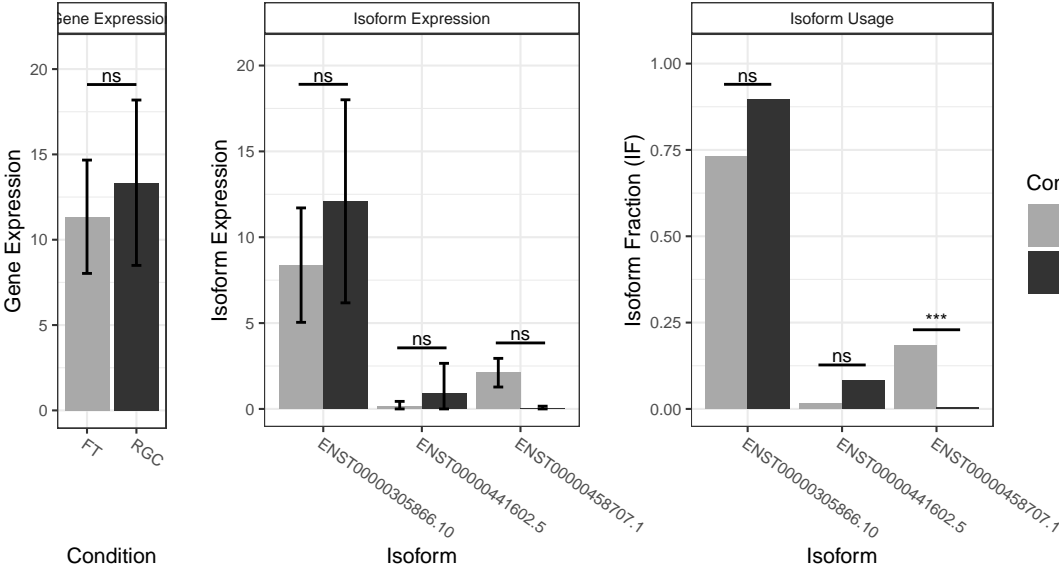
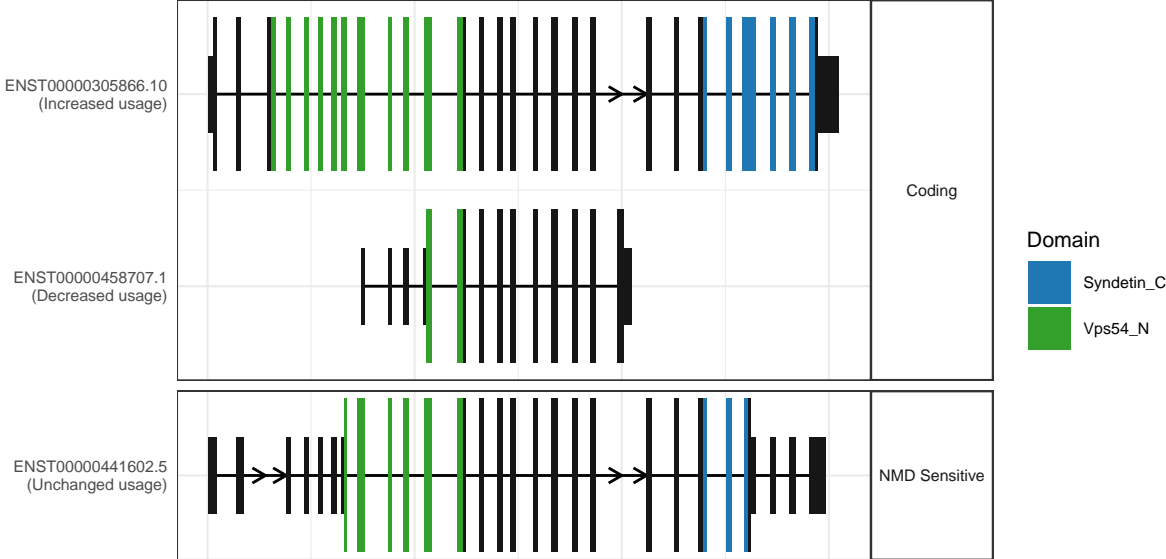
Domain







The isoform switch in VPS50 (FT vs RGC)



Domain

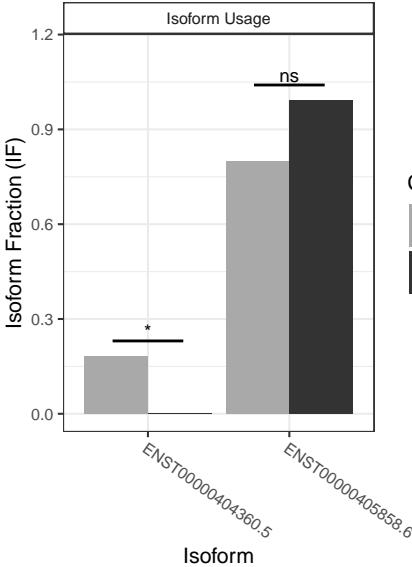
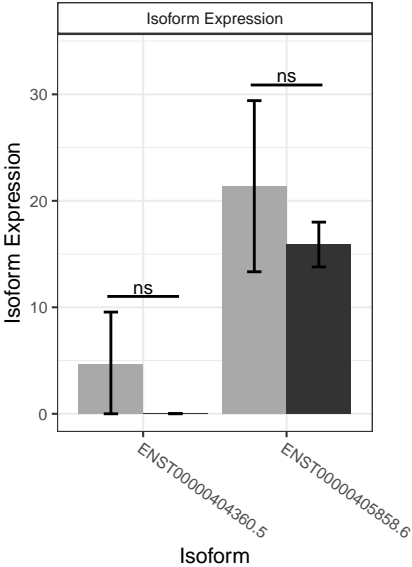
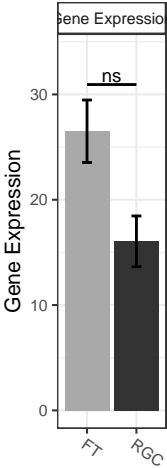
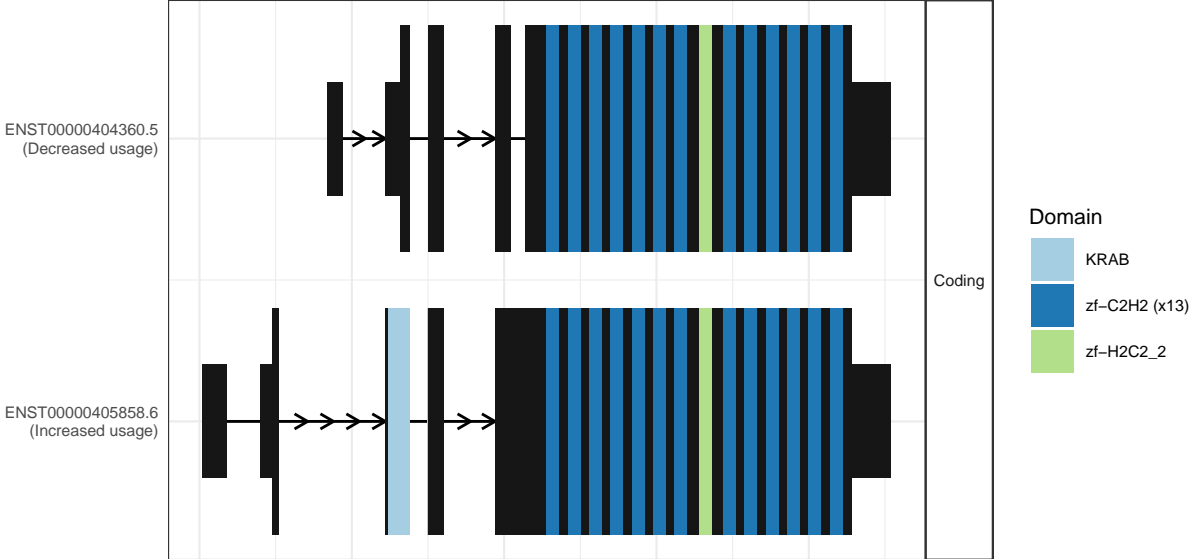


Syndetin_C






Vps54_N

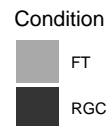
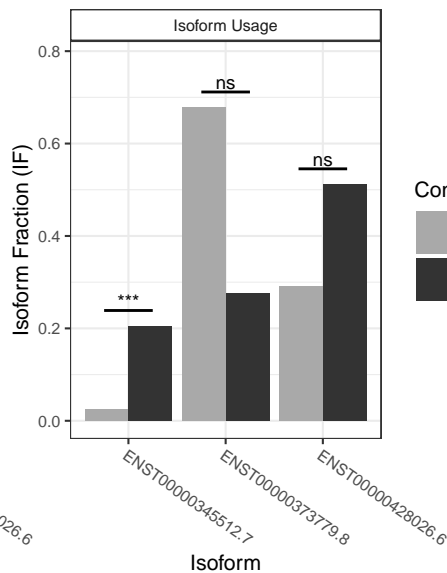
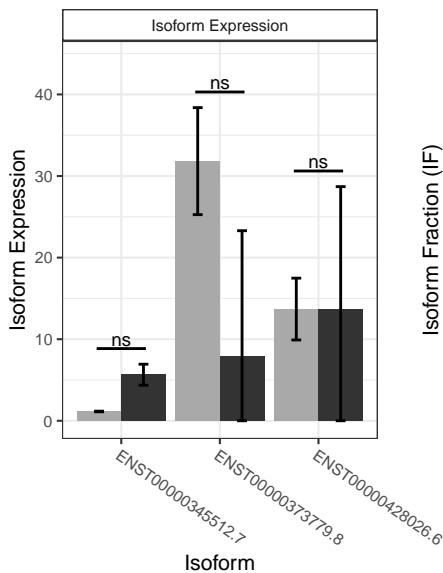
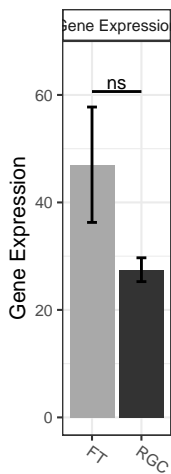
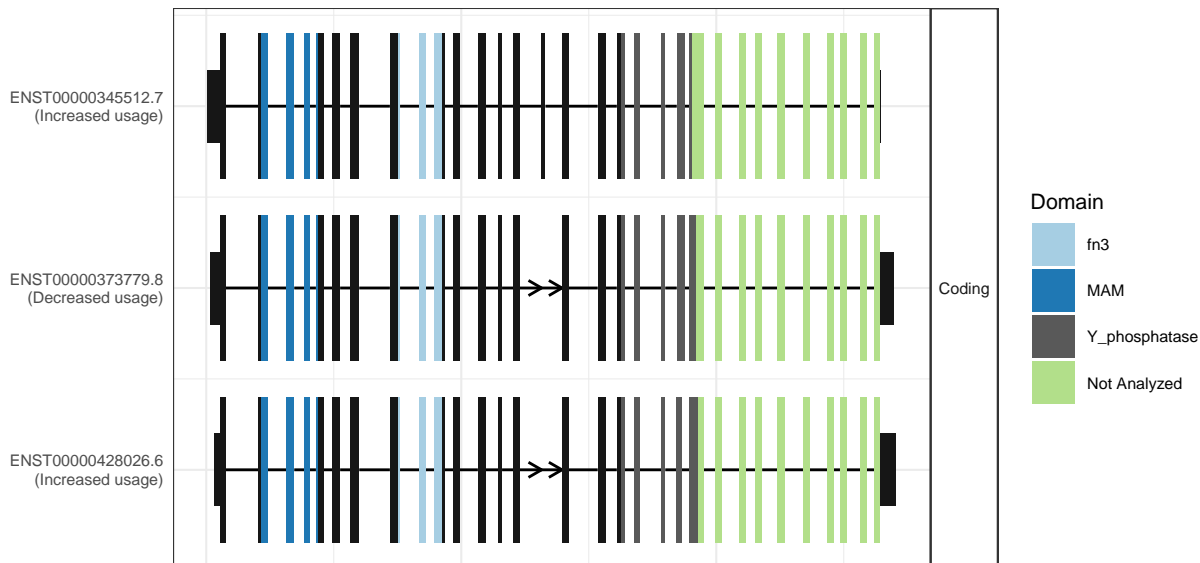
The isoform switch in ZNF12 (FT vs RGC)



Domain

	KRAB
	zf-C2H2 (x13)
	zf-H2C2_2

The isoform switch in PTPRU (FT vs RGC)



Domain



fn3



MAM

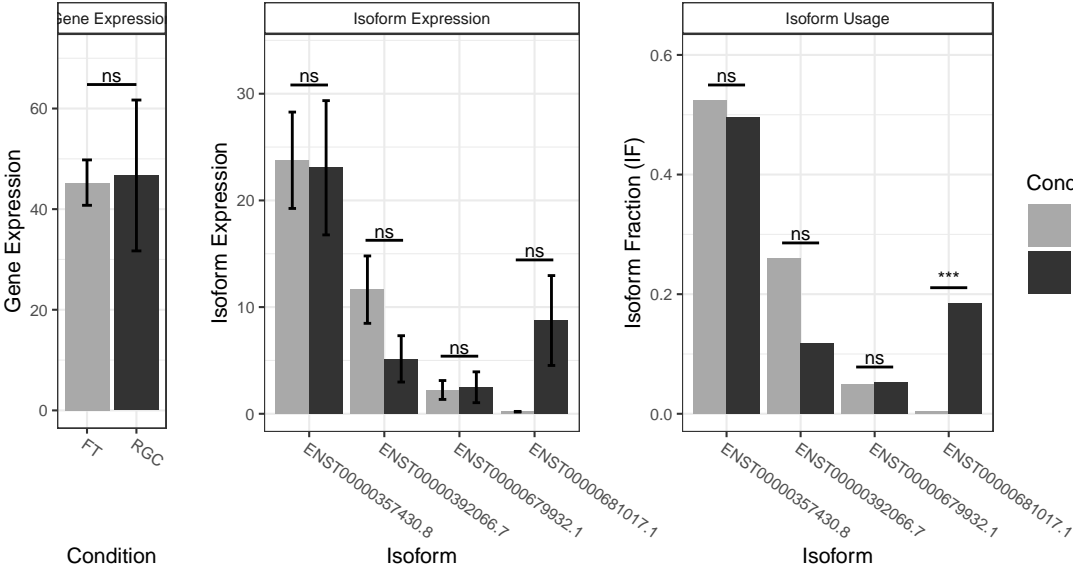
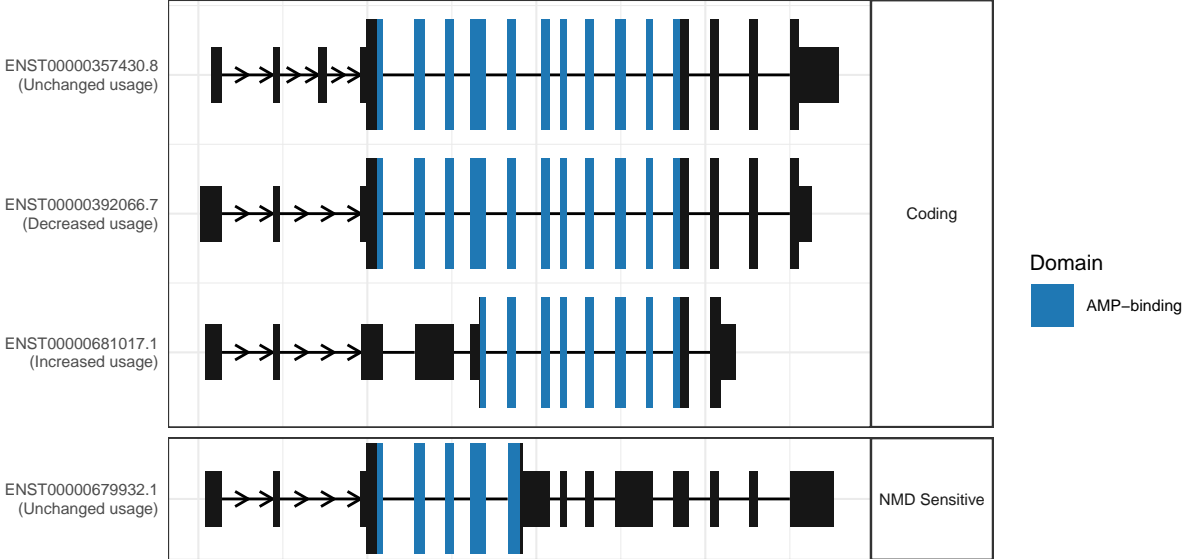


Y_phosphatase



Not Analyzed

The isoform switch in ACSL3 (FT vs RGC)

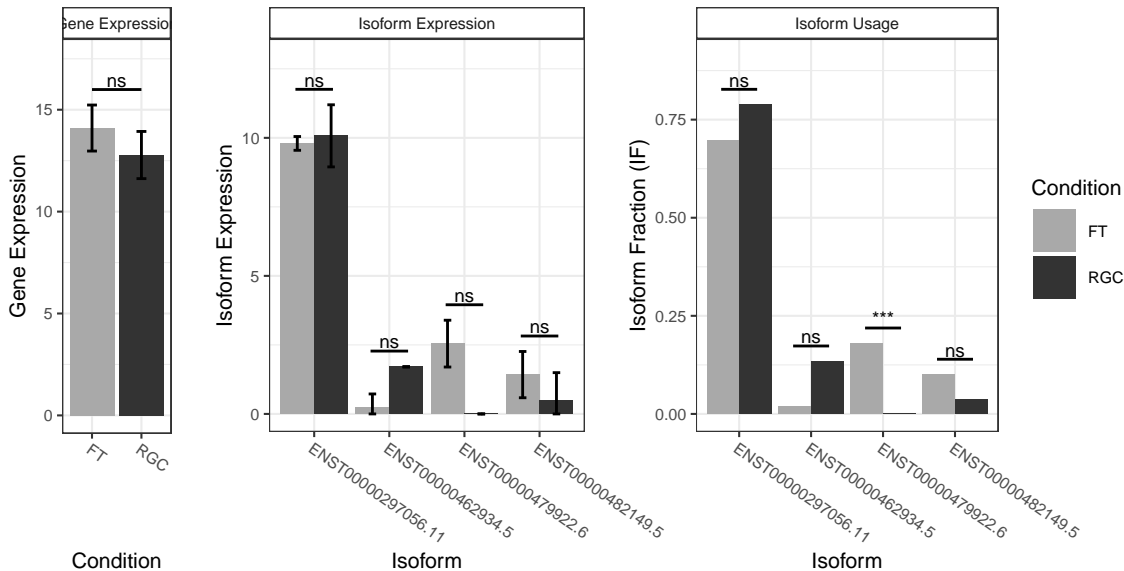
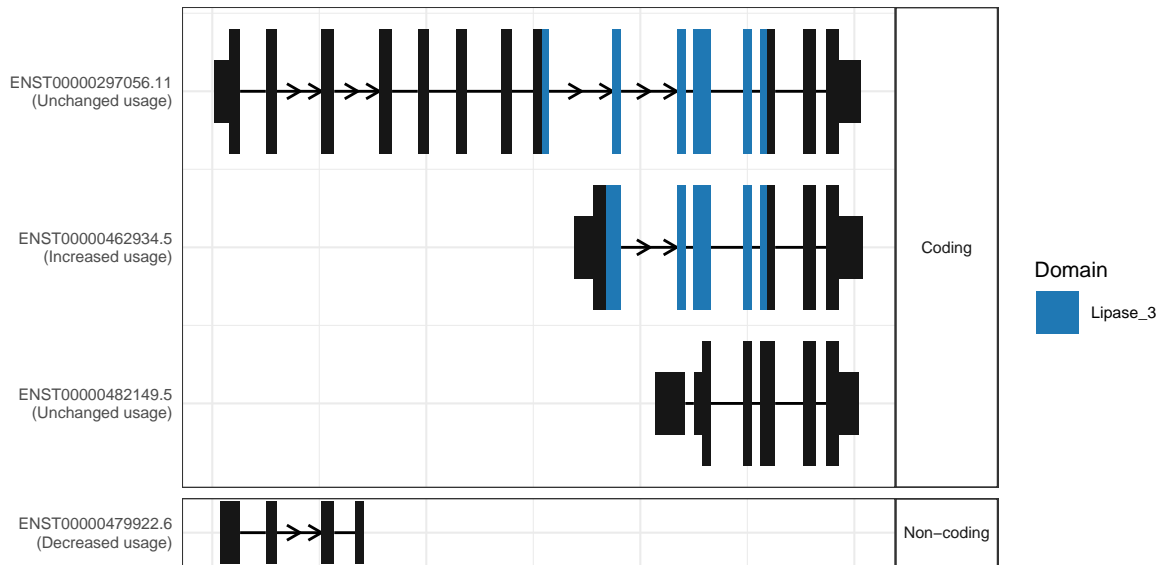


Domain



AMP-binding

The isoform switch in DAGLB (FT vs RGC)

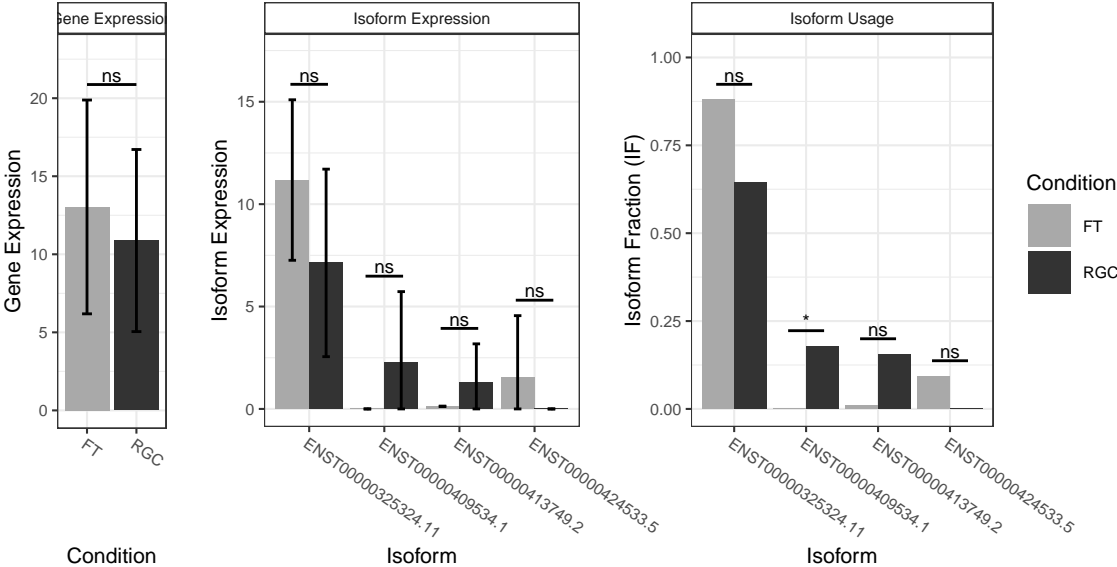
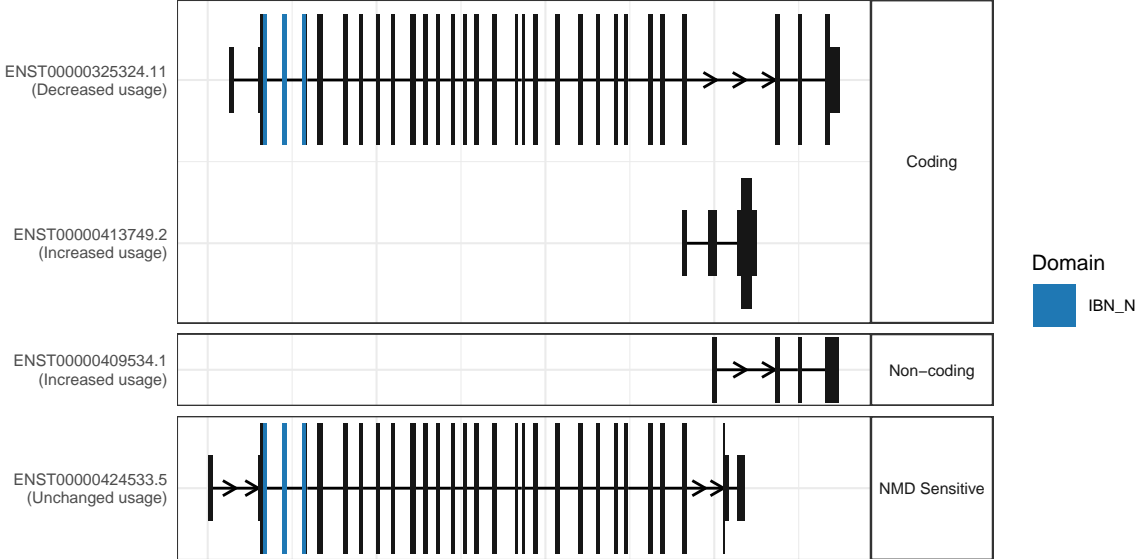


Domain



Lipase_3

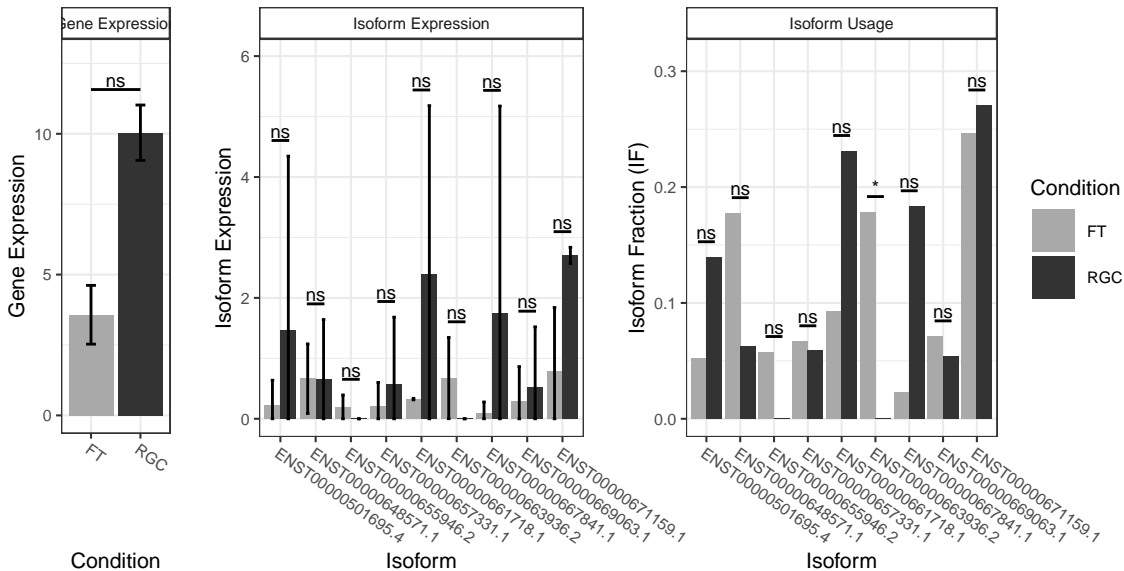
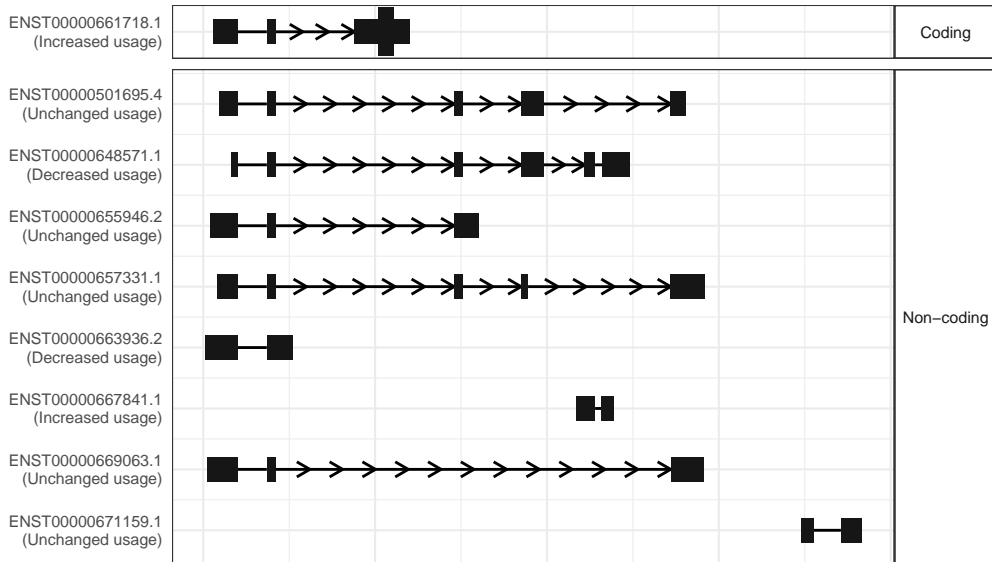
The isoform switch in IPO11 (FT vs RGC)



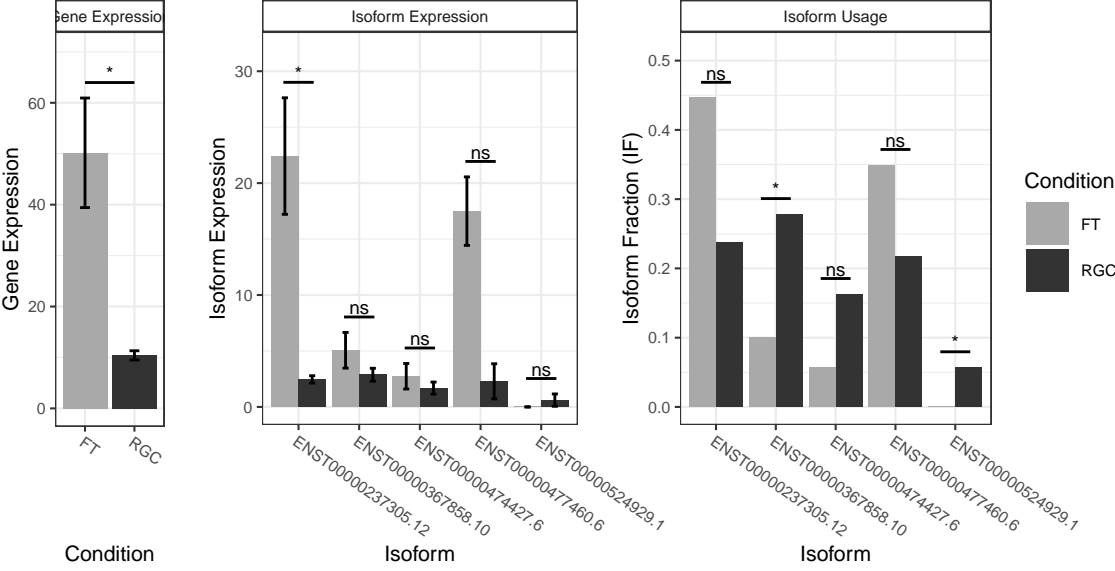
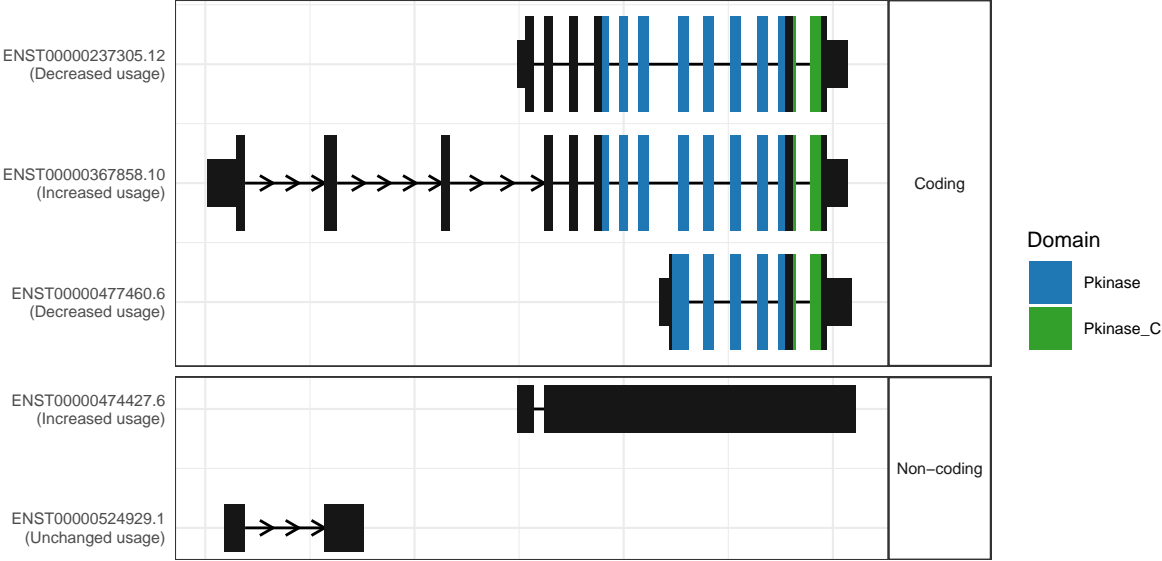
Domain

 IBN_N

The isoform switch in FBXO38-DT (FT vs RGC)



The isoform switch in SGK1 (FT vs RGC)



Domain

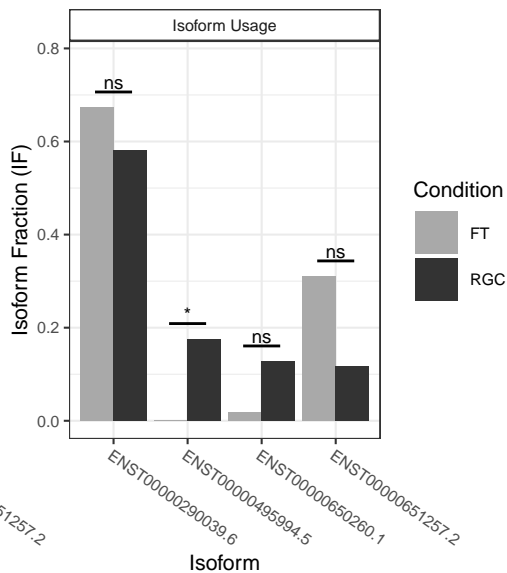
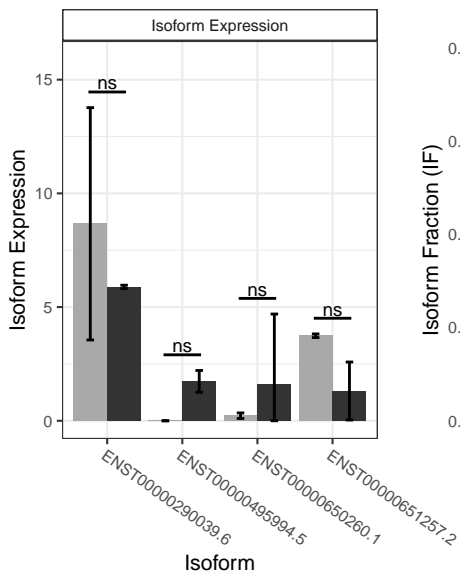
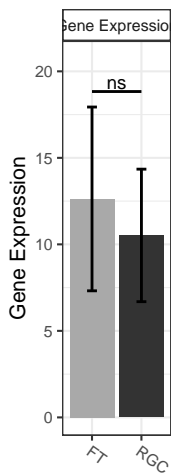
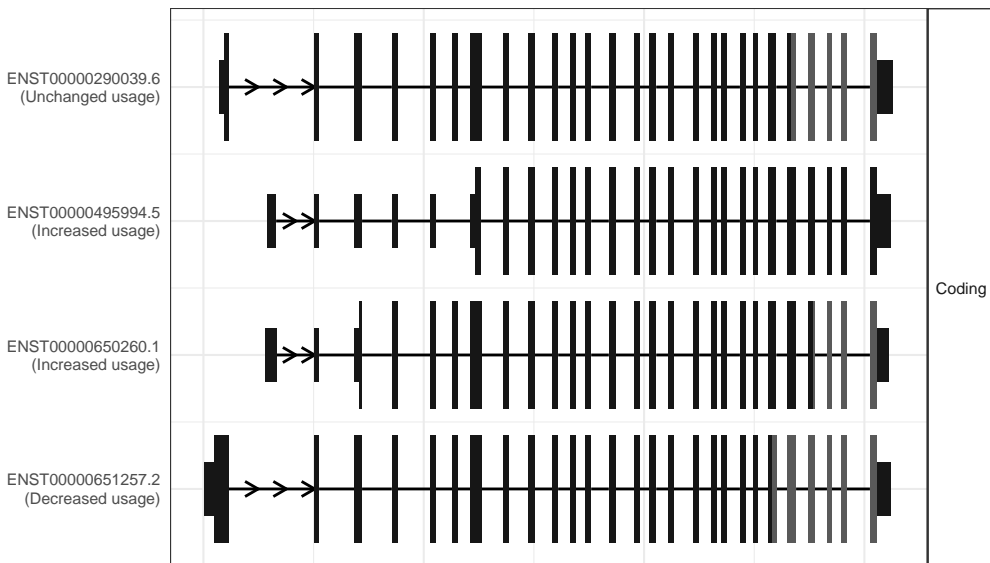


Pkinase



Pkinase_C

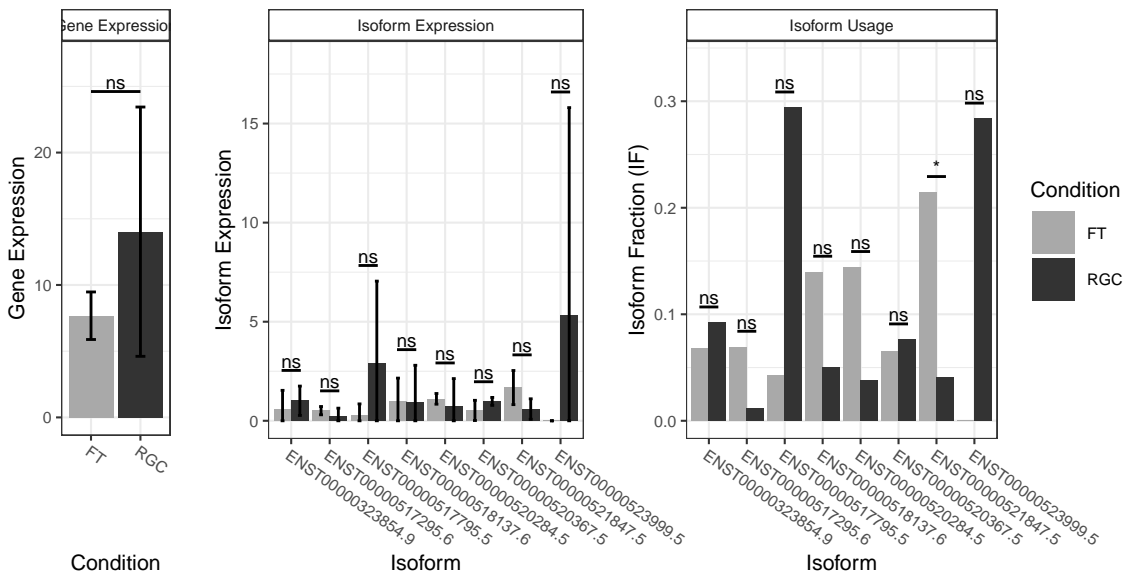
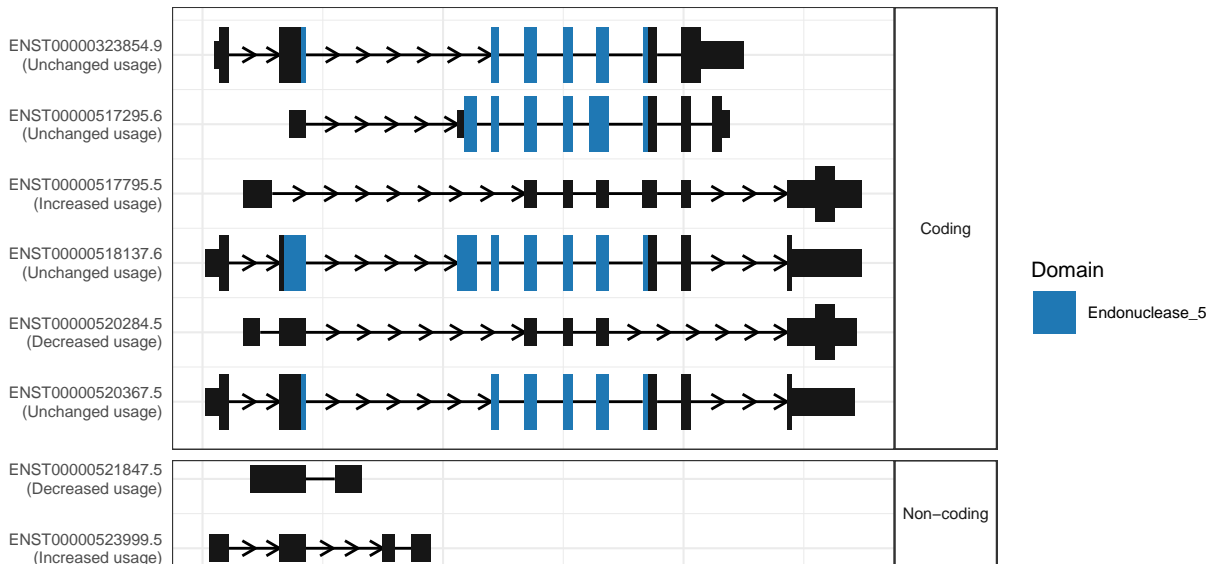
The isoform switch in CACHD1 (FT vs RGC)



Condition



The isoform switch in ENDOV (FT vs RGC)

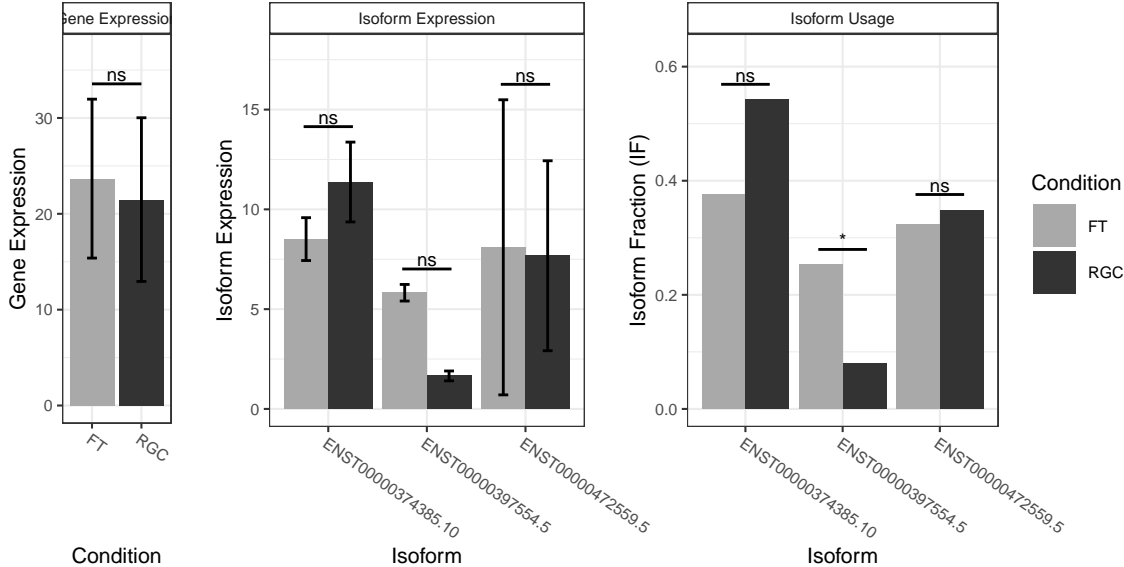
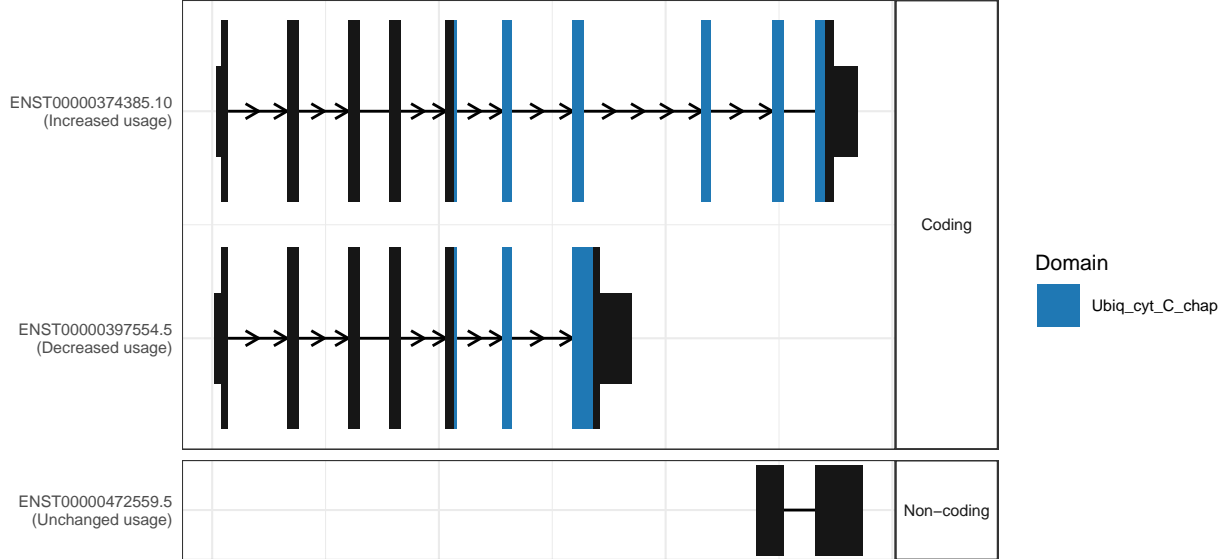


Domain



Endonuclease_5

The isoform switch in UQCC1 (FT vs RGC)

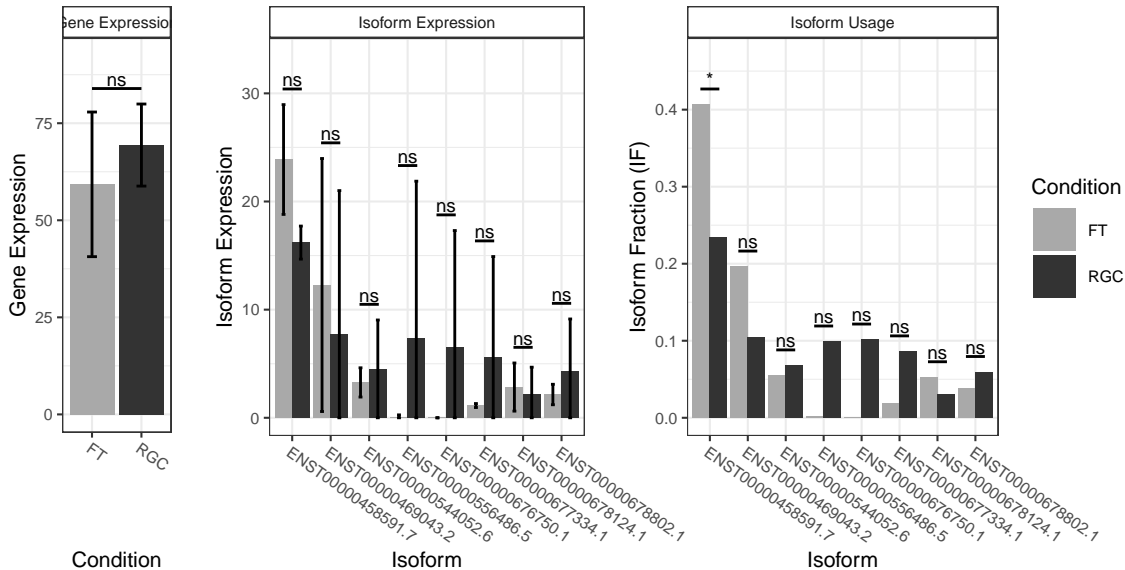
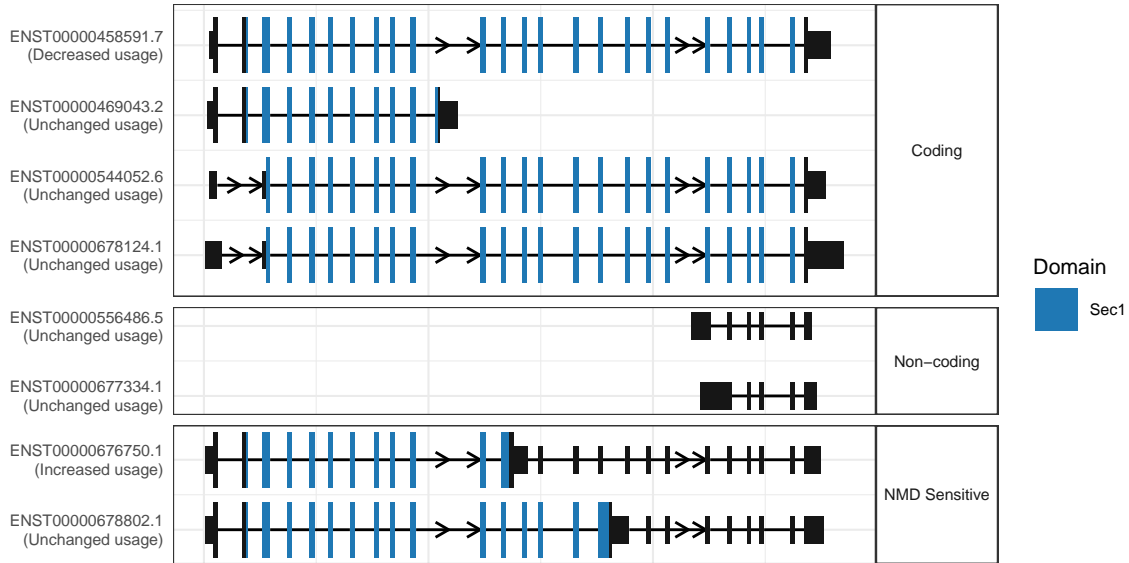


Domain



Ubiqu_cyt_C_chap

The isoform switch in SCFD1 (FT vs RGC)

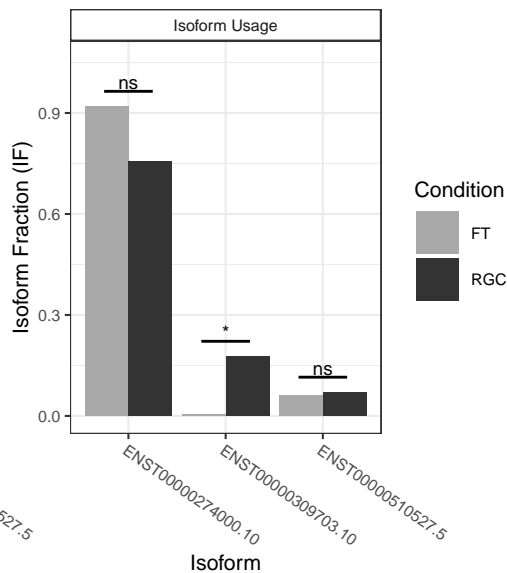
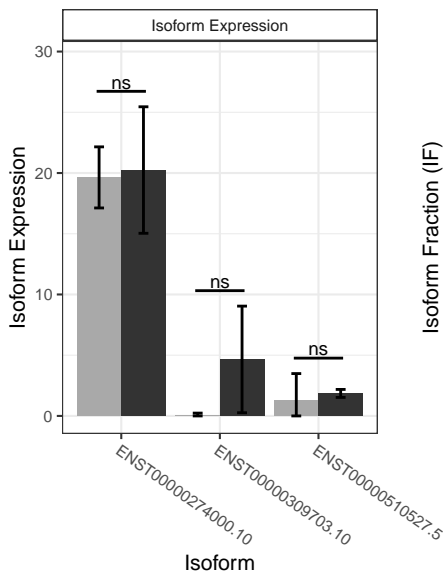
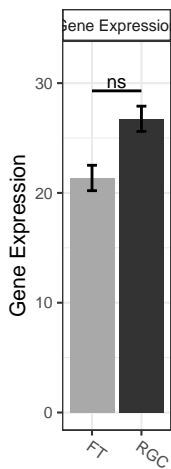
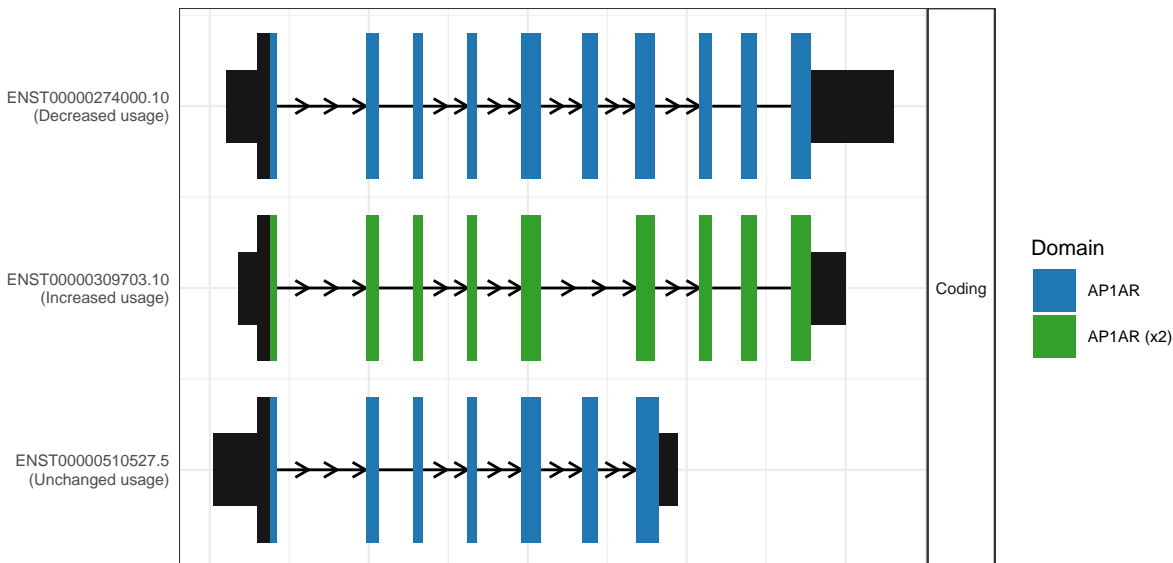


Domain



Sec1

The isoform switch in AP1AR (FT vs RGC)



Domain

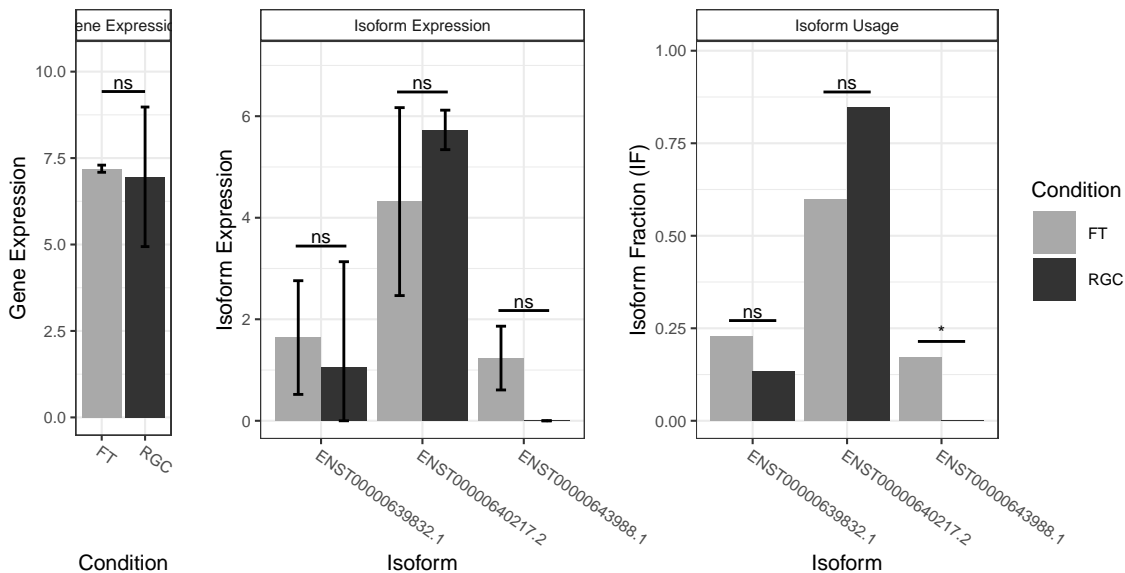
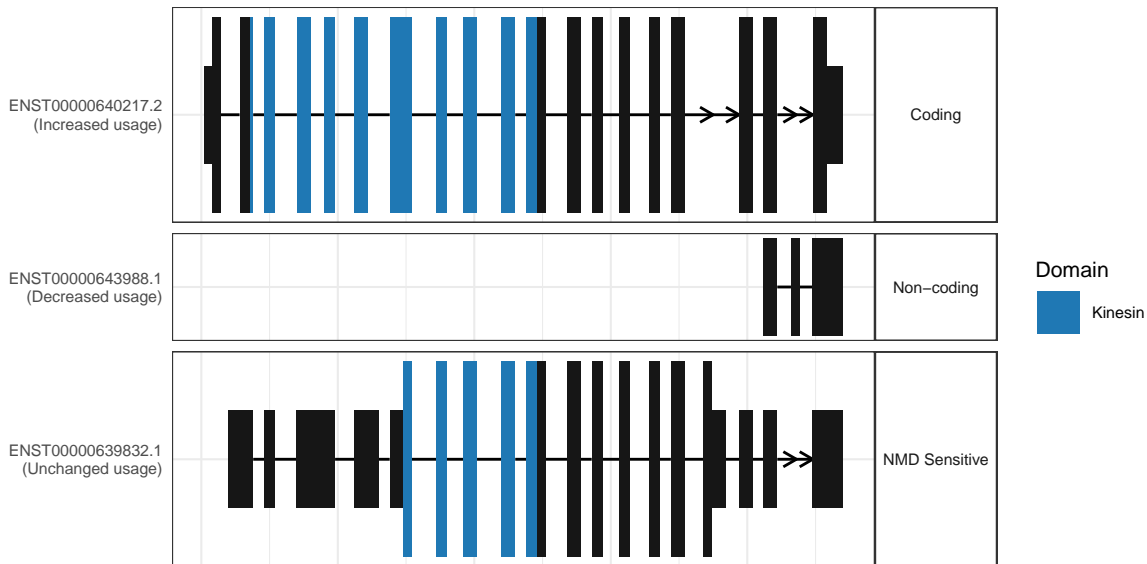


AP1AR



AP1AR (x2)

The isoform switch in KIF12 (FT vs RGC)

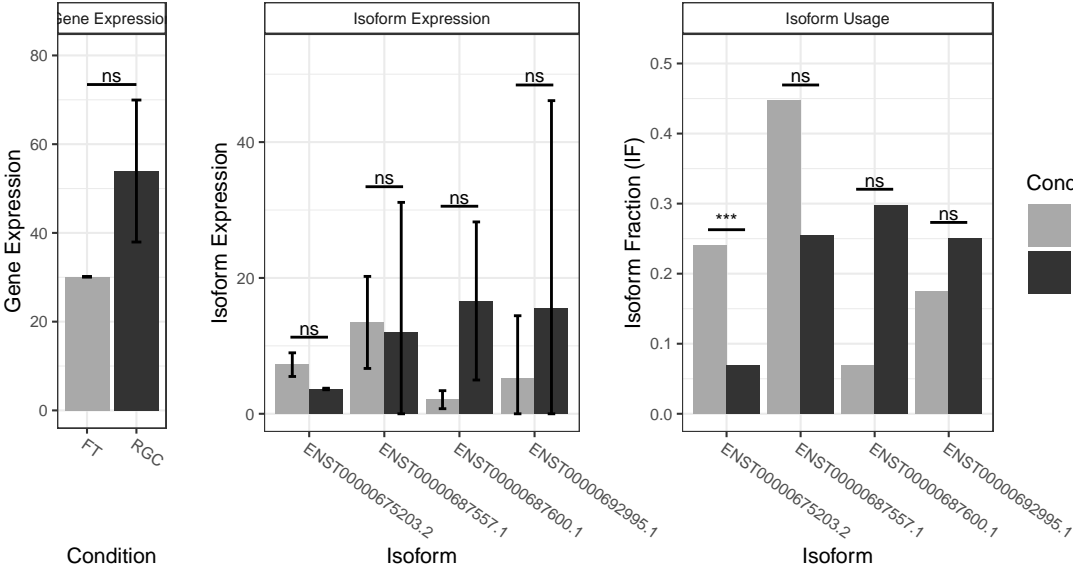
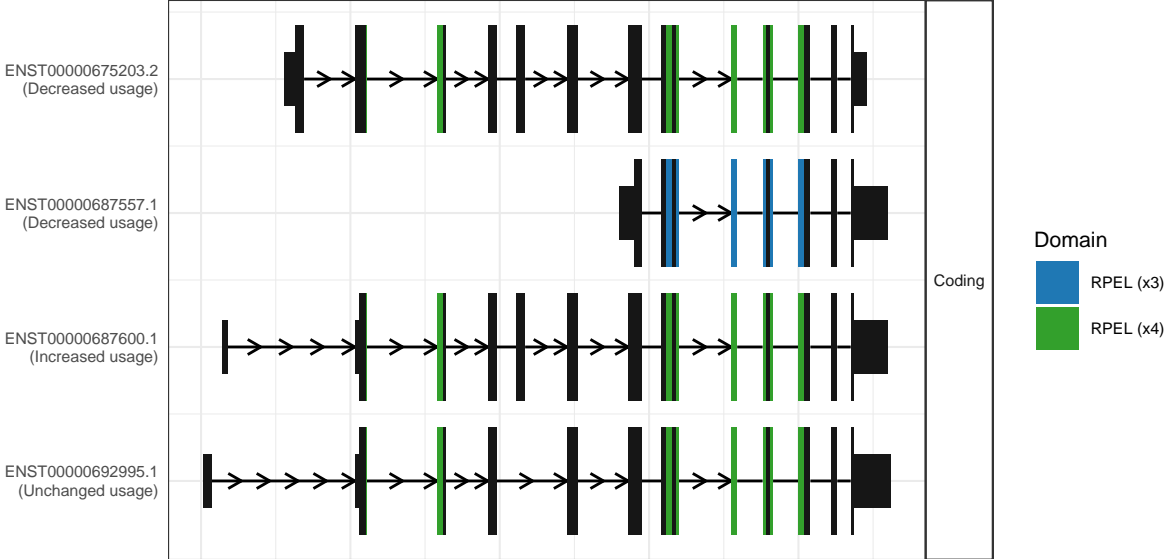


Domain



Kinesin

The isoform switch in PHACTR1 (FT vs RGC)



Domain

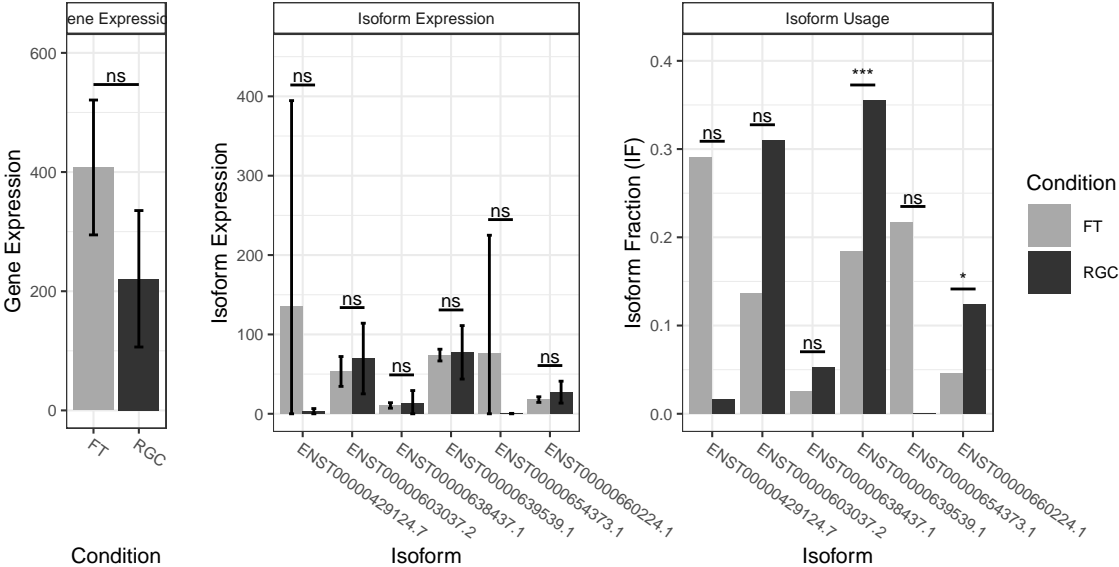
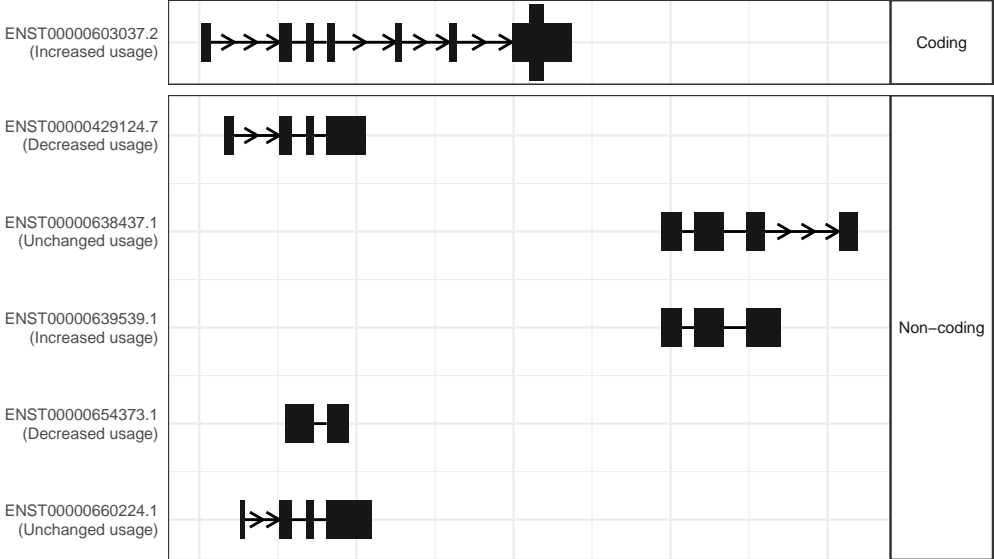


RPEL (x3)

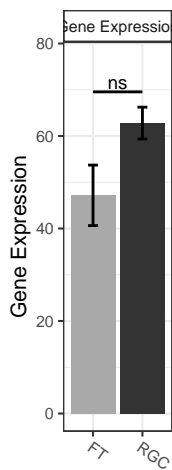
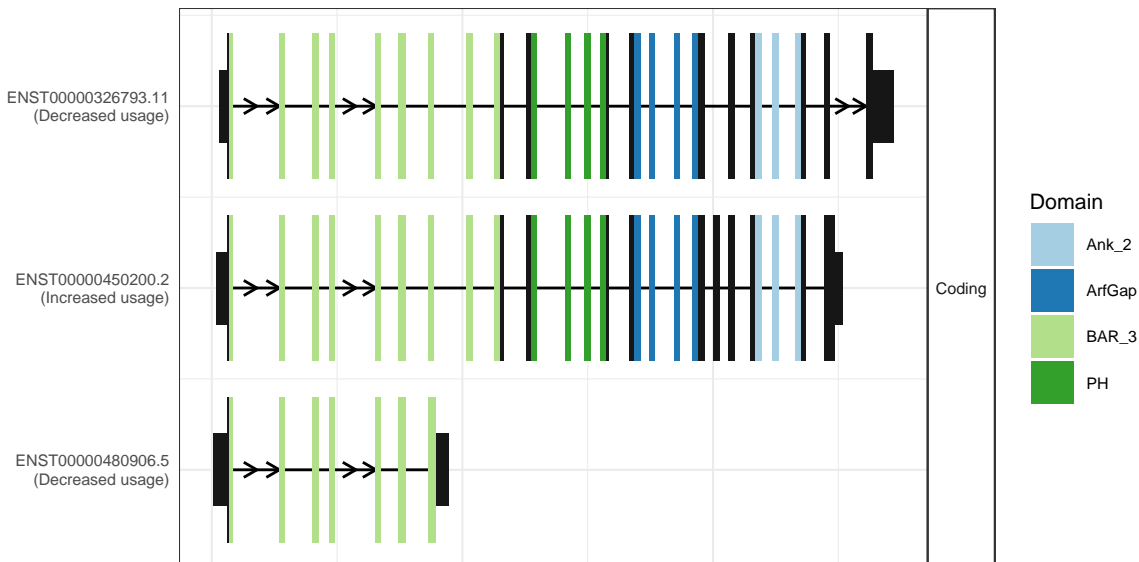


RPEL (x4)

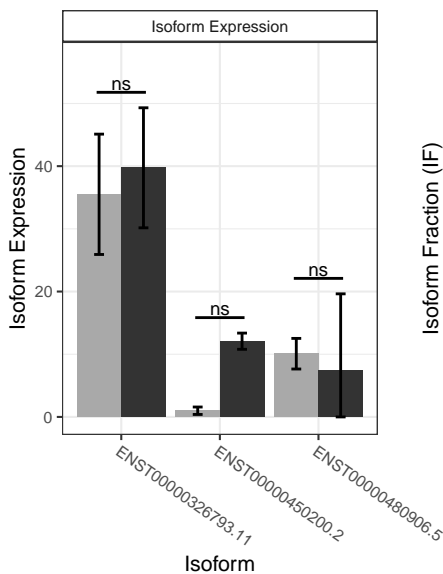
The isoform switch in FTX (FT vs RGC)



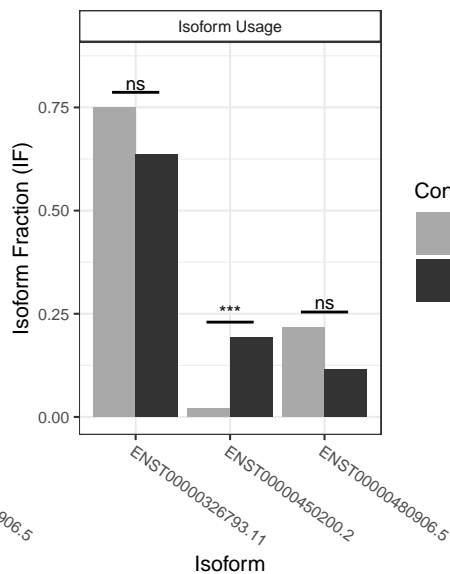
The isoform switch in ACAP2 (FT vs RGC)



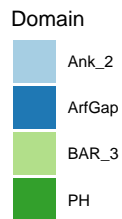
Condition



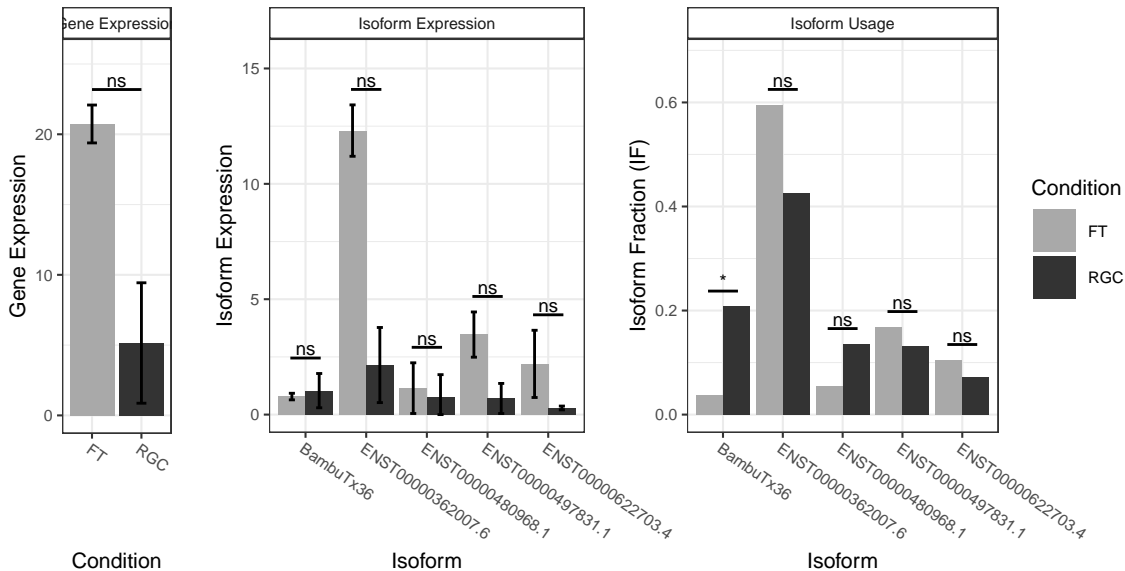
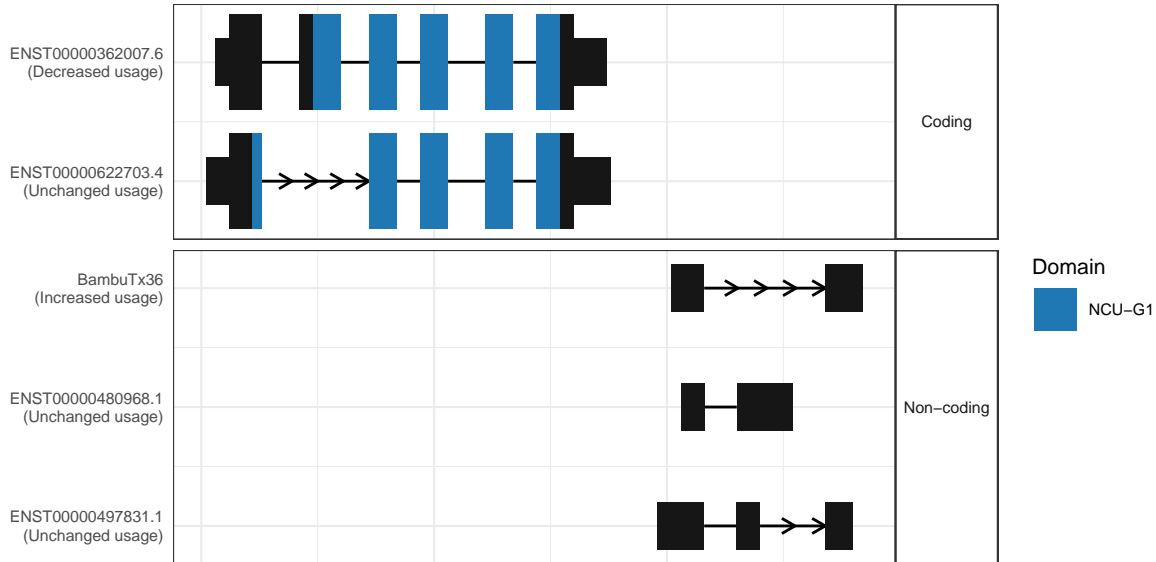
Isoform



Isoform



The isoform switch in GLMP (FT vs RGC)

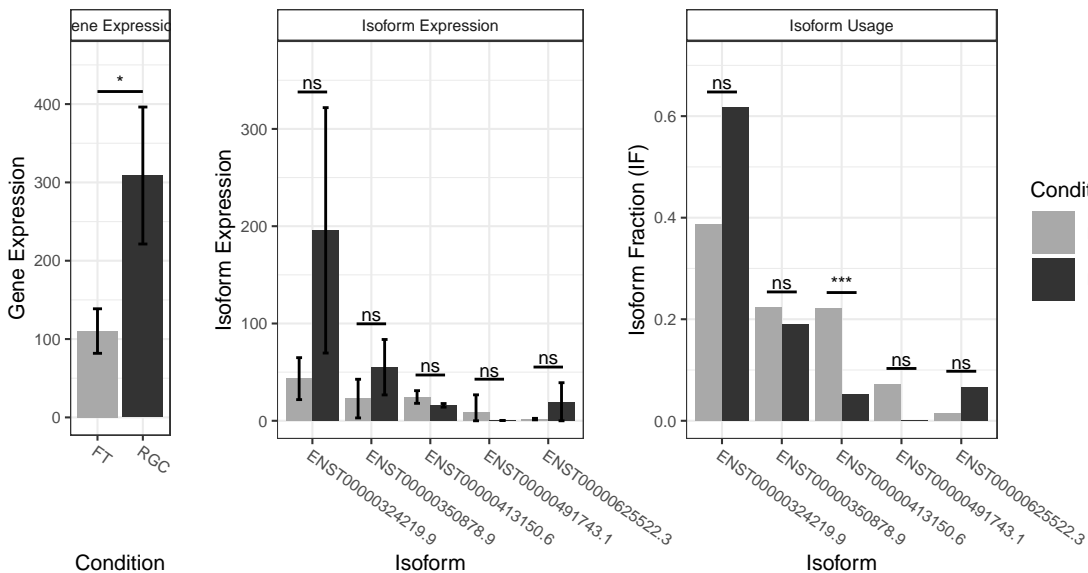
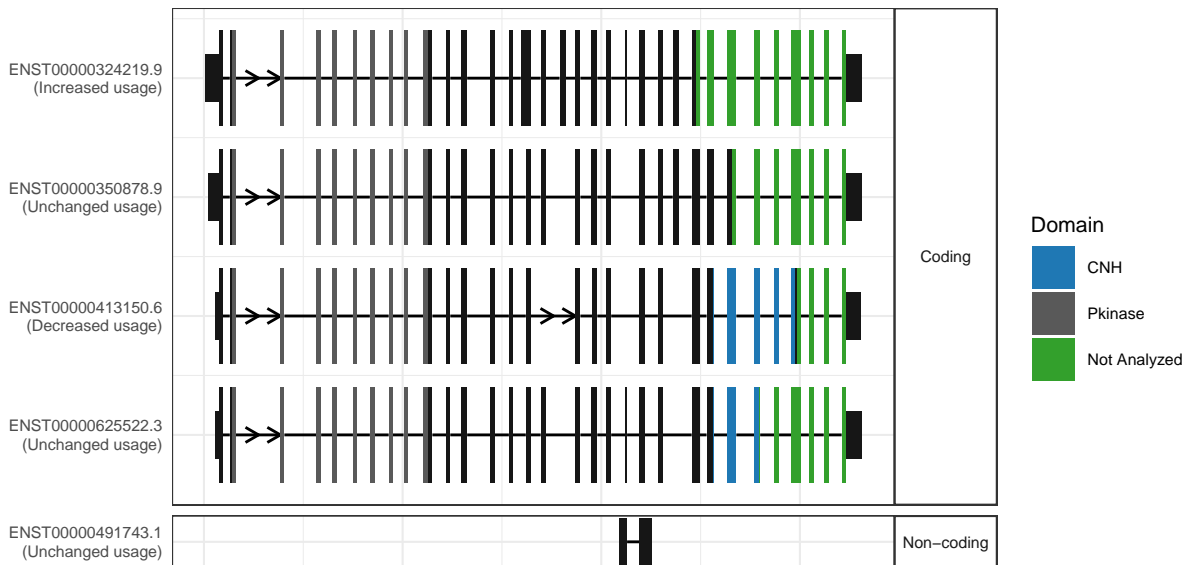


Domain



NCU-G1

The isoform switch in MAP4K4 (FT vs RGC)



Domain



CNH

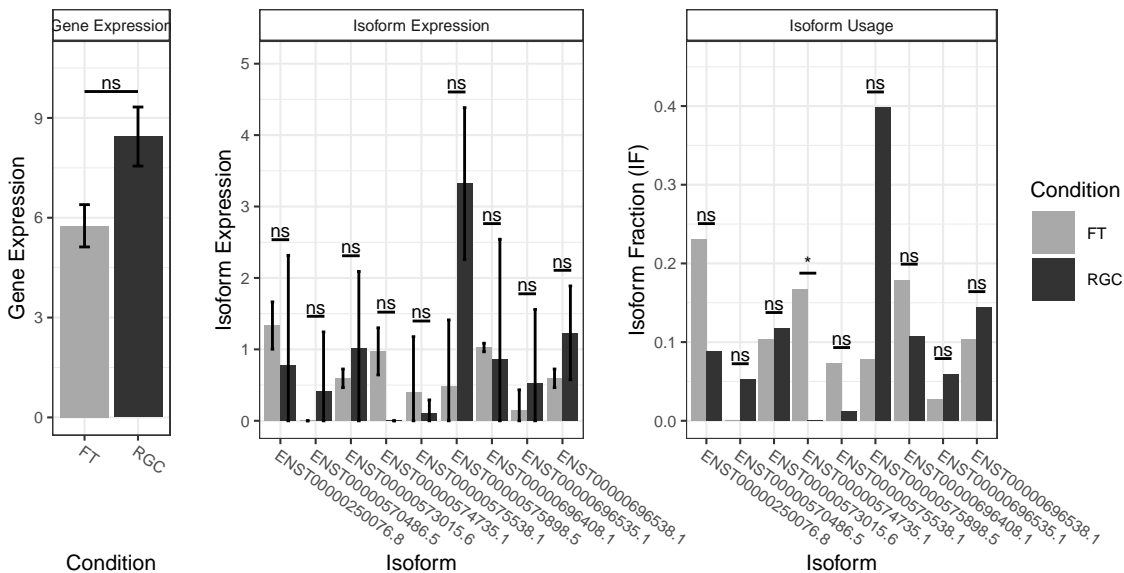
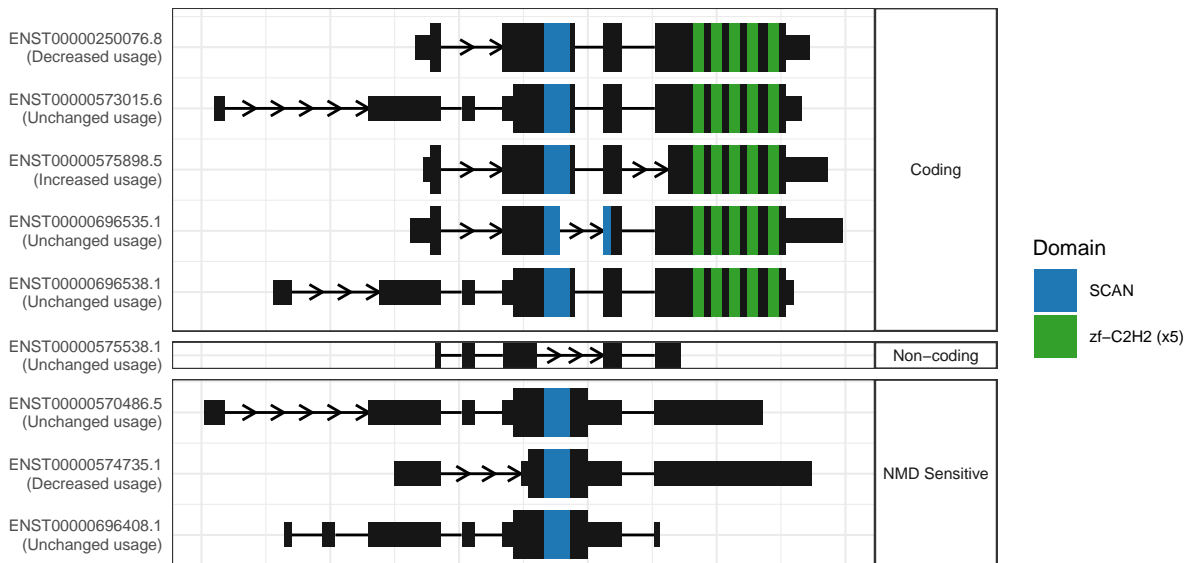


Pkinase



Not Analyzed

The isoform switch in ZNF232 (FT vs RGC)



Domain

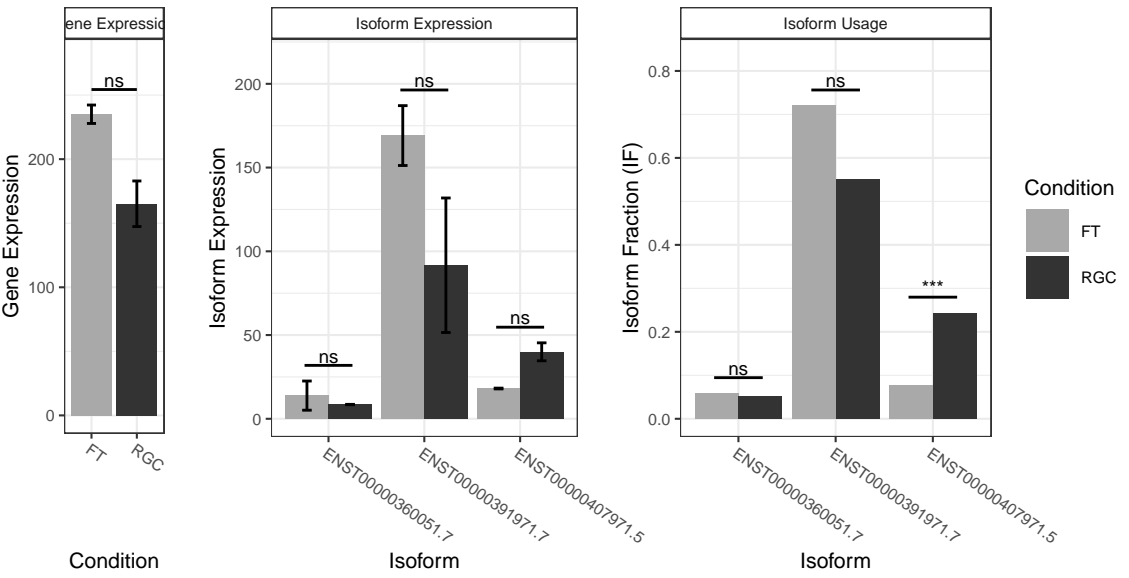
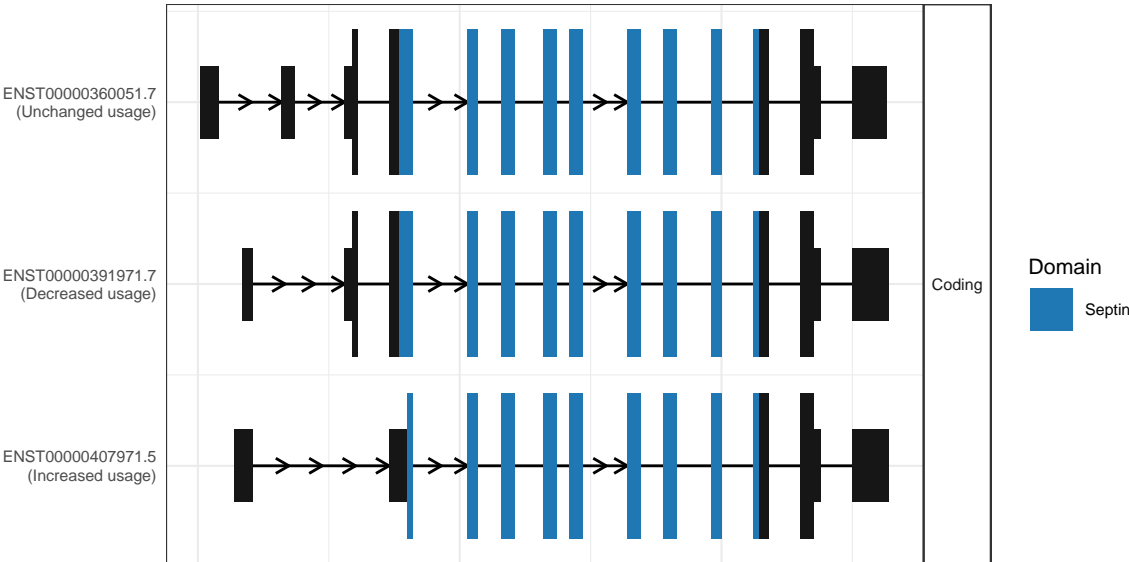


SCAN



zf-C2H2 (x5)

The isoform switch in SEPTIN2 (FT vs RGC)

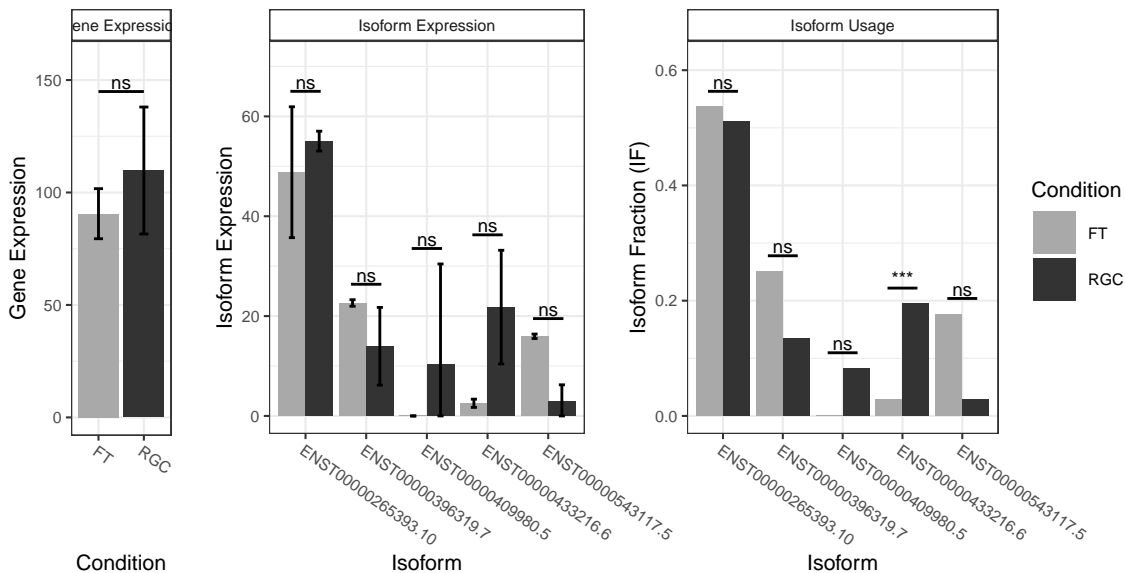
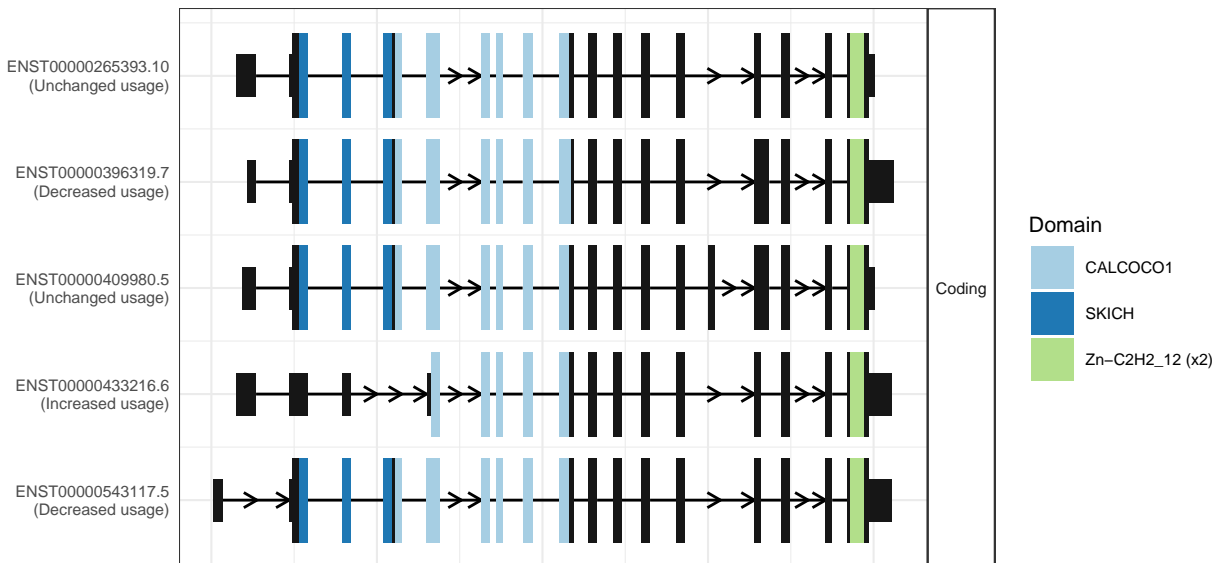


Domain



Septin

The isoform switch in TAX1BP1 (FT vs RGC)



Domain



CALCOCO1

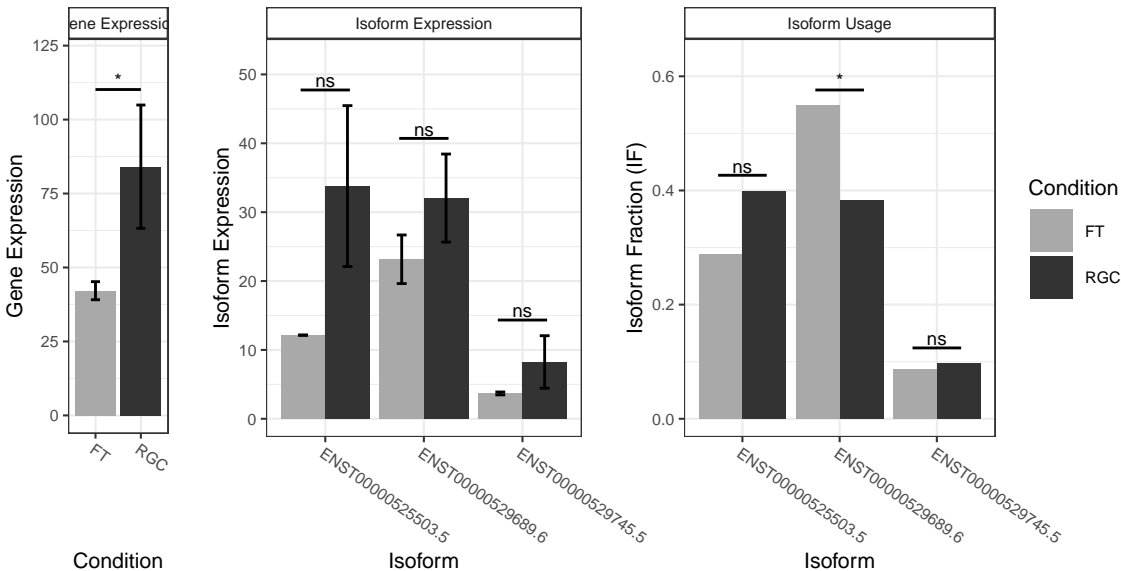
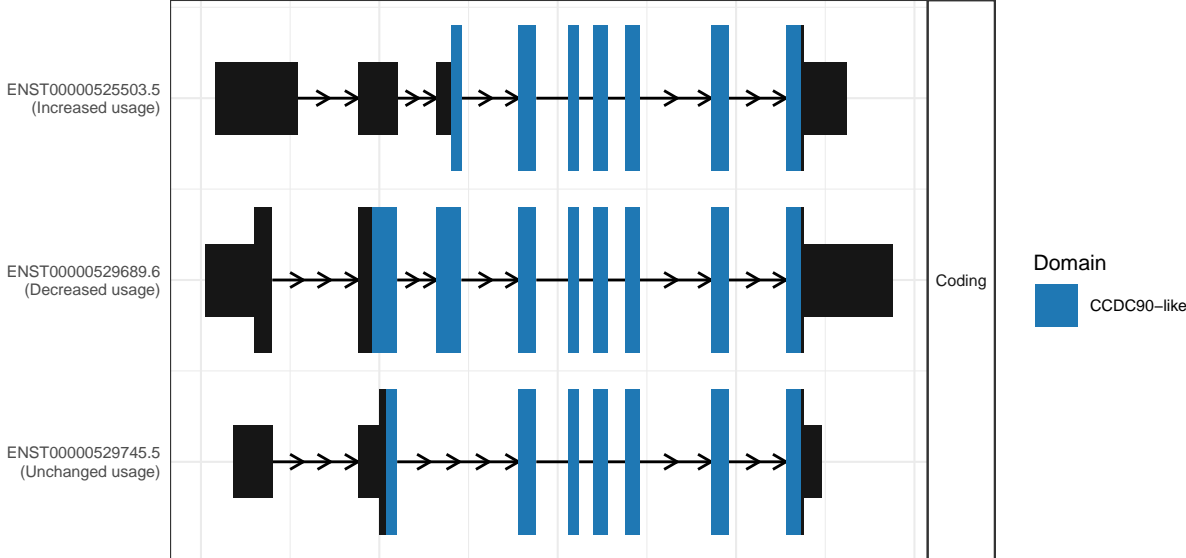


SKICH



Zn-C2H2_12 (x2)

The isoform switch in CCDC90B (FT vs RGC)

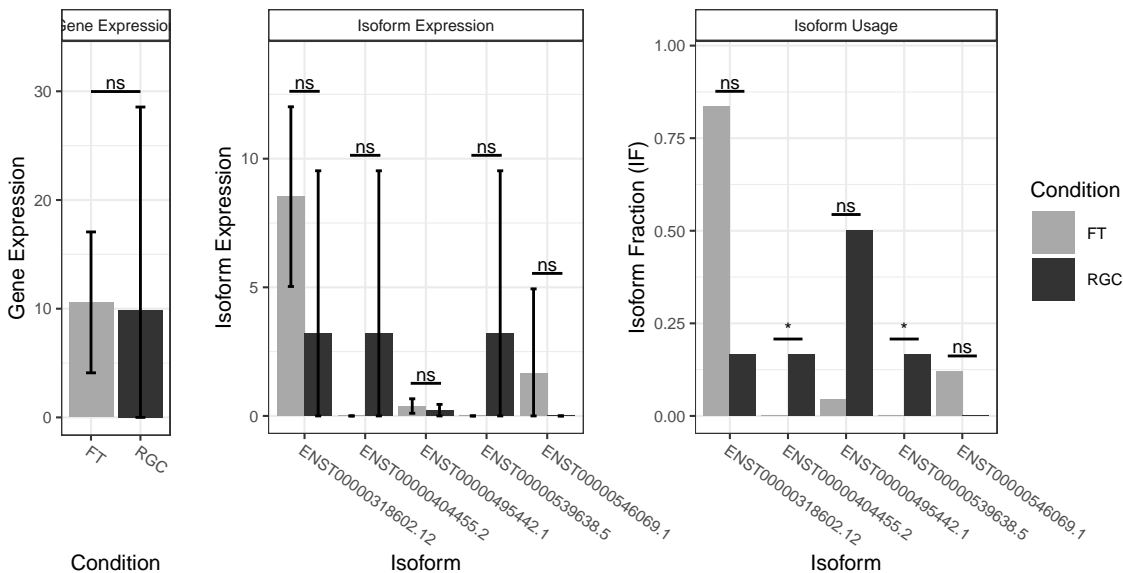
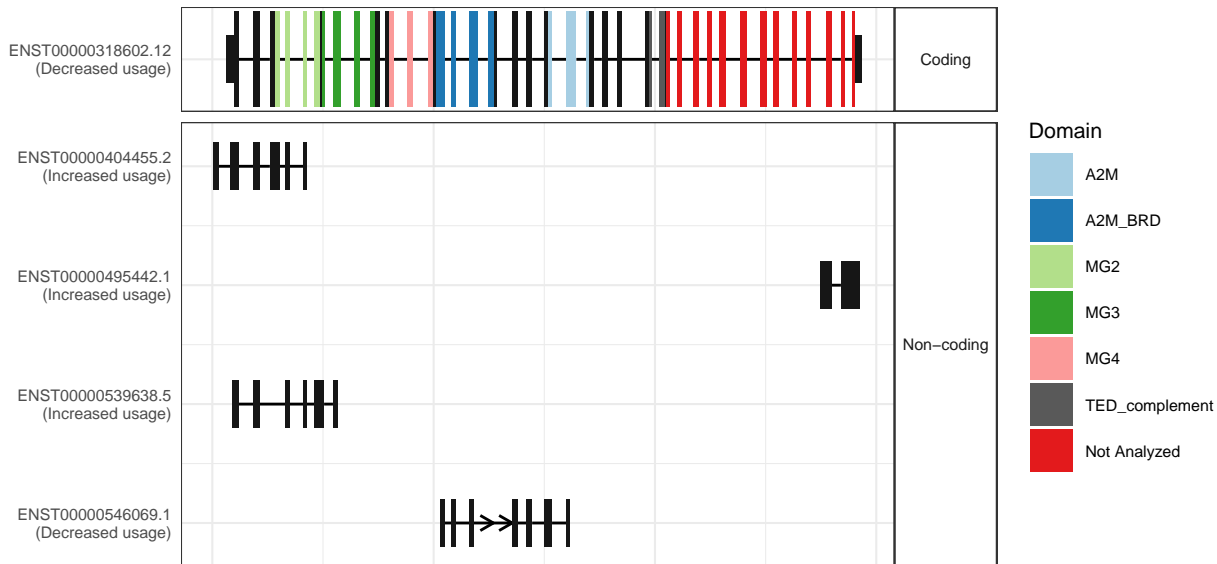


Domain

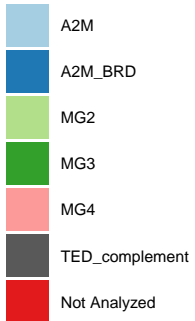


CCDC90-like

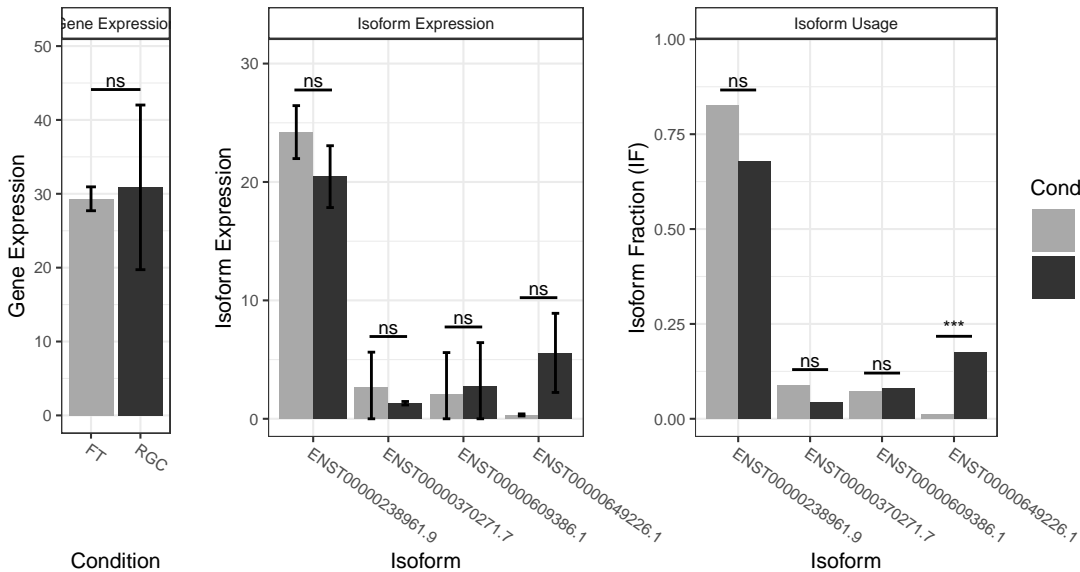
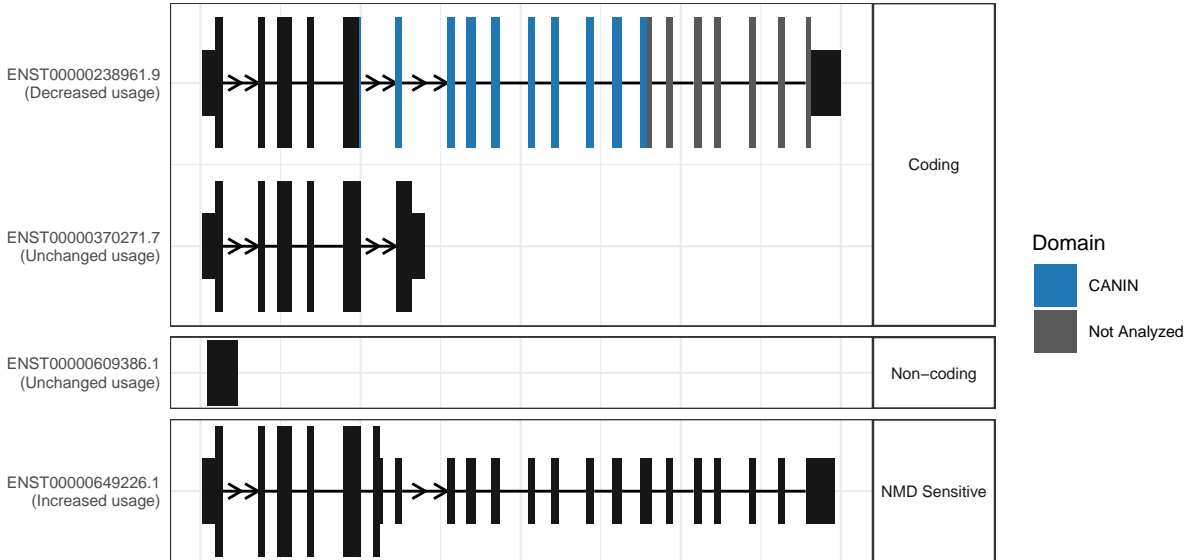
The isoform switch in A2M (FT vs RGC)



Domain



The isoform switch in SLF2 (FT vs RGC)



Domain

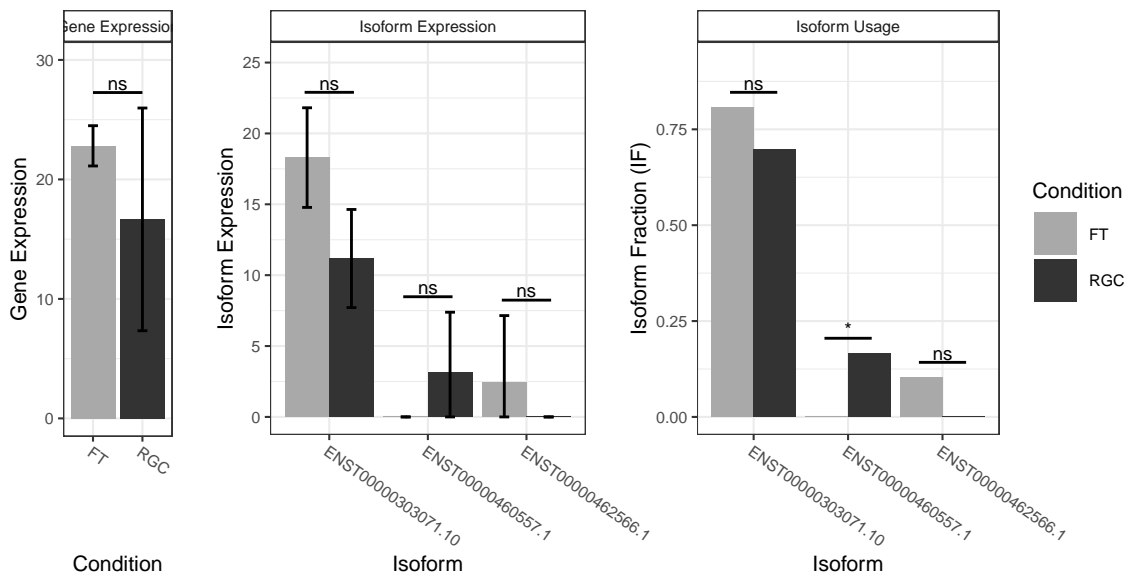
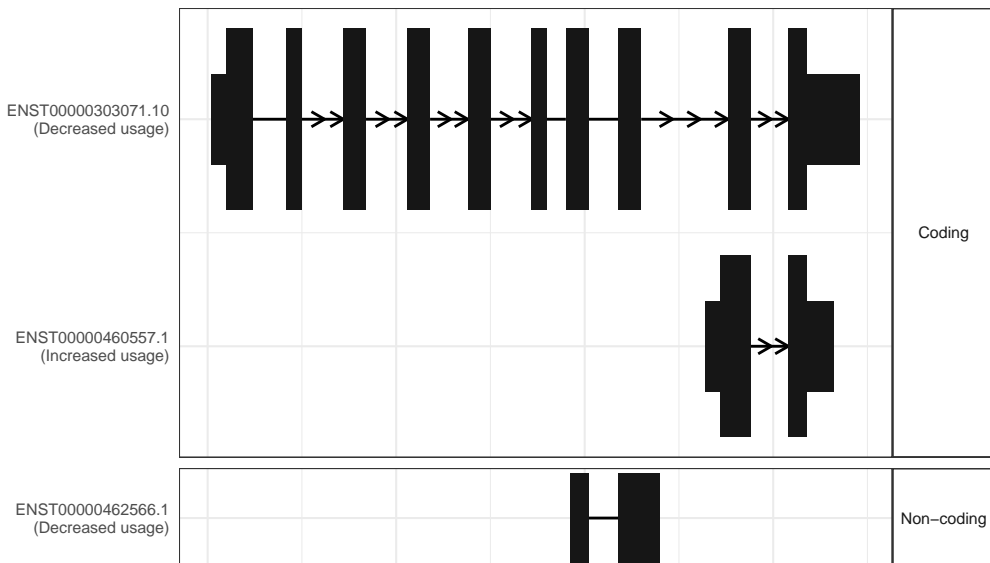


CANIN

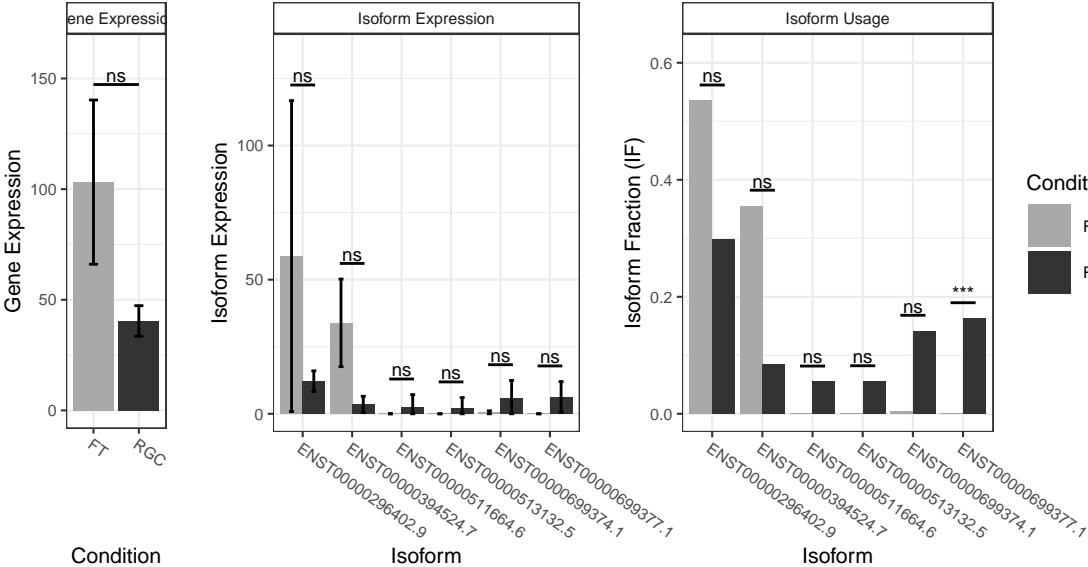
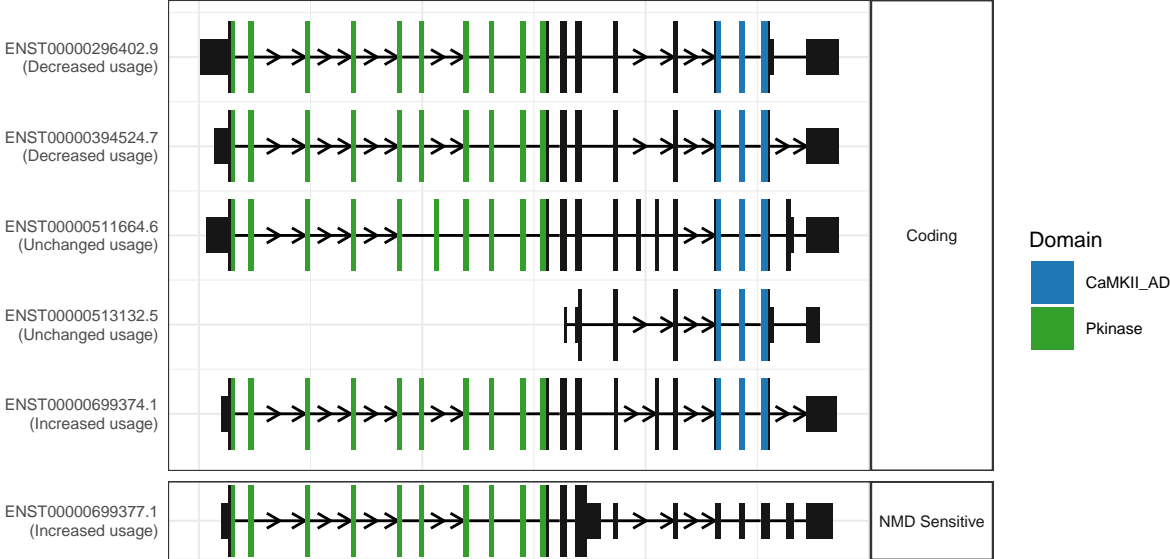


Not Analyzed

The isoform switch in DONSON (FT vs RGC)



The isoform switch in CAMK2D (FT vs RGC)



Domain

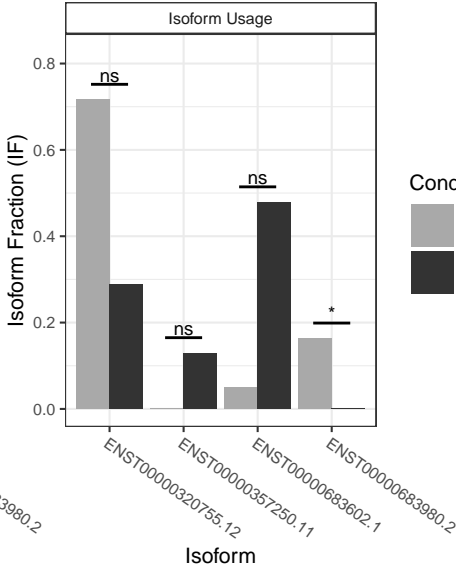
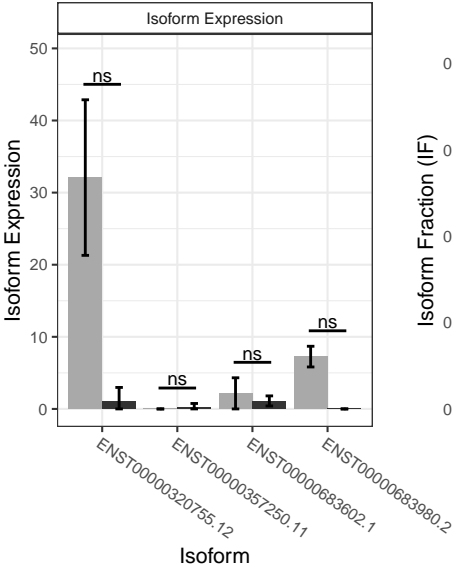
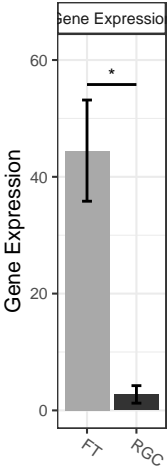
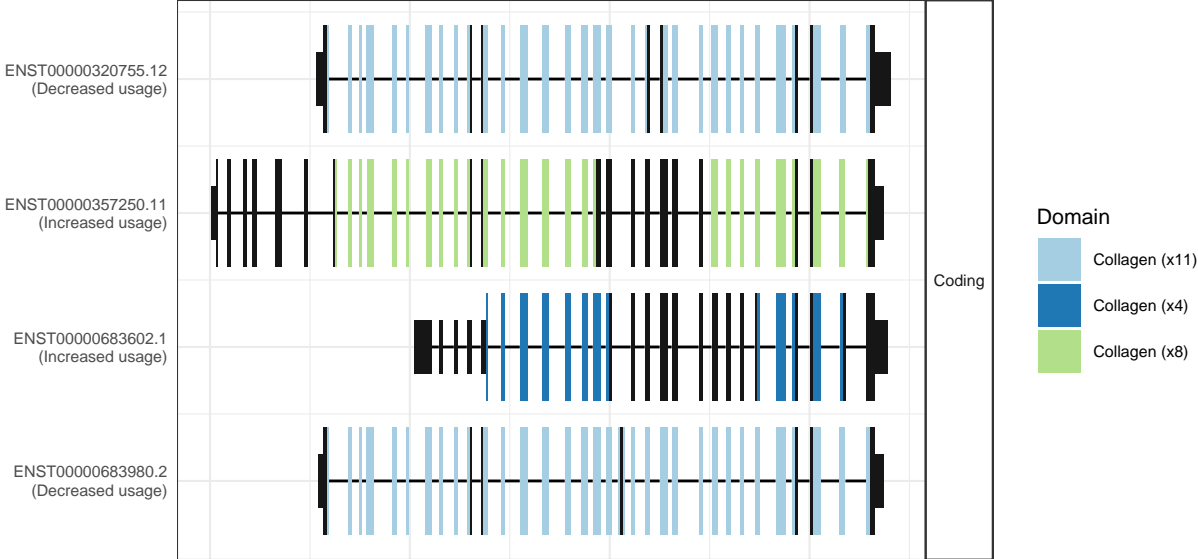


CaMKII_AD



Pkinase

The isoform switch in COL9A1(FT vs RGC)



Domain



Collagen (x11)

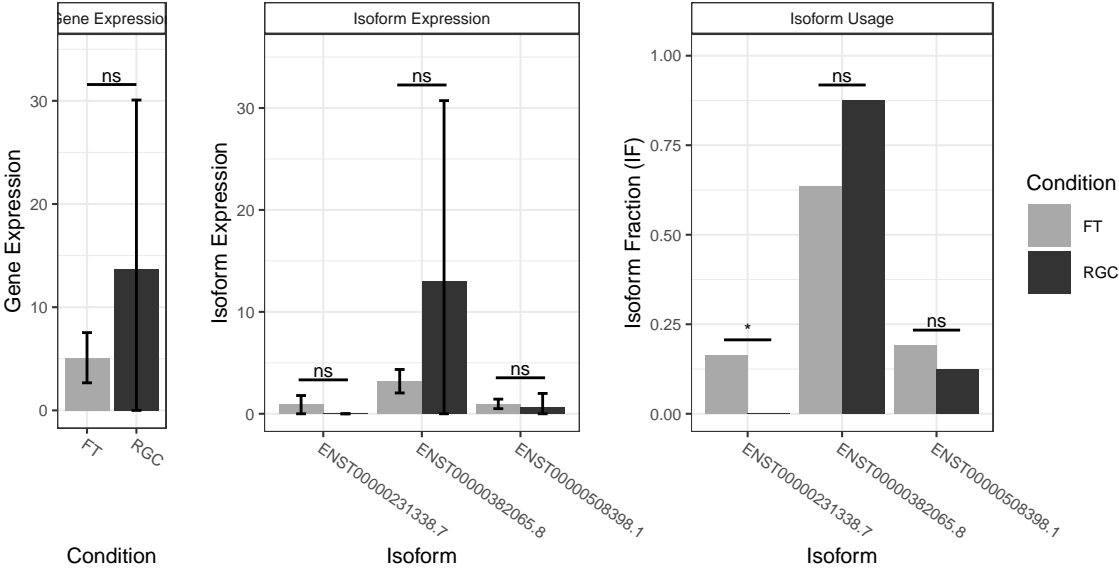
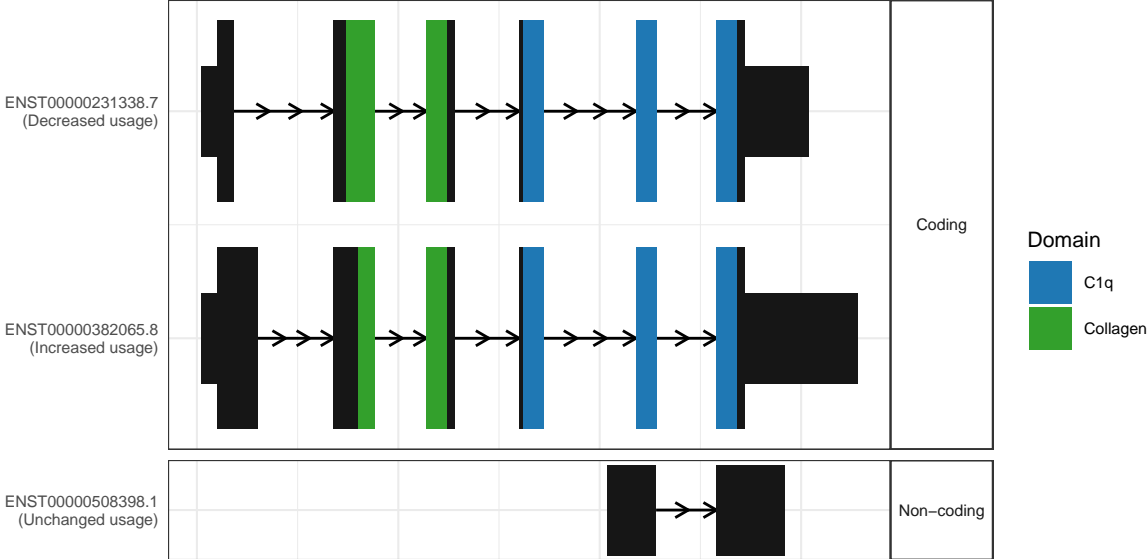


Collagen (x4)



Collagen (x8)

The isoform switch in C1QTNF3 (FT vs RGC)



Domain

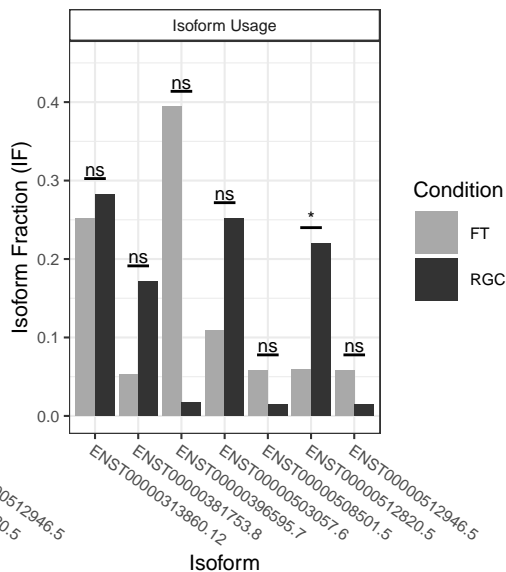
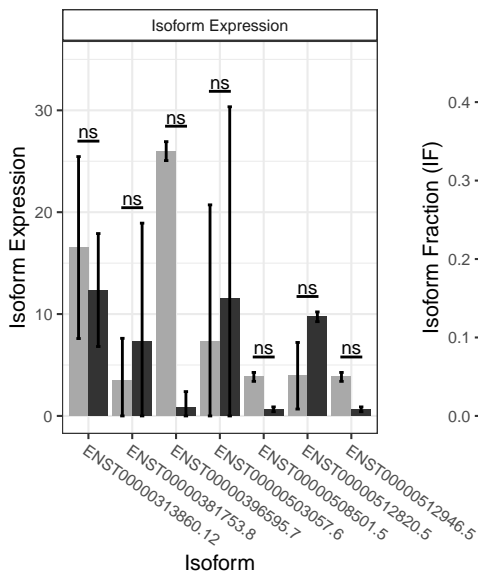
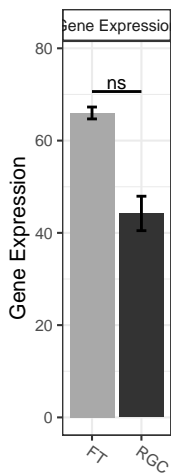
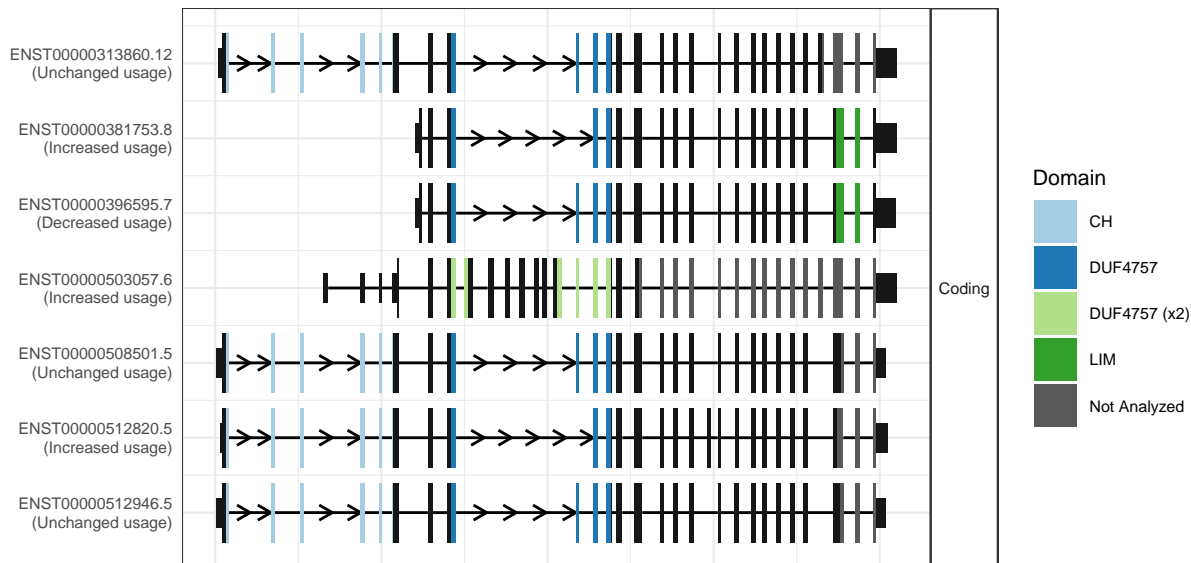


C1q








Collagen

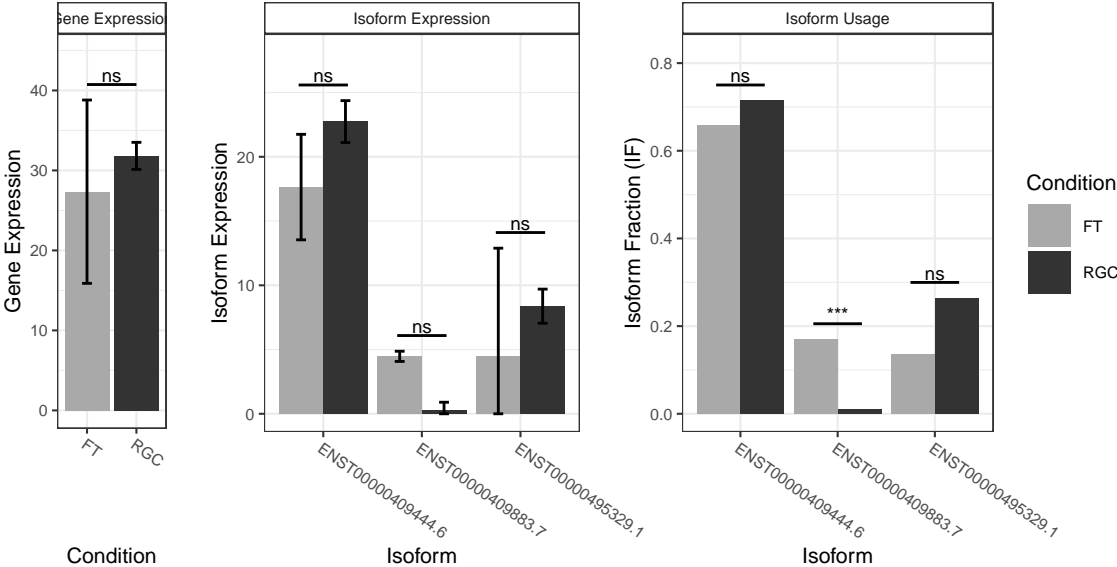
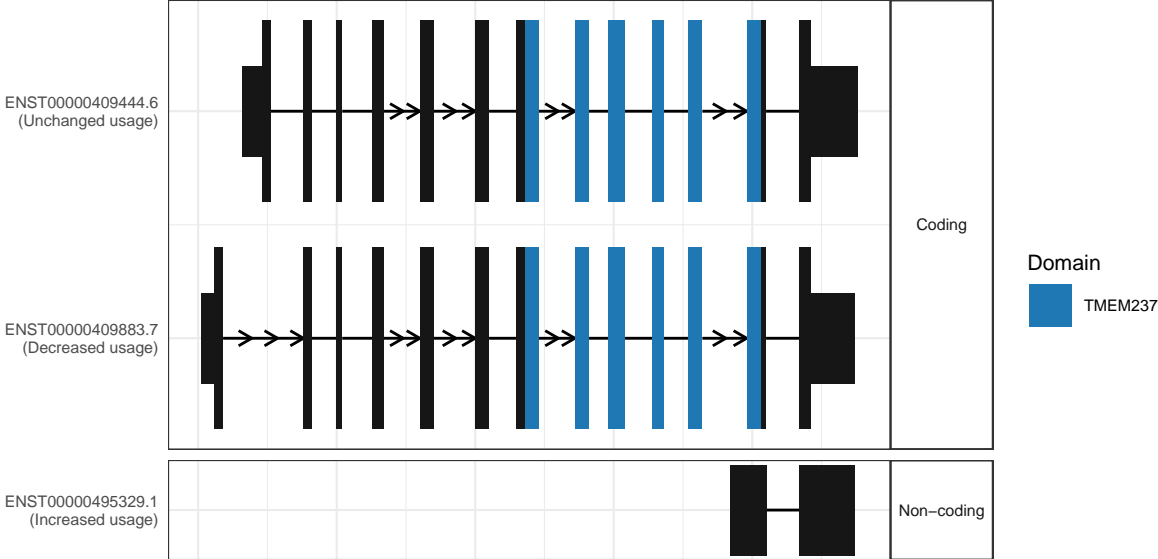
The isoform switch in LIMCH1 (FT vs RGC)



Domain

	CH
	DUF4757
	DUF4757 (x2)
	LIM
	Not Analyzed

The isoform switch in TMEM237 (FT vs RGC)

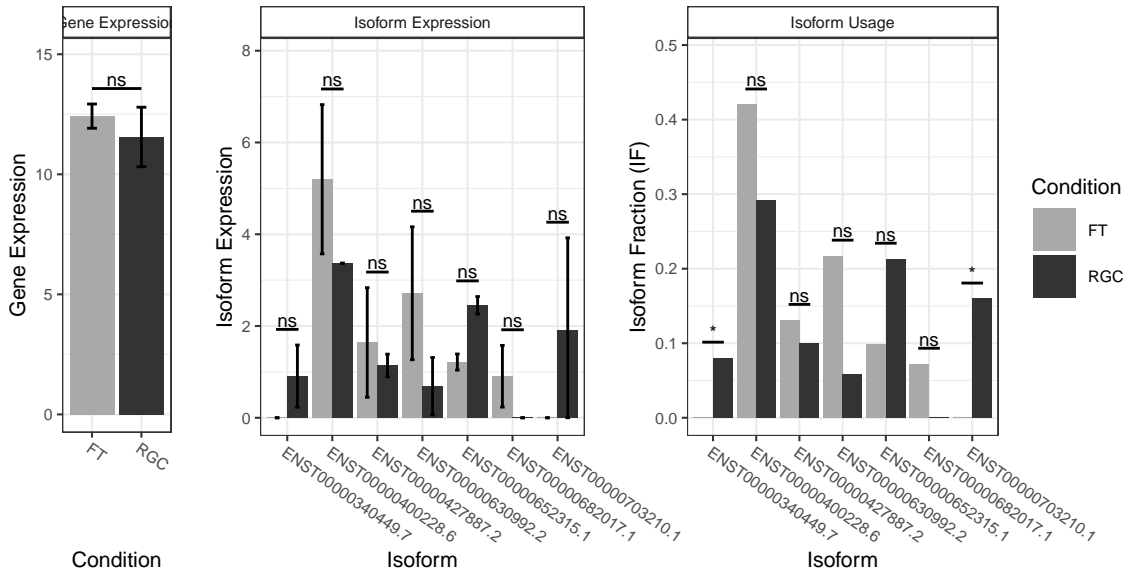
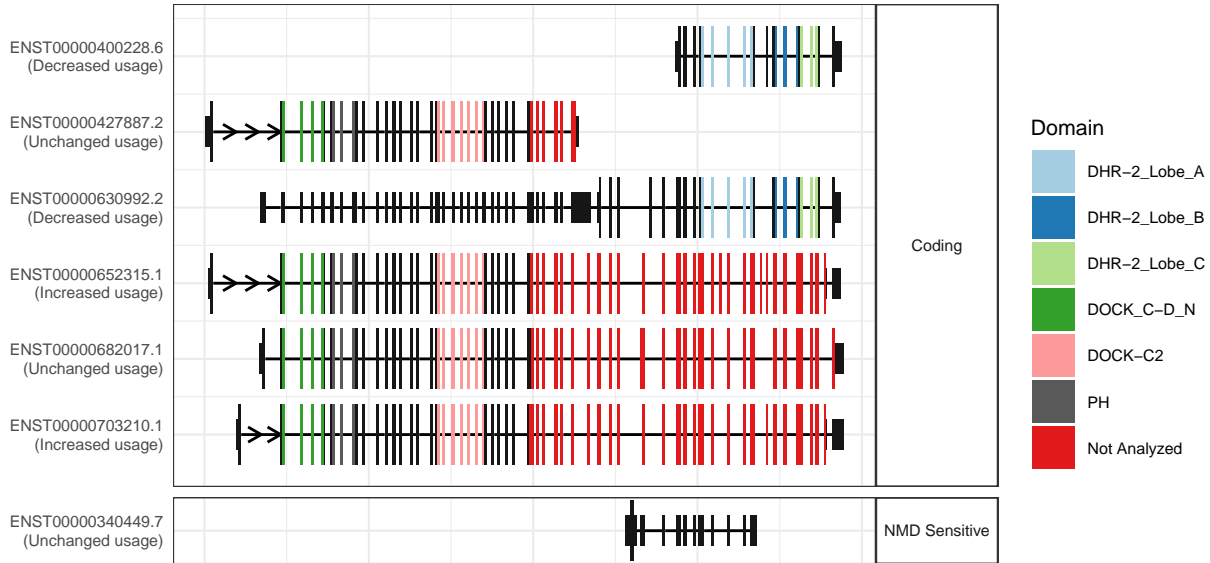


Domain










TMEM237

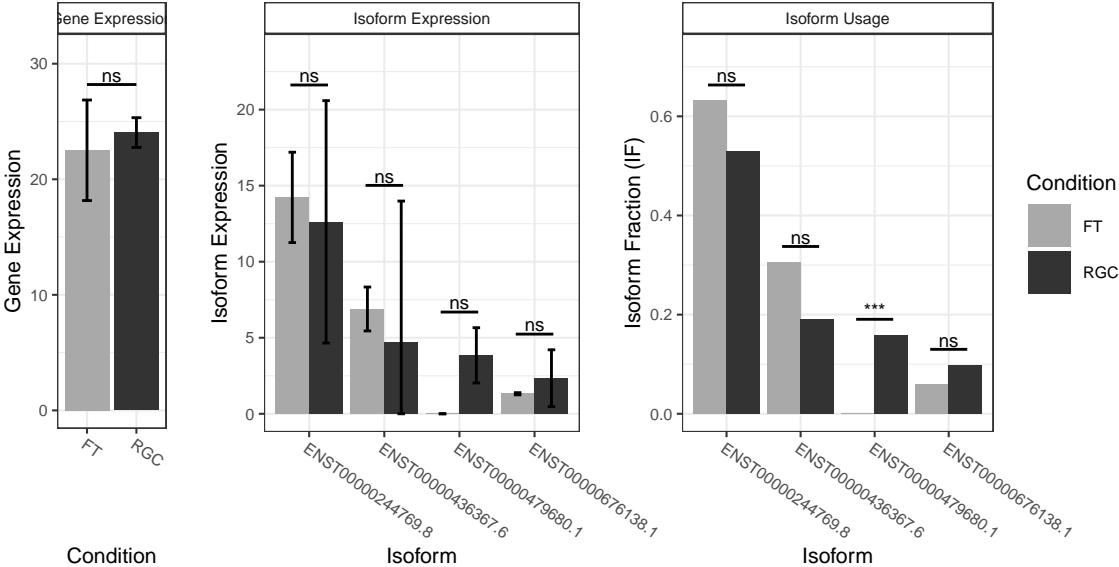
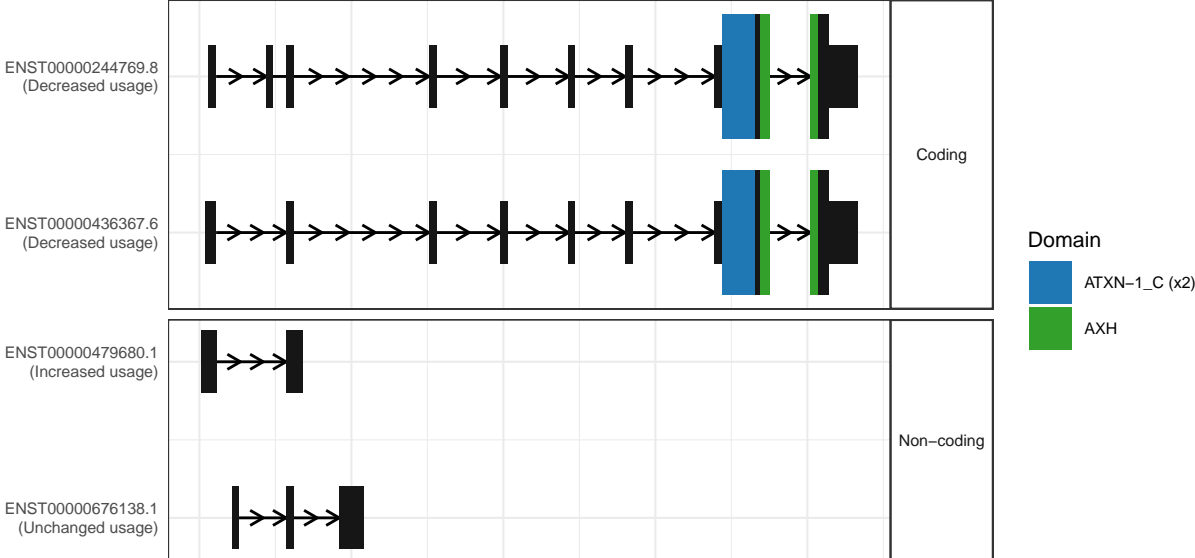
The isoform switch in DOCK9 (FT vs RGC)



Domain

	DHR~2_Lobe_A
	DHR~2_Lobe_B
	DHR~2_Lobe_C
	DOCK_C-D_N
	DOCK-C2
	PH
	Not Analyzed

The isoform switch in ATXN1 (FT vs RGC)



Domain

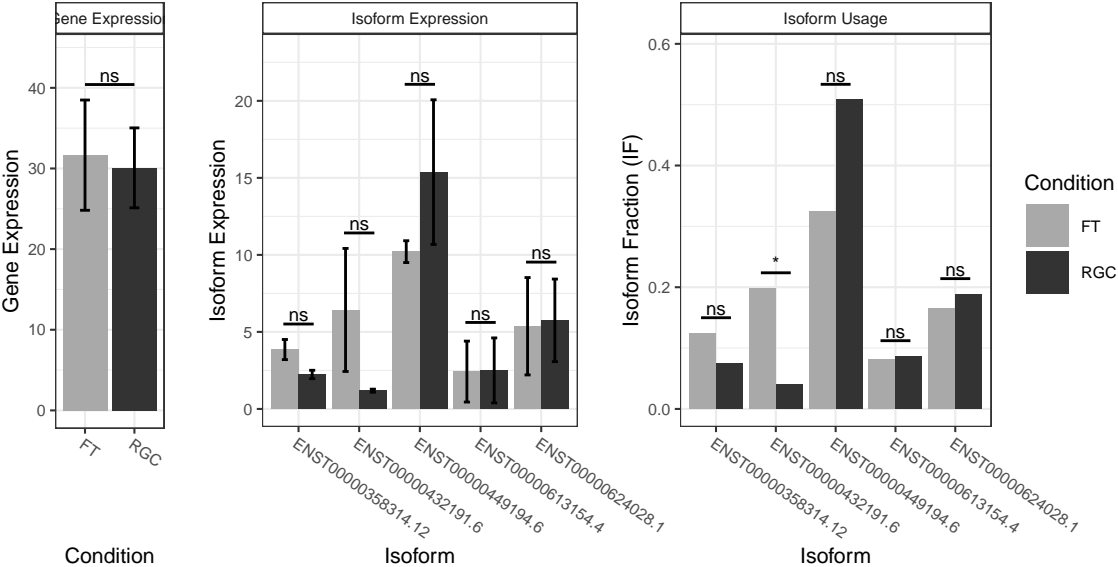
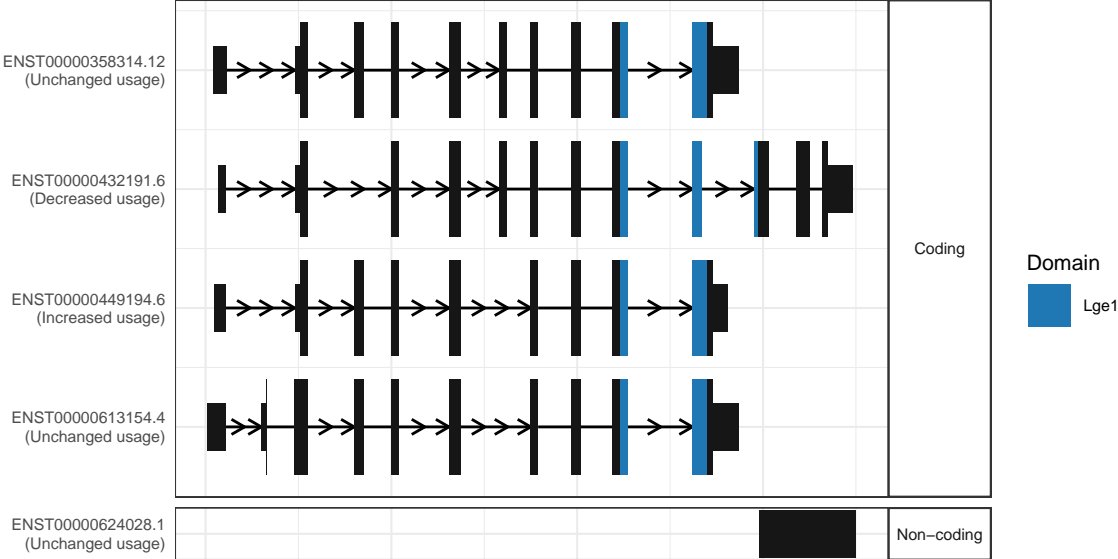


ATXN-1_C (x2)



AXH

The isoform switch in PPHLN1(FT vs RGC)

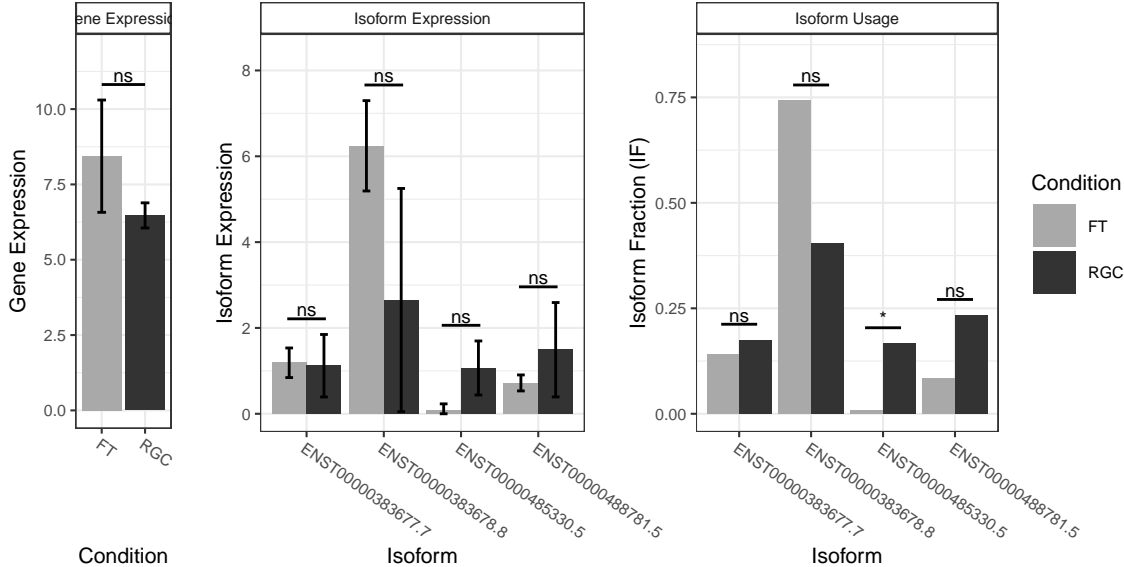
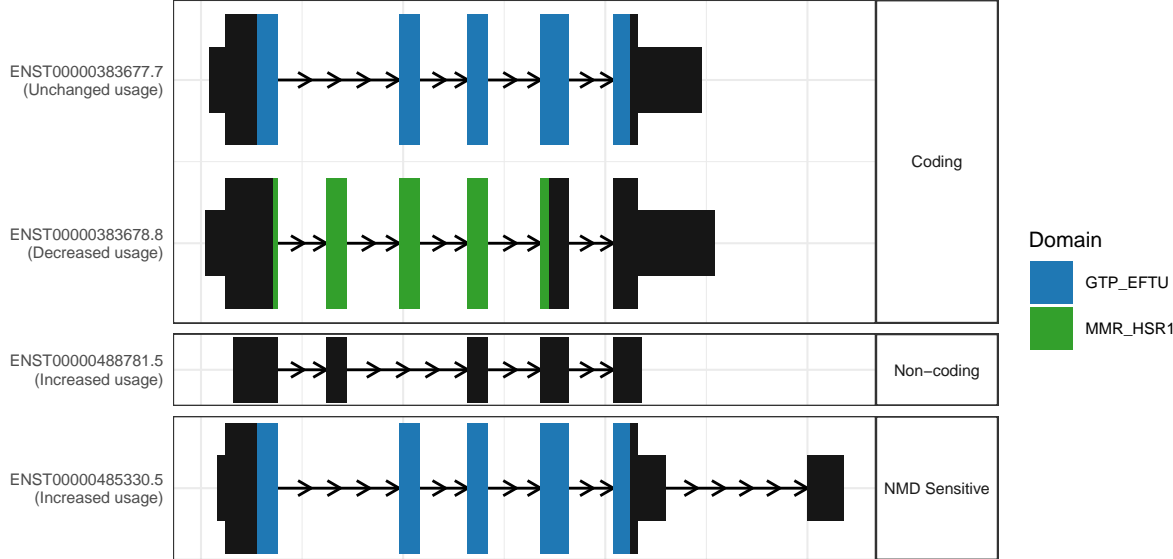


Domain



Lge1

The isoform switch in GTPBP8 (FT vs RGC)



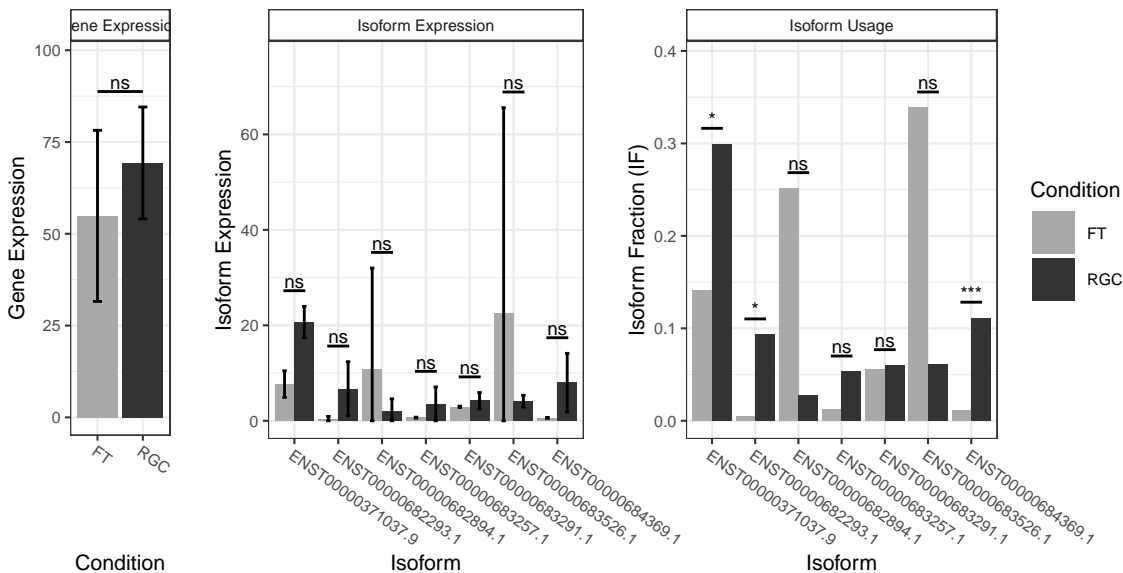
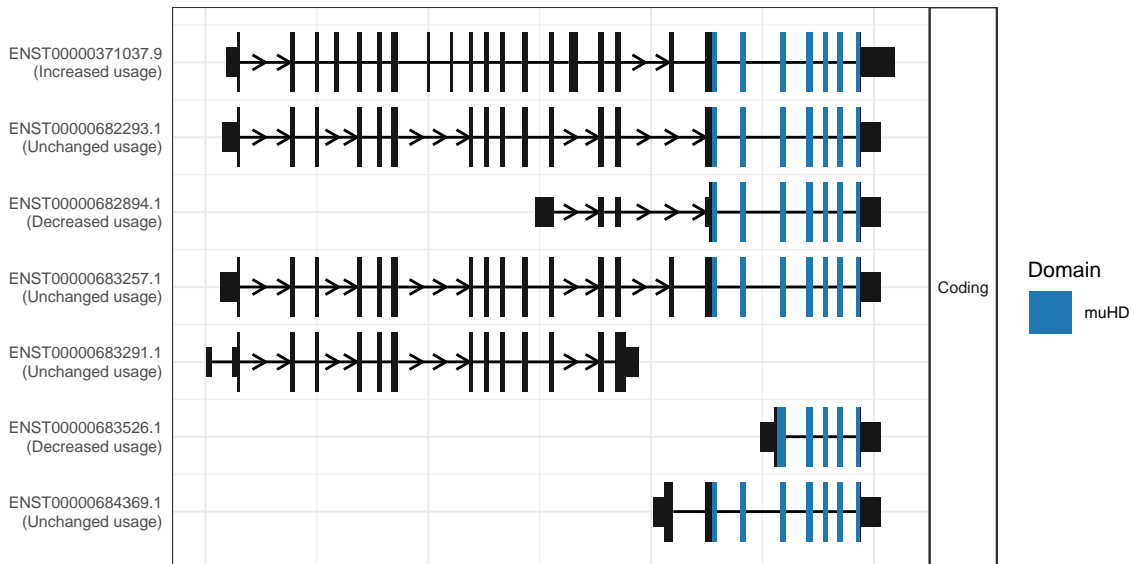
Domain



GTP_EFTU

MMR_HSR1

The isoform switch in SGIP1 (FT vs RGC)

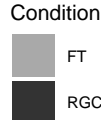
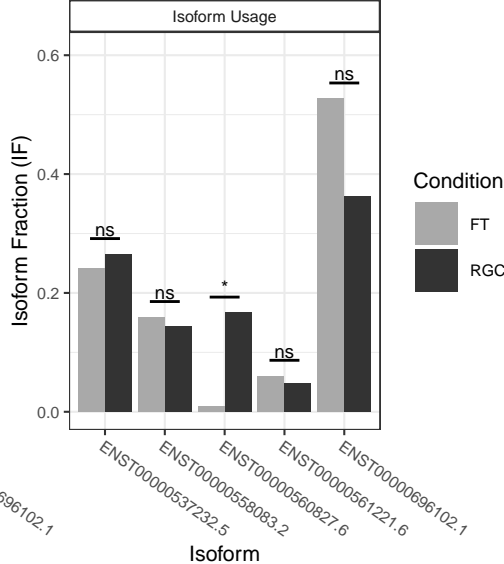
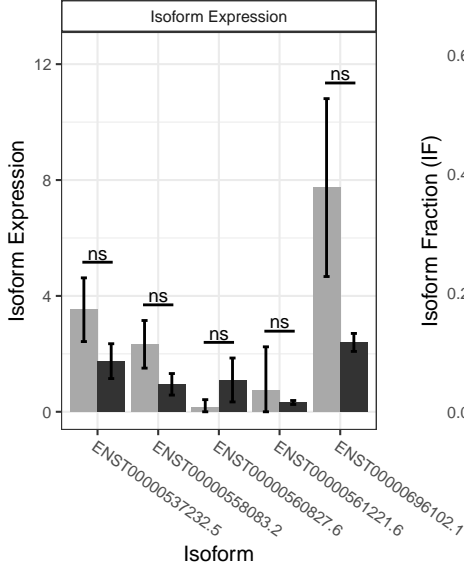
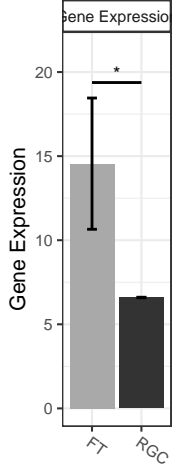
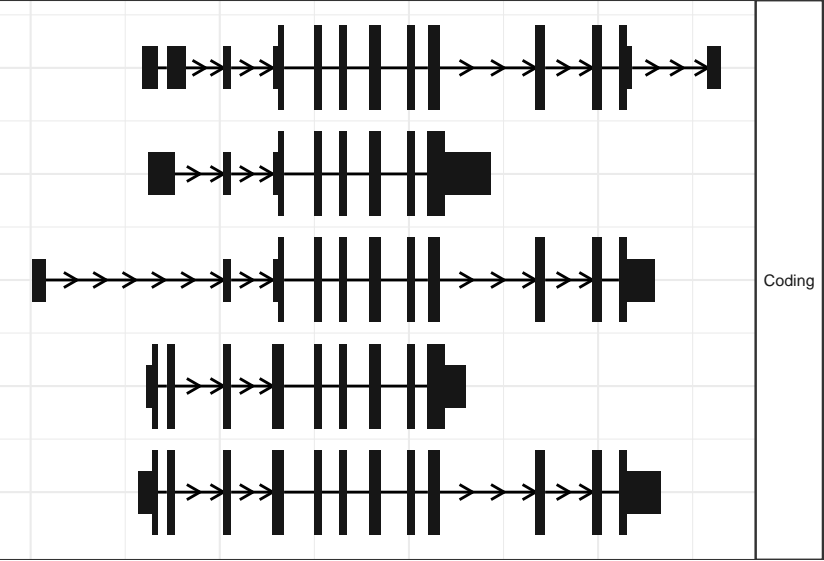


Domain

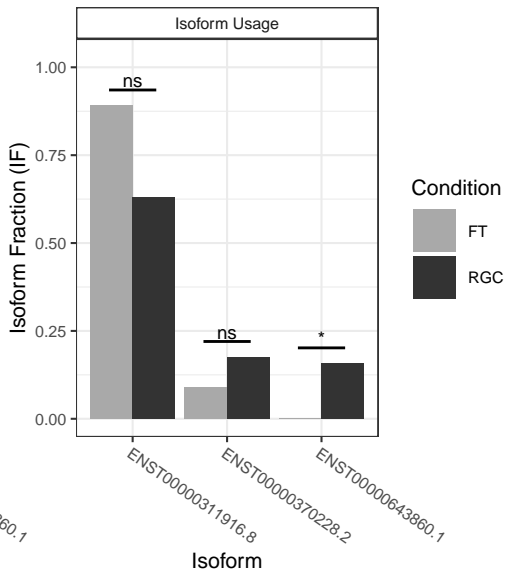
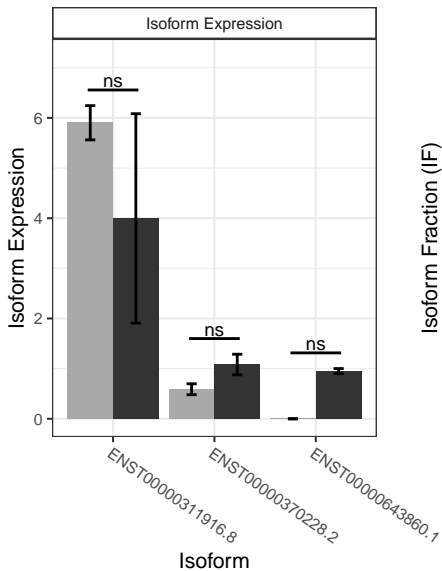
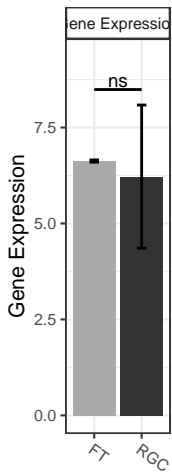
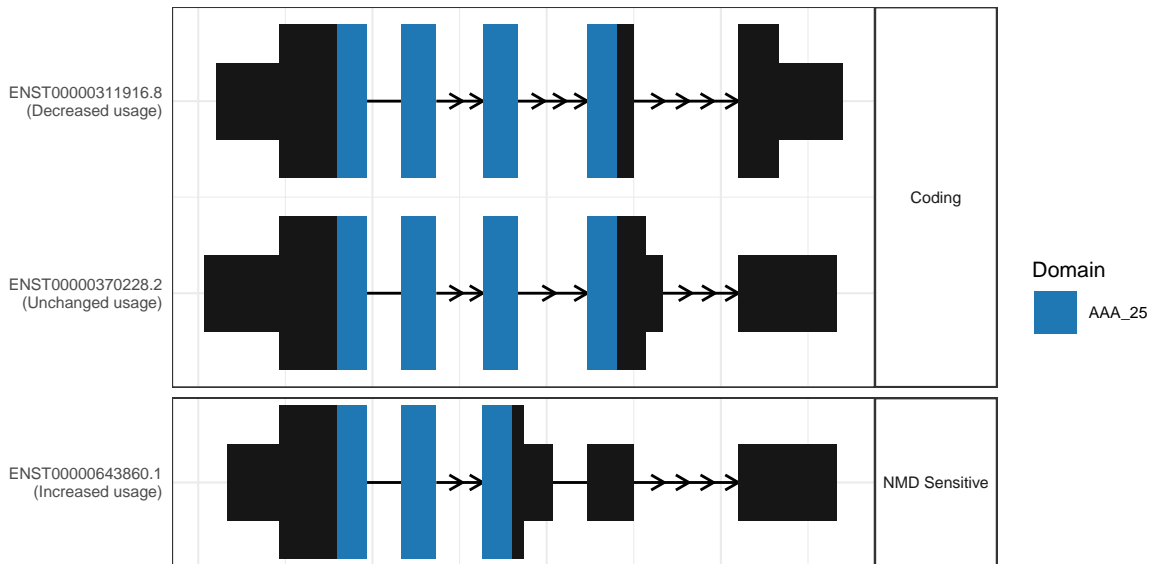


muHD

The isoform switch in TEX9 (FT vs RGC)



The isoform switch in TWNK (FT vs RGC)

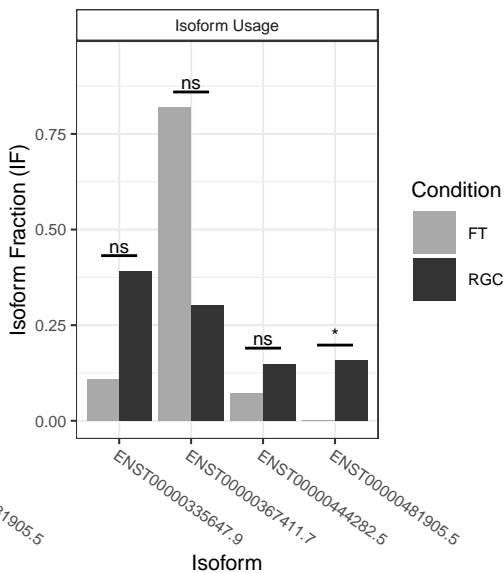
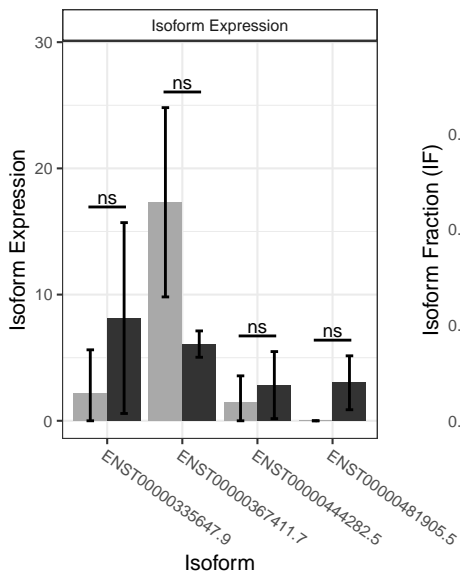
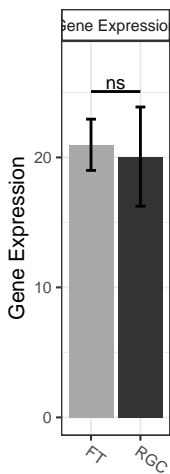
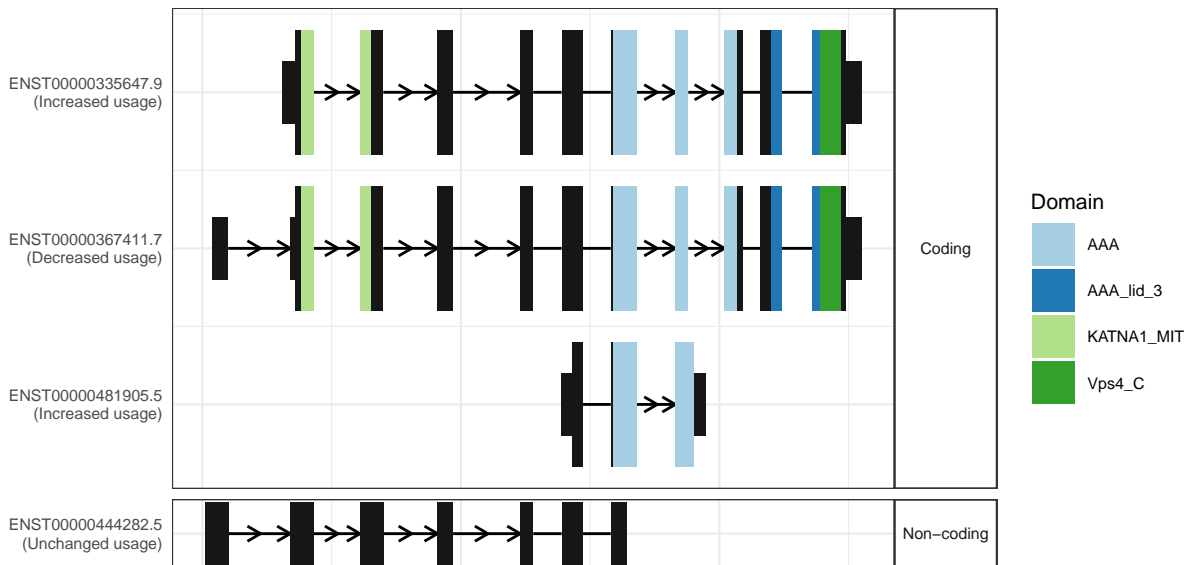


Domain



AAA_25

The isoform switch in KATNA1 (FT vs RGC)



Domain



AAA



AAA_lid_3

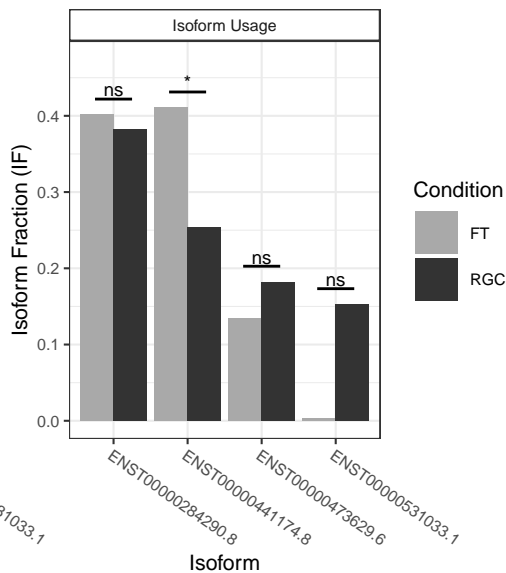
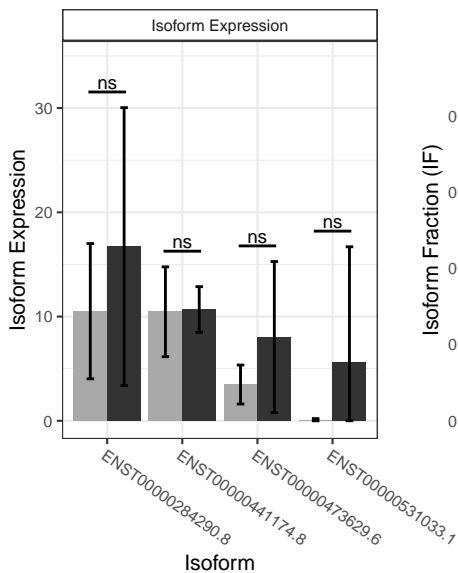
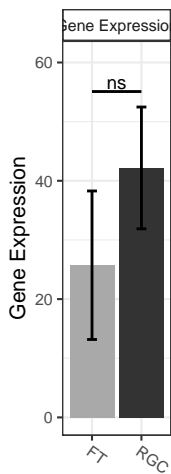
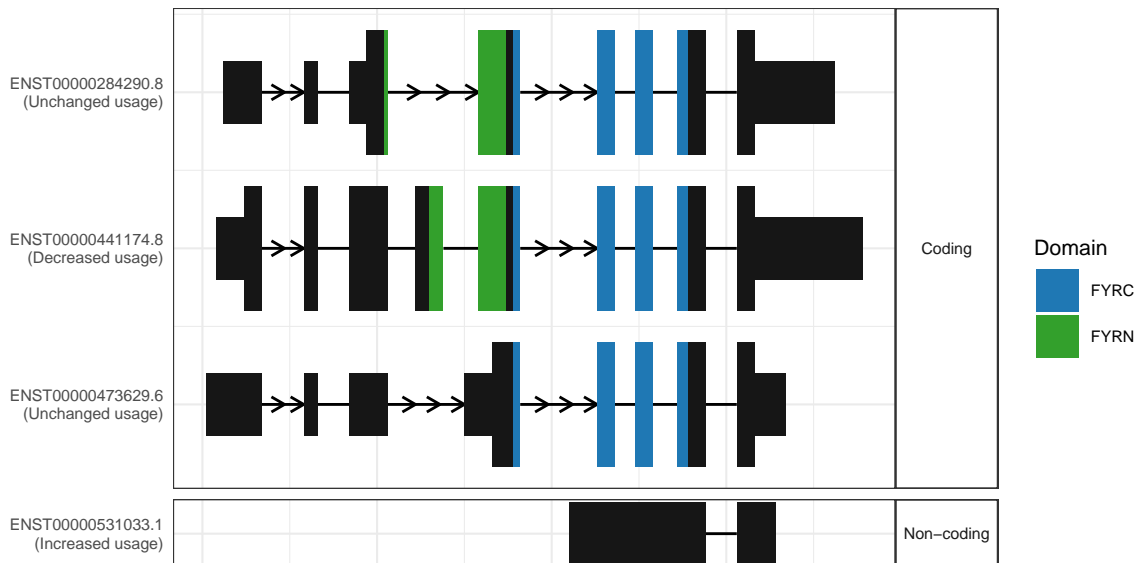


KATNA1_MIT



Vps4_C

The isoform switch in TBRG1 (FT vs RGC)



Domain

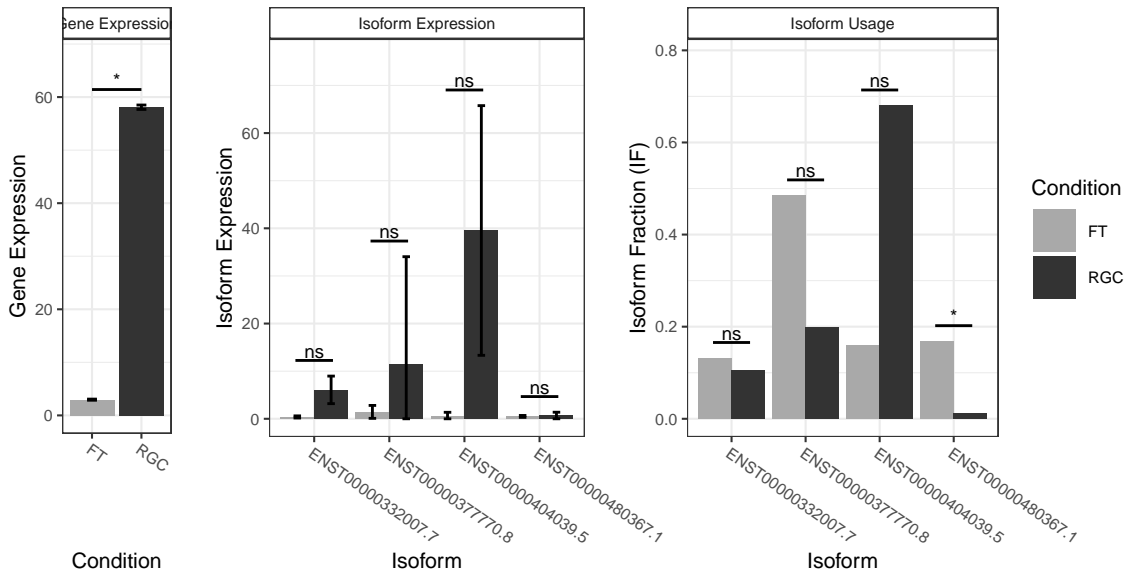
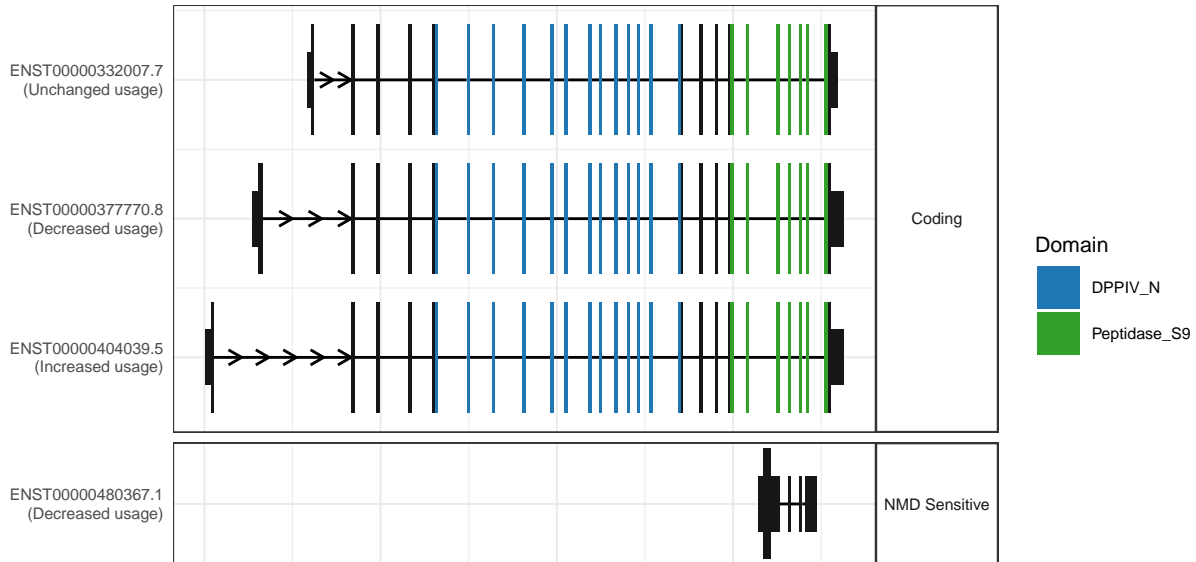


FYRC



FYRN

The isoform switch in DPP6 (FT vs RGC)



Domain

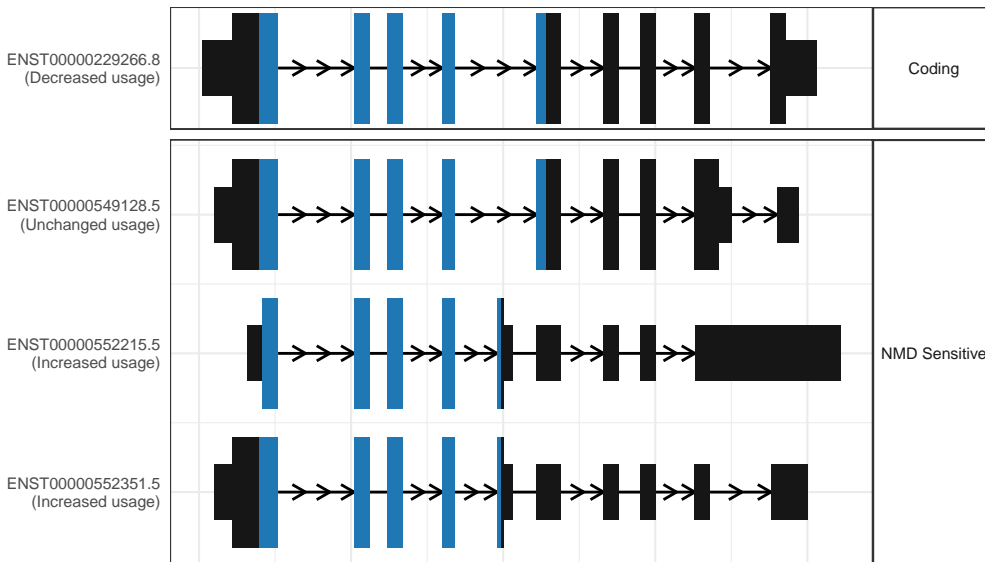


DPPIV_N



Peptidase_S9

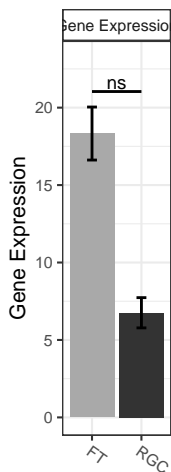
The isoform switch in CHPT1 (FT vs RGC)



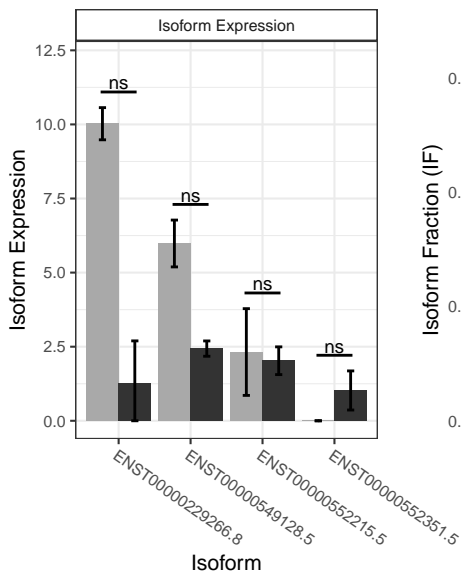
Domain

CDP-OH_P_transf

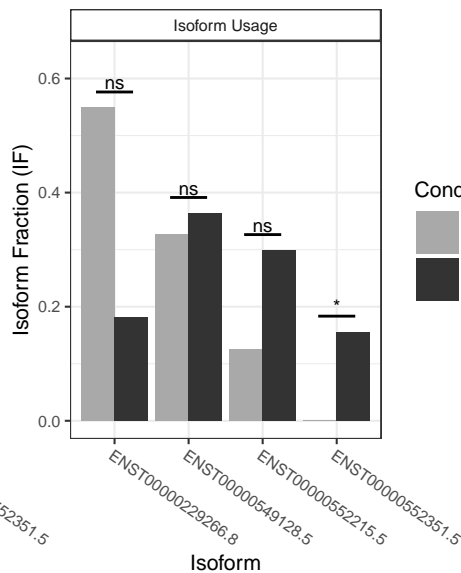
NMD Sensitive



Condition



Isoform



Isoform

Condition

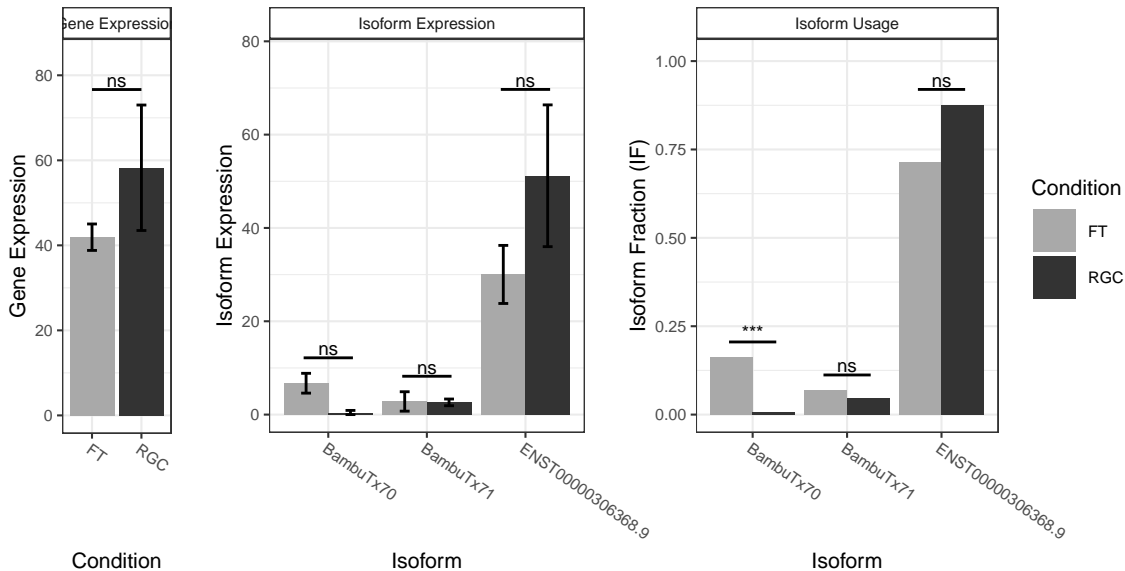
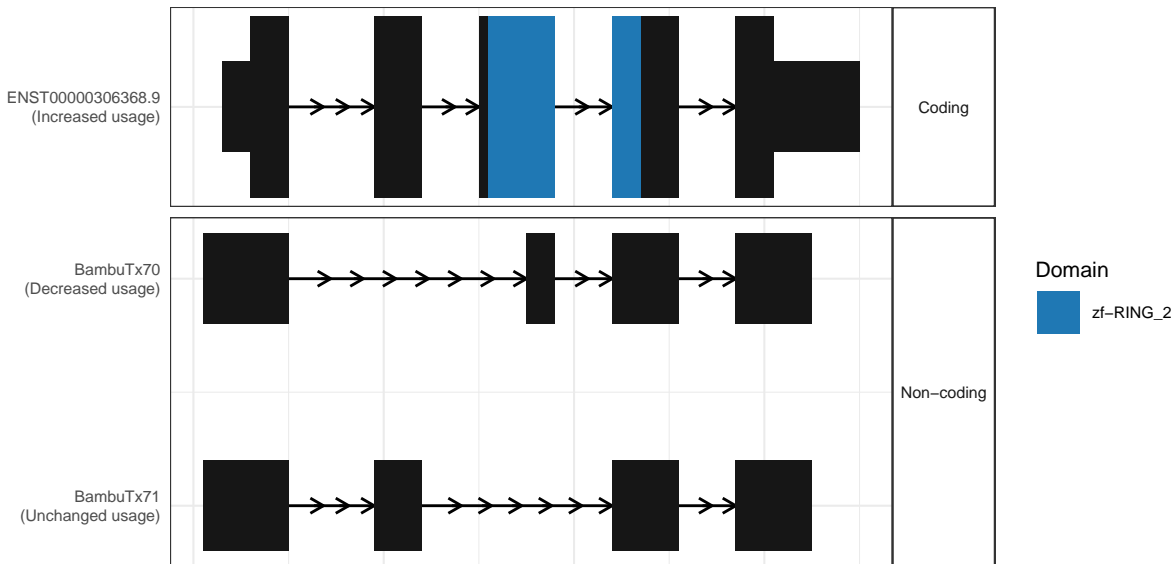
FT
RGC

Domain



CDP-OH_P_transf

The isoform switch in RNF181 (FT vs RGC)

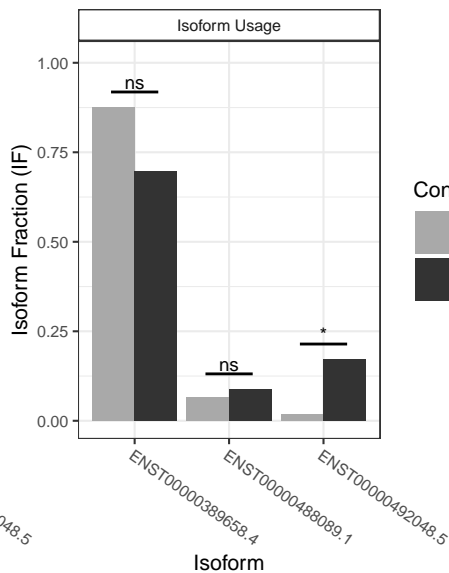
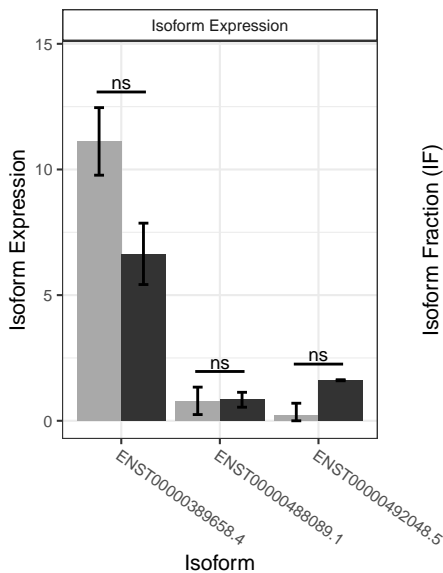
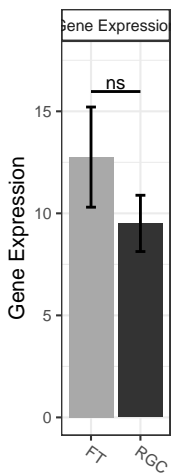
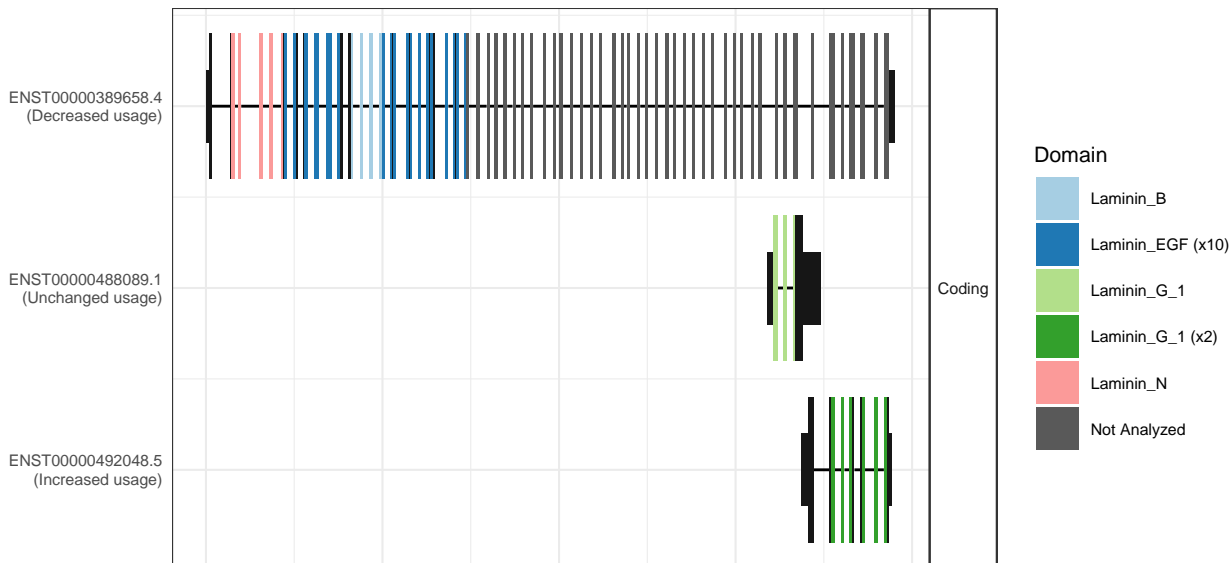


Domain









zf-RING_2

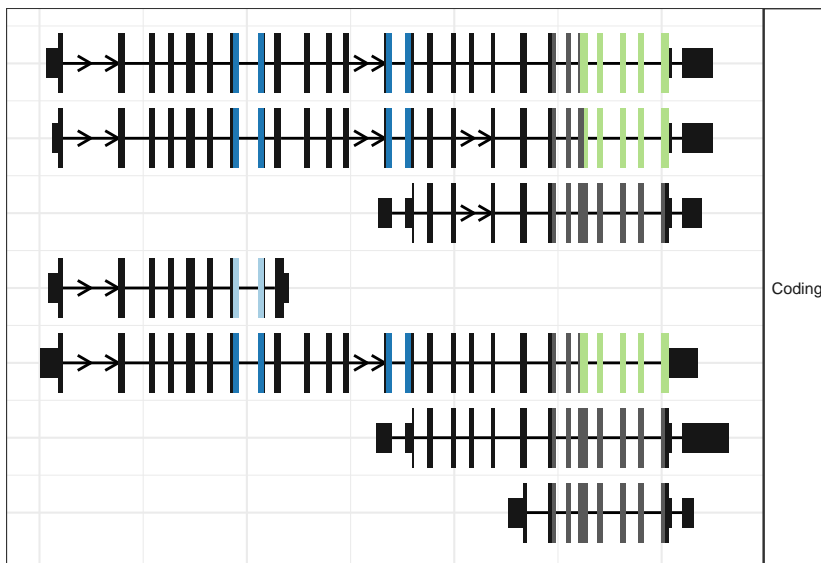
The isoform switch in LAMA1 (FT vs RGC)



Domain

	Laminin_B
	Laminin_EGF (x10)
	Laminin_G_1
	Laminin_G_1 (x2)
	Laminin_N
	Not Analyzed

The isoform switch in PTPRO (FT vs RGC)



Domain

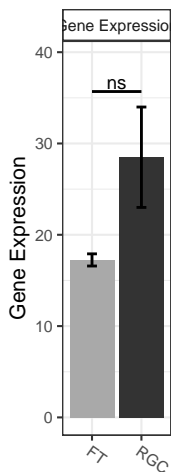
fn3

fn3 (x2)

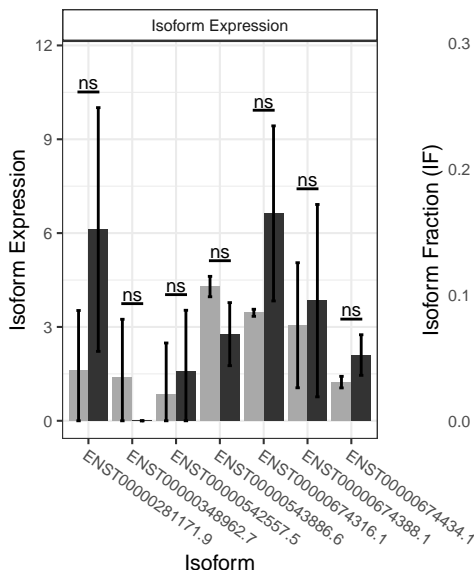
Y_phosphatase

Not Analyzed

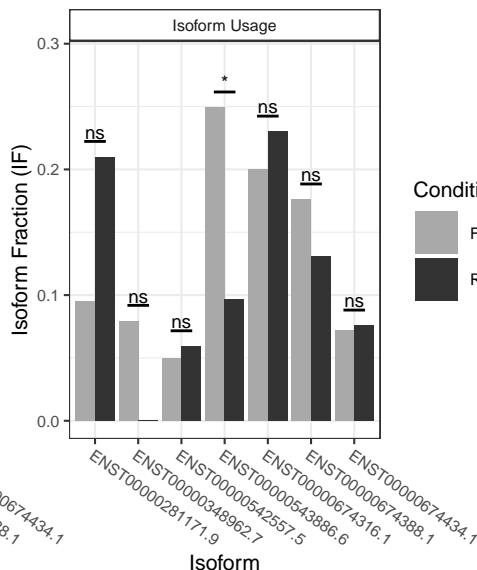
Coding



Condition



Isoform



Isoform

Condition

FT

RGC

Domain



fn3



fn3 (x2)

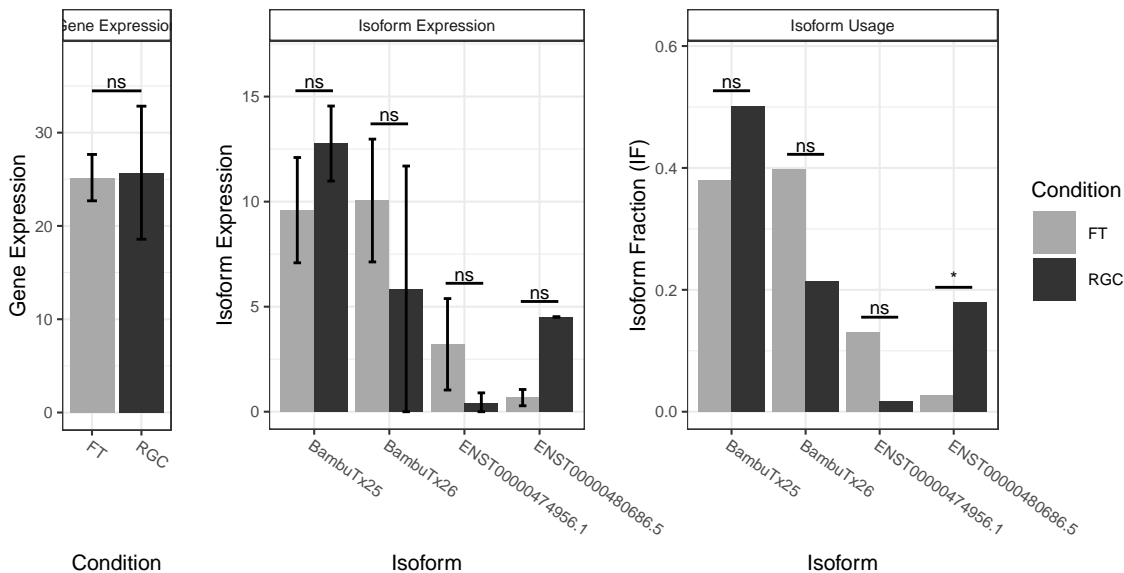
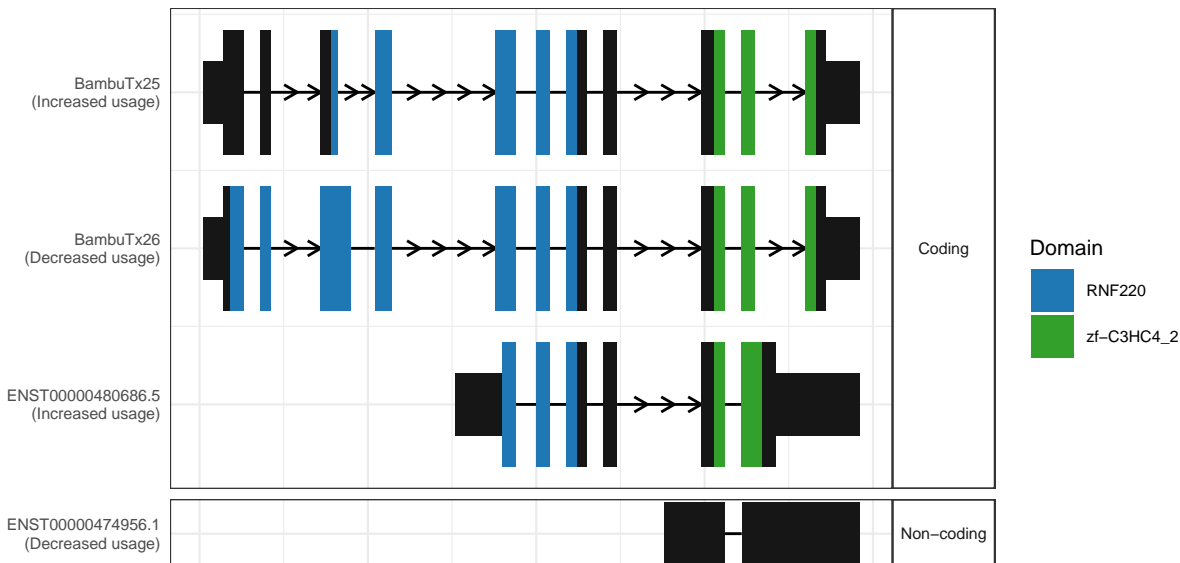


Y_phosphatase



Not Analyzed

The isoform switch in RNF220 (FT vs RGC)



Domain

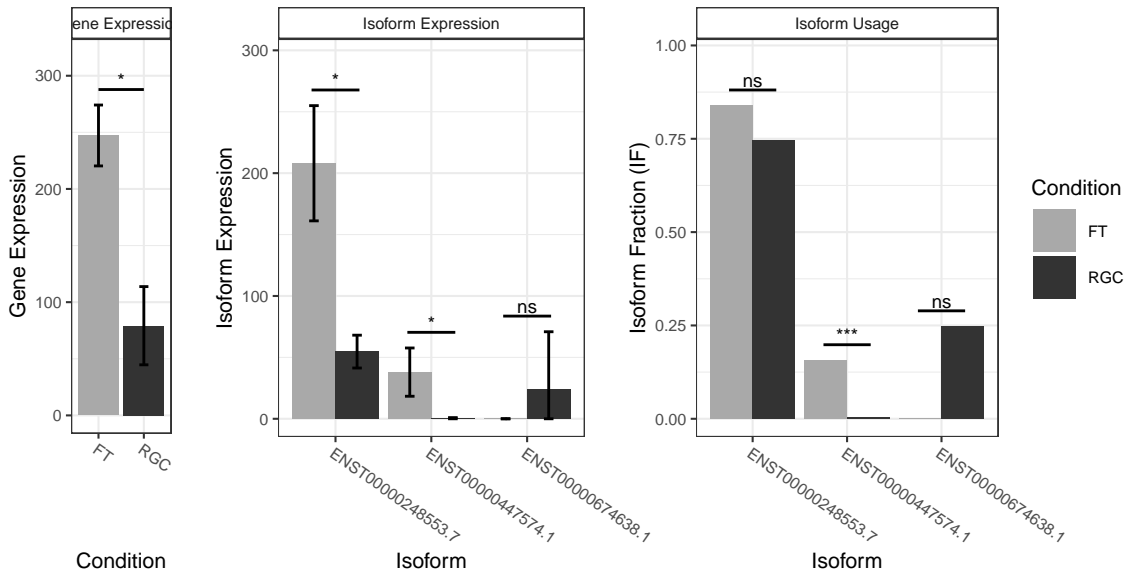
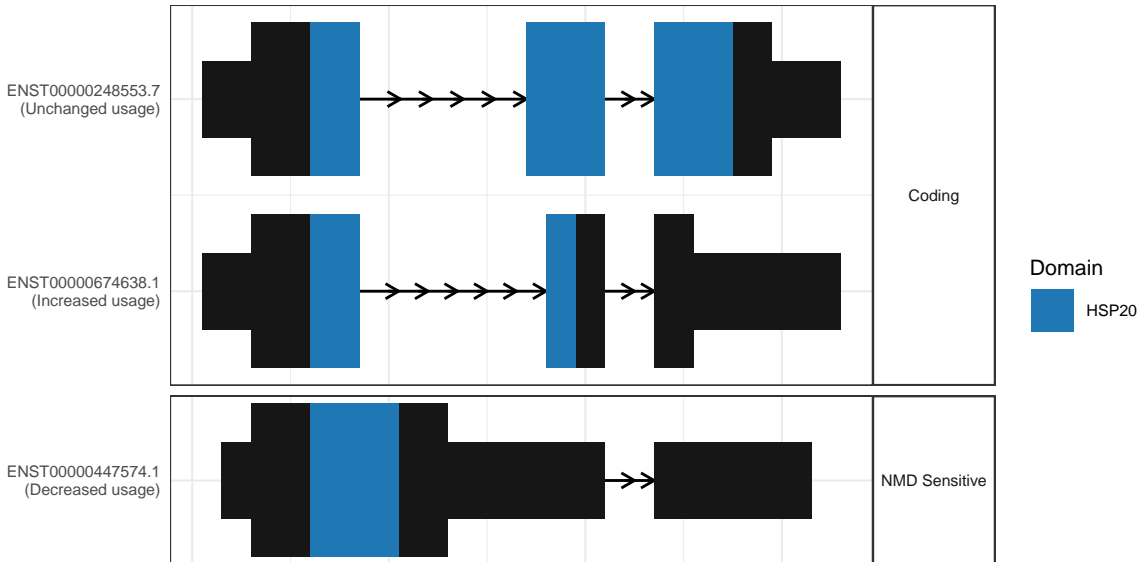


RNF220



zf-C3HC4_2

The isoform switch in HSPB1 (FT vs RGC)

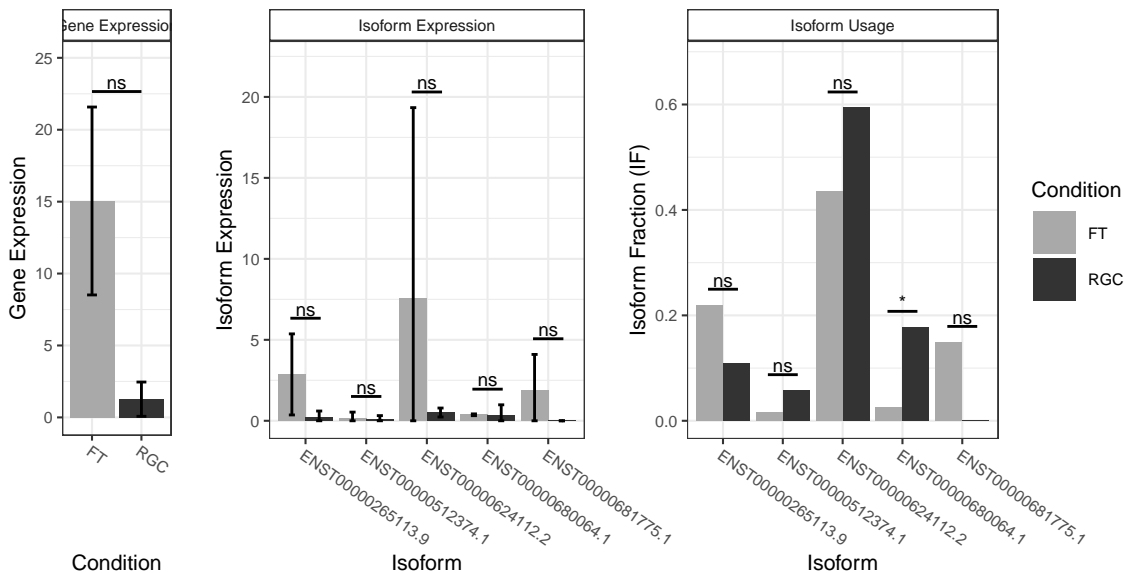
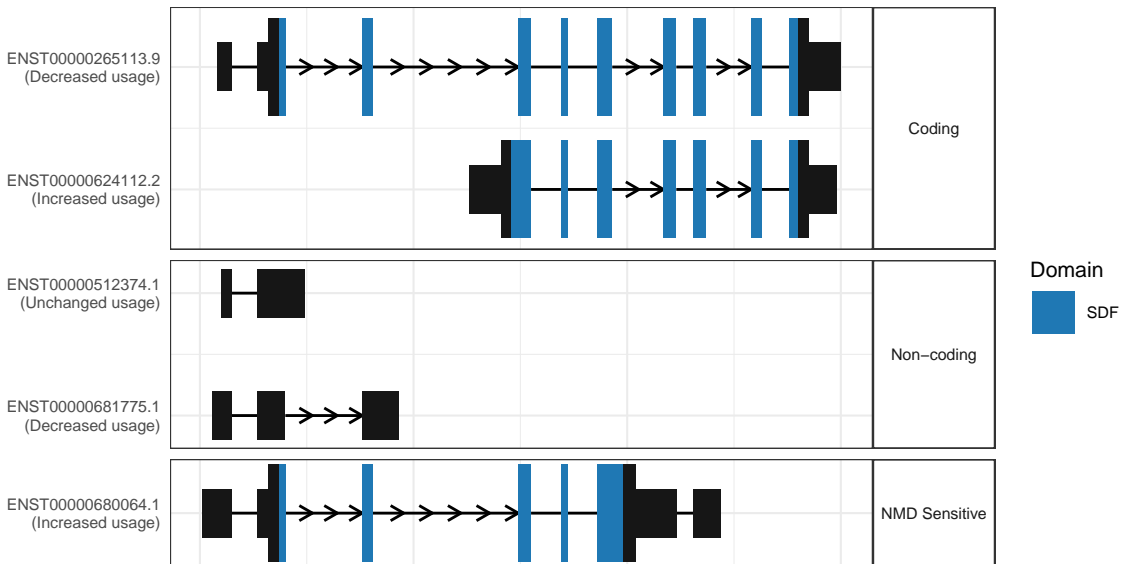


Domain



HSP20

The isoform switch in SLC1A3 (FT vs RGC)

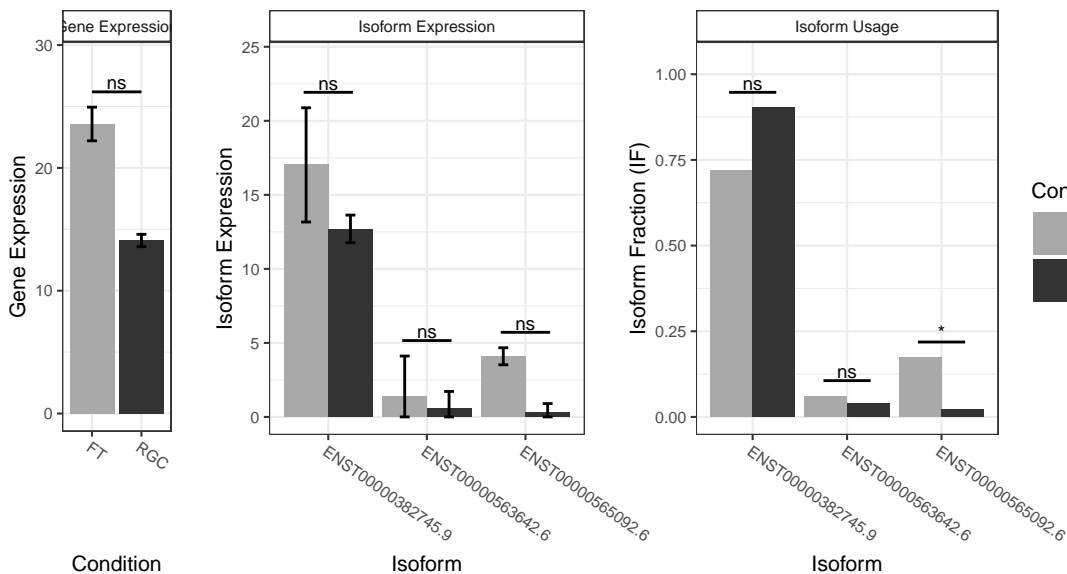
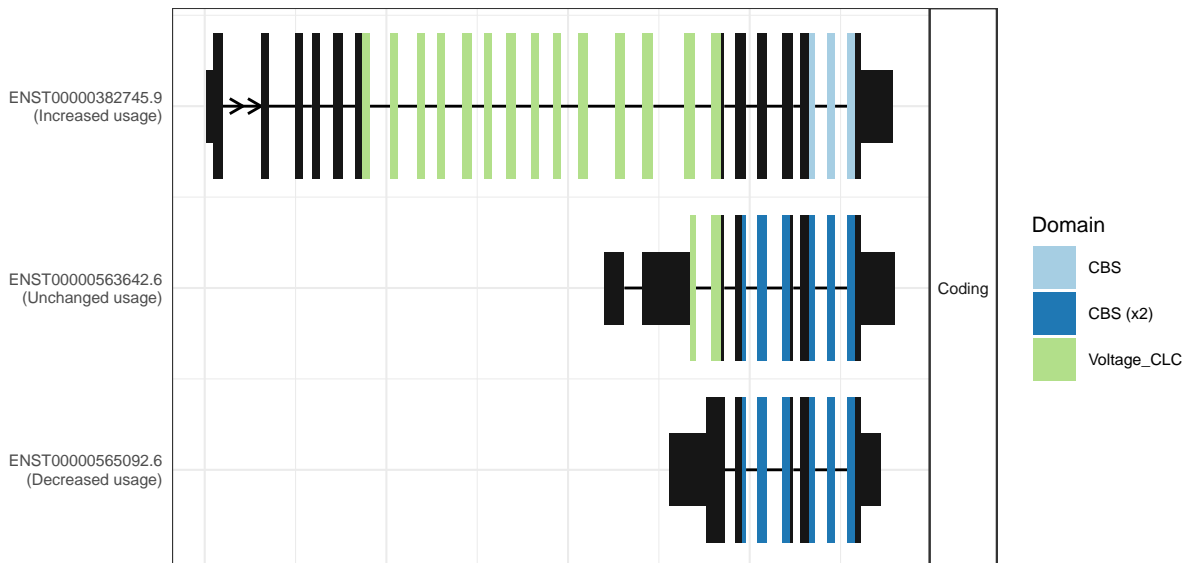


Domain



SDF

The isoform switch in CLCN7 (FT vs RGC)



Domain



CBS

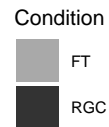
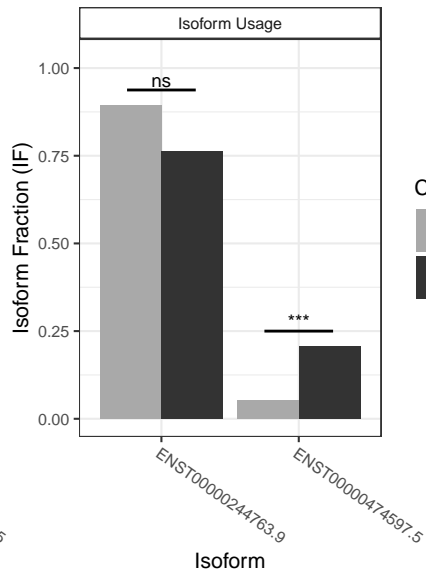
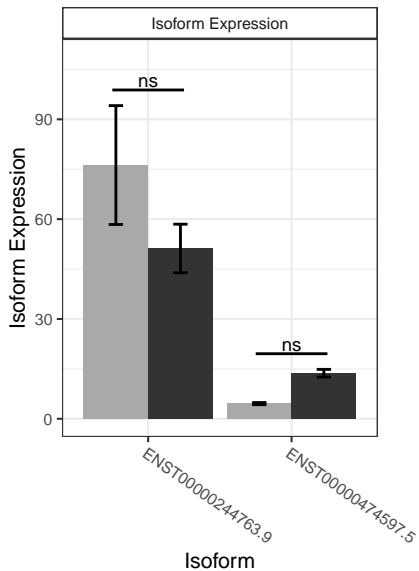
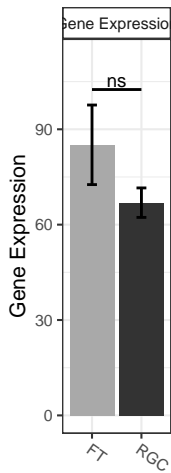
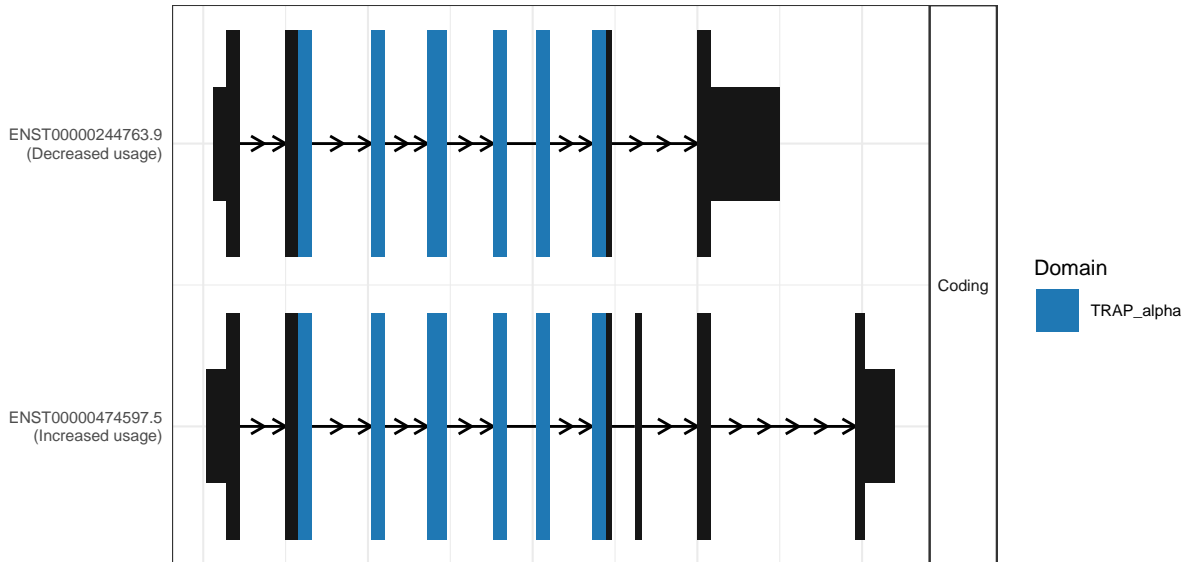


CBS (x2)



Voltage_CLC

The isoform switch in SSR1 (FT vs RGC)

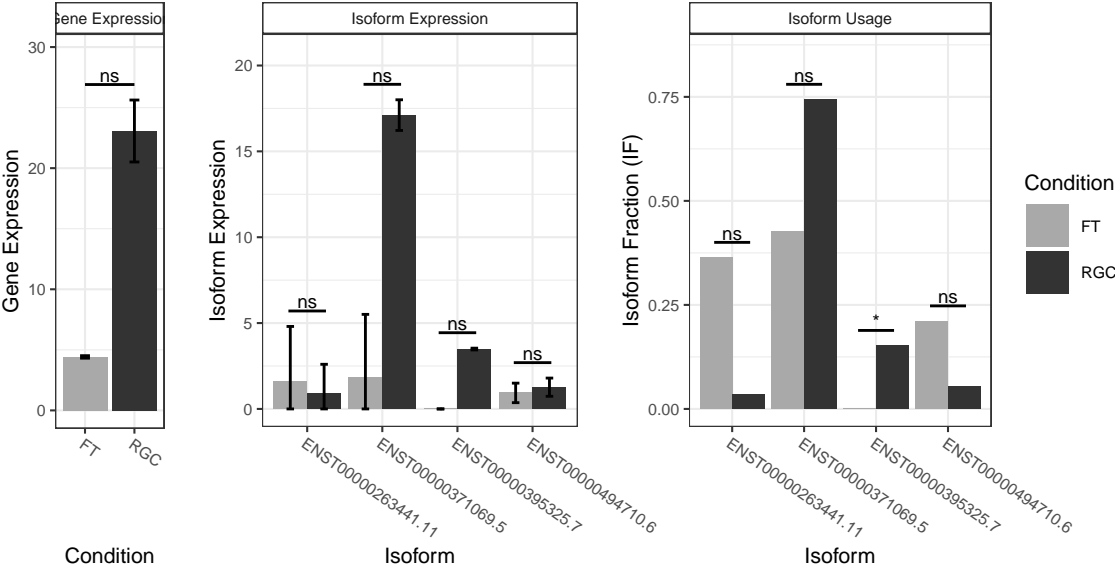
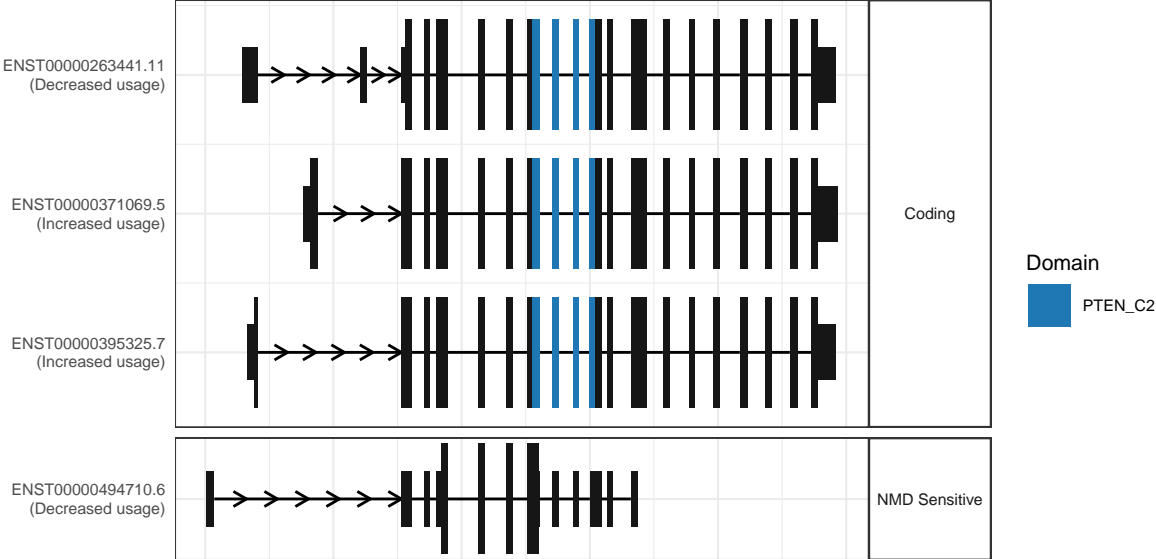


Domain



TRAP_alpha

The isoform switch in DNAJC6 (FT vs RGC)

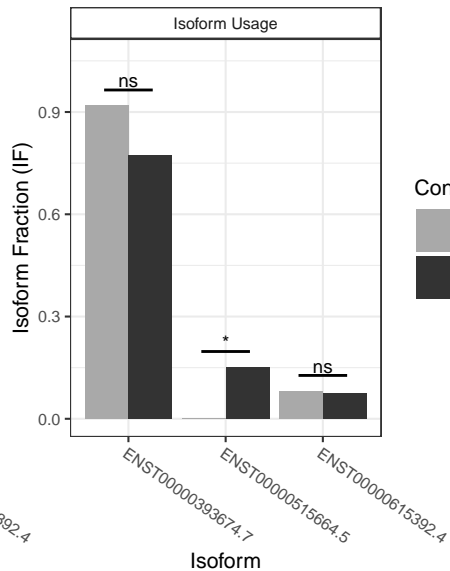
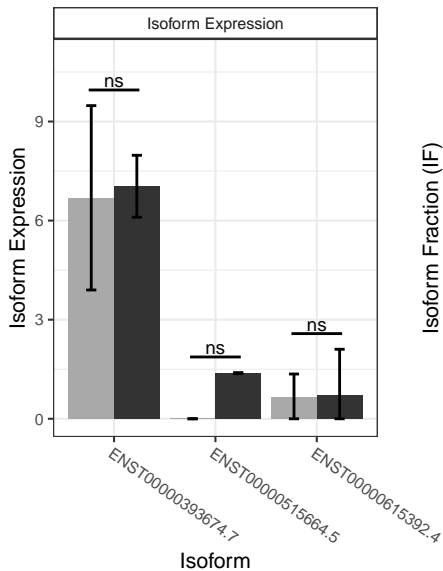
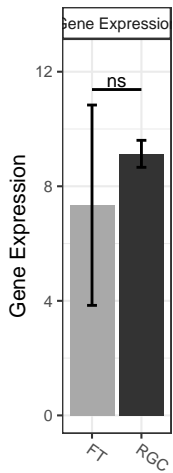
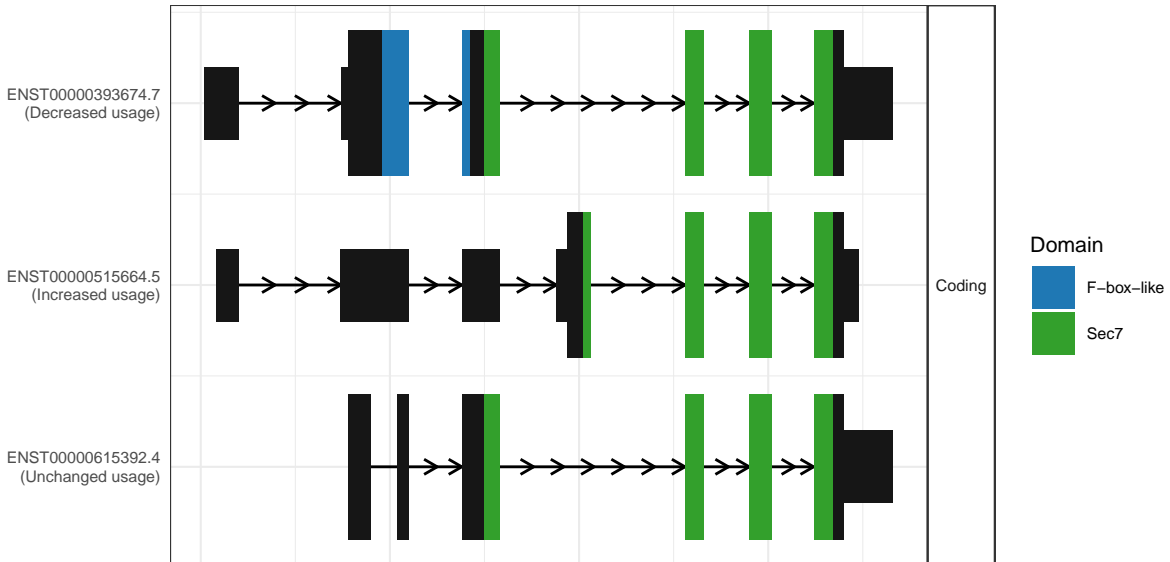


Domain



PTEN_C2

The isoform switch in FBXO8 (FT vs RGC)



Domain

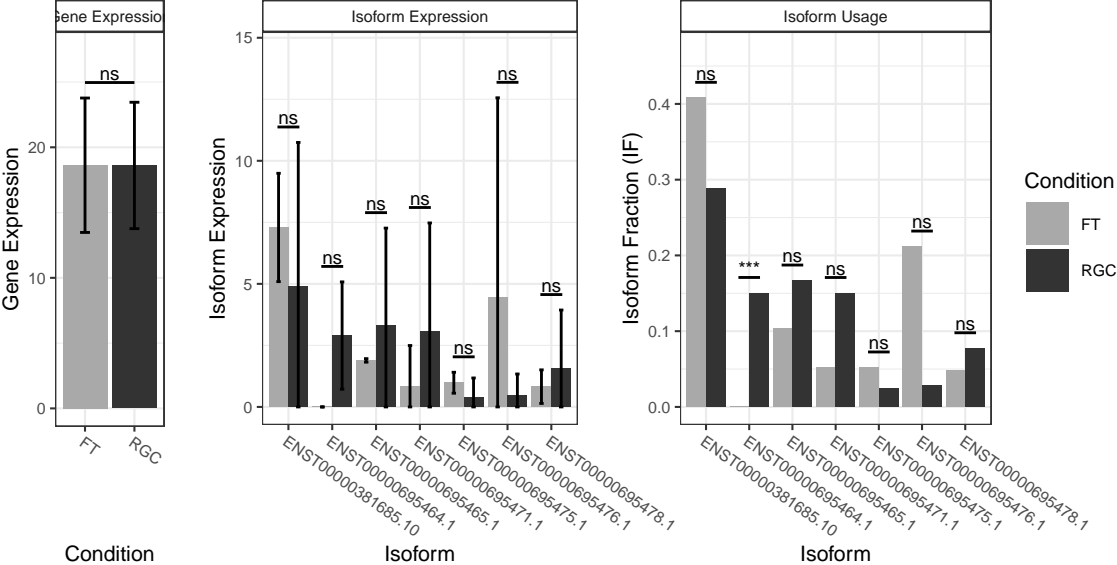
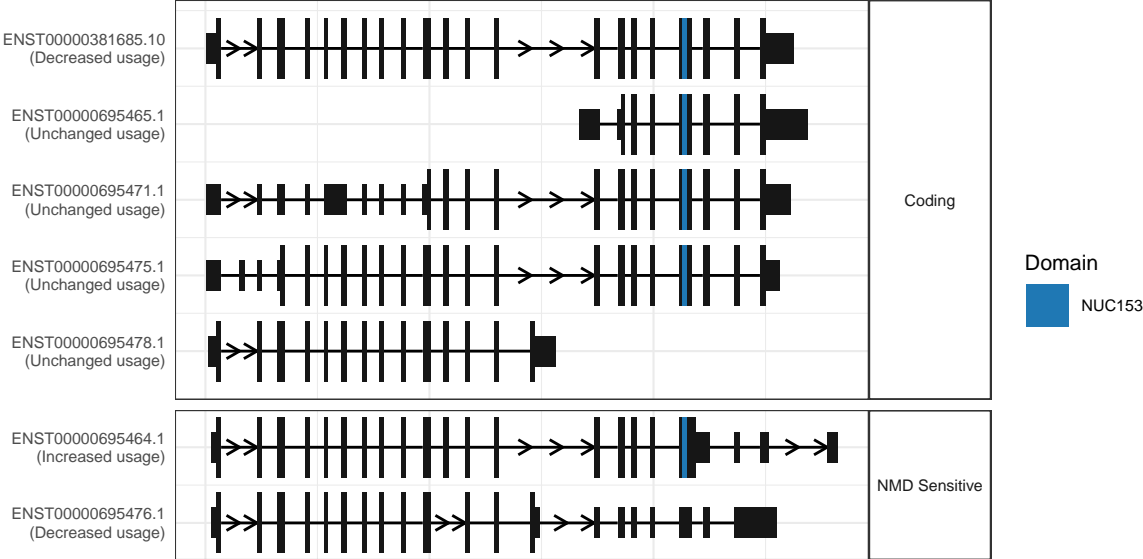


F-box-like



Sec7

The isoform switch in NOL10 (FT vs RGC)

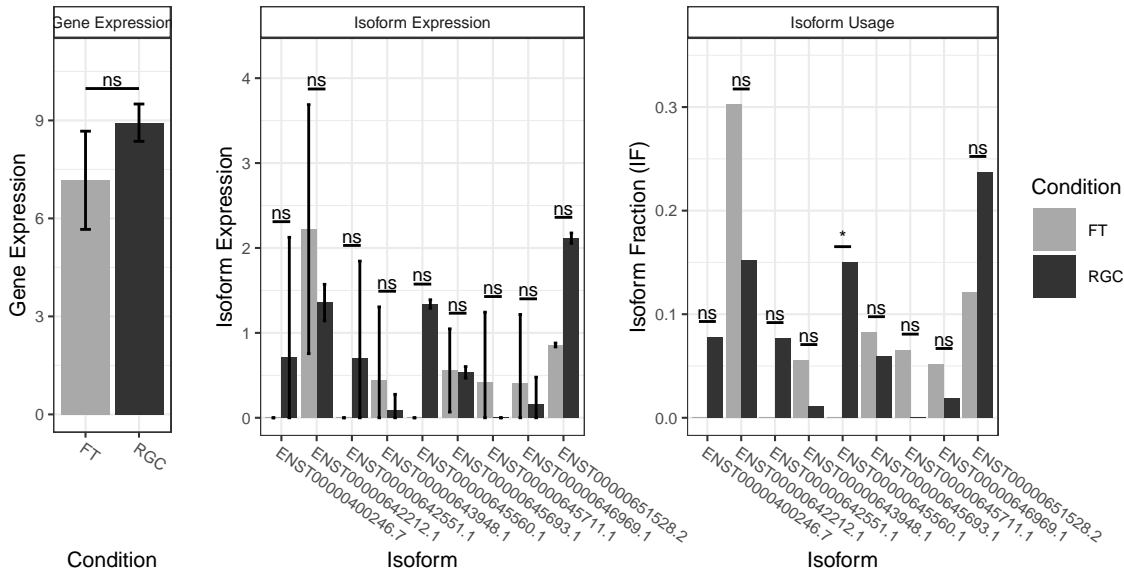
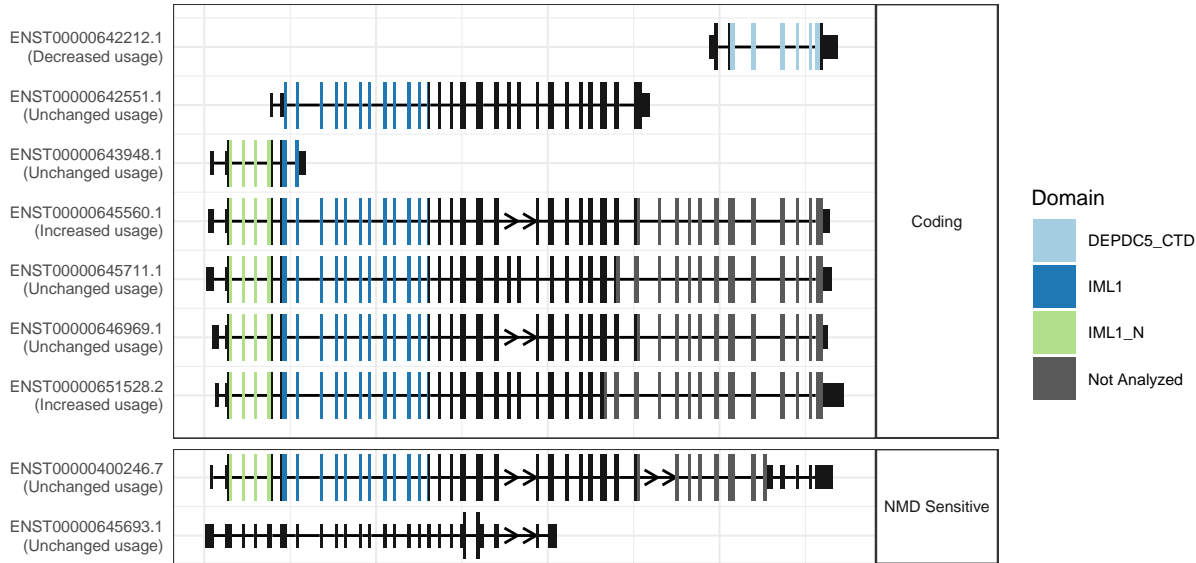


Domain

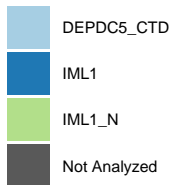


NUC153

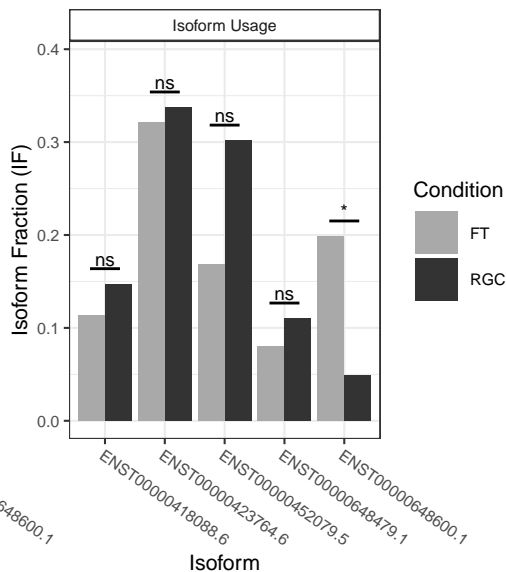
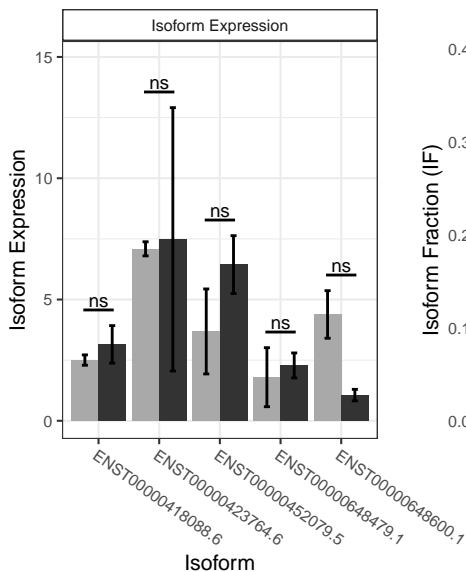
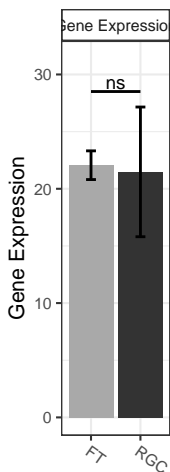
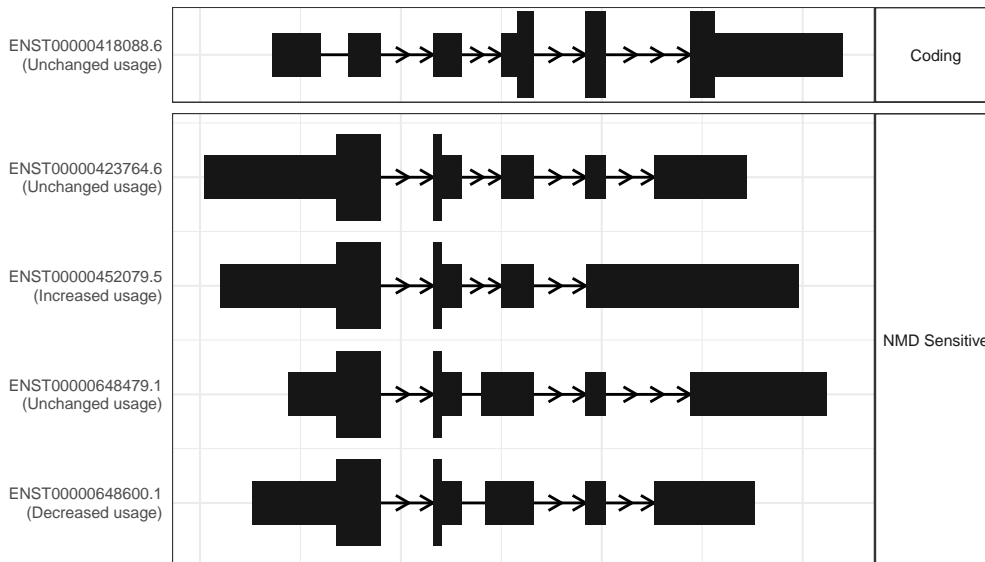
The isoform switch in DEPDC5 (FT vs RGC)



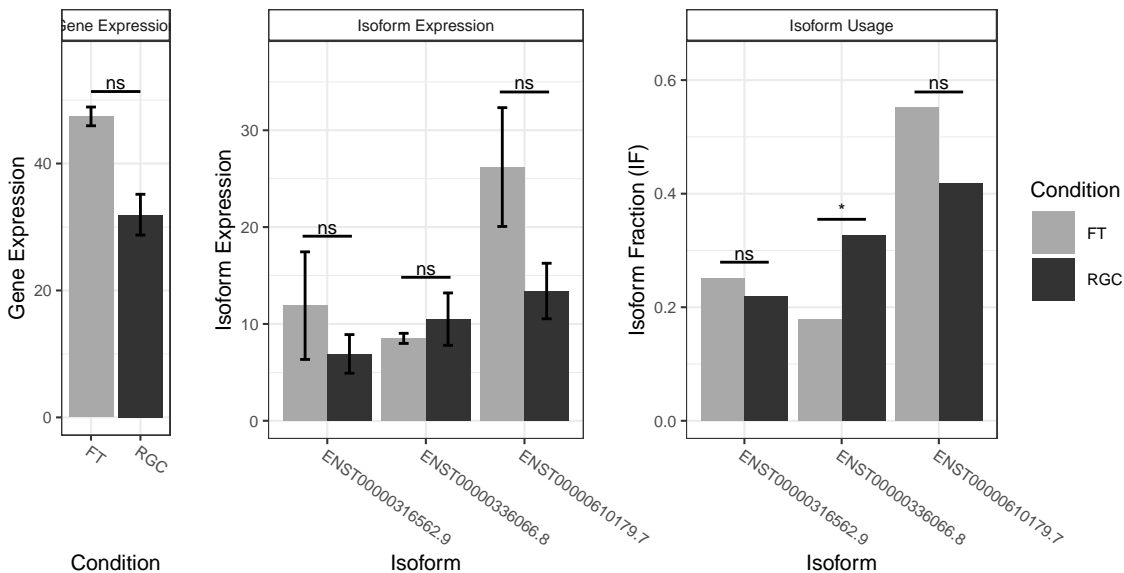
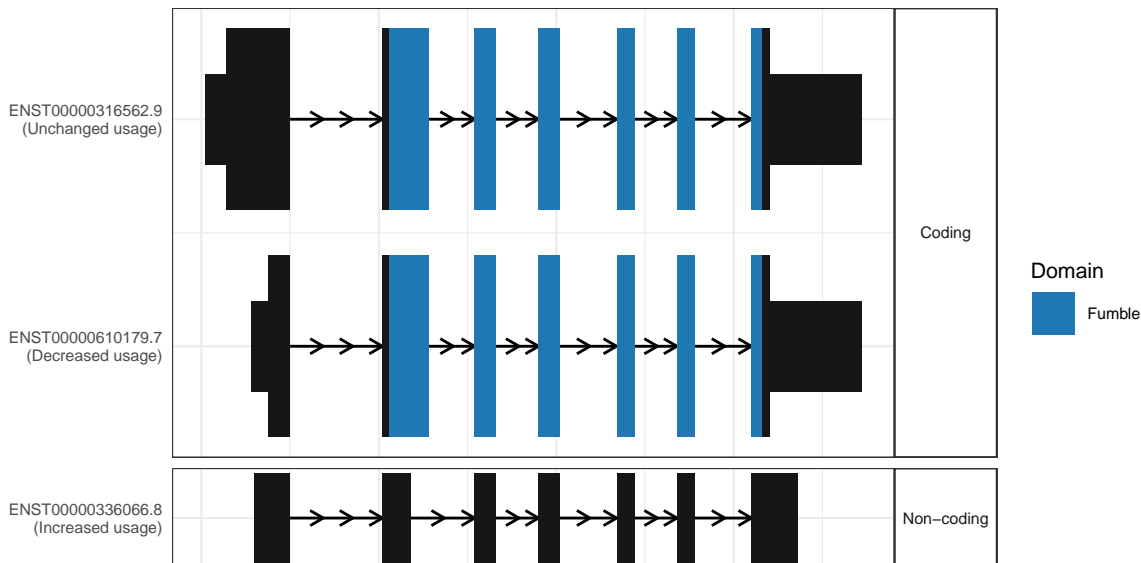
Domain



The isoform switch in GFOD3P (FT vs RGC)



The isoform switch in PANK2 (FT vs RGC)

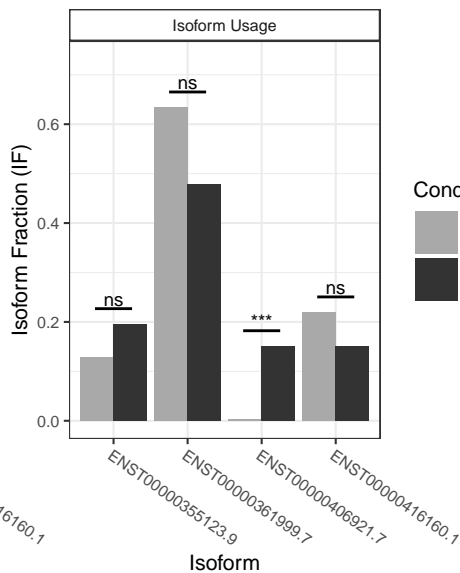
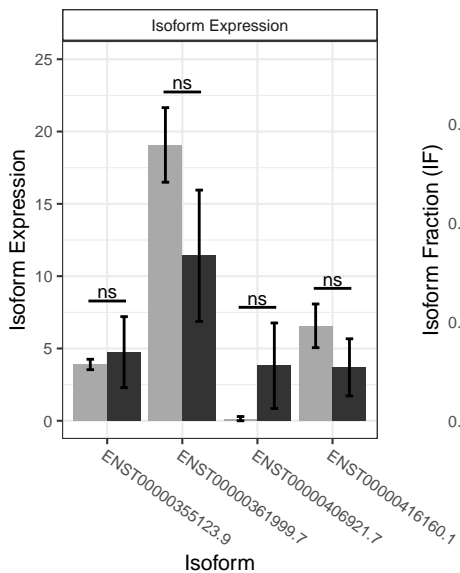
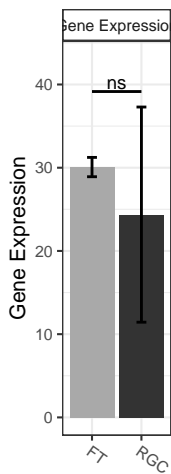
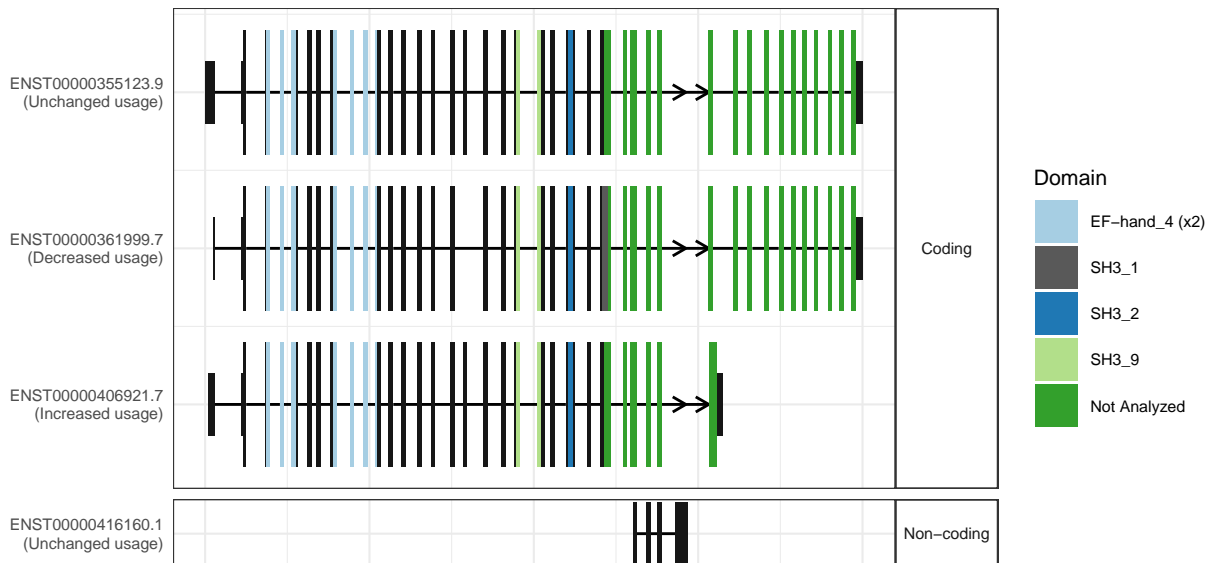


Domain

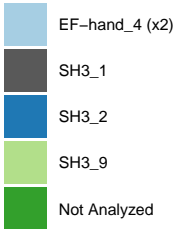


Fumble

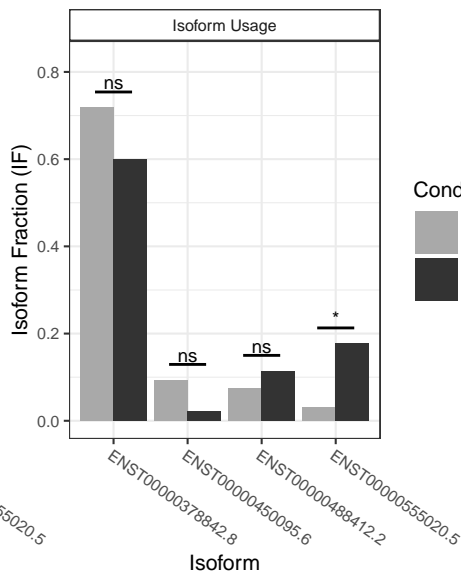
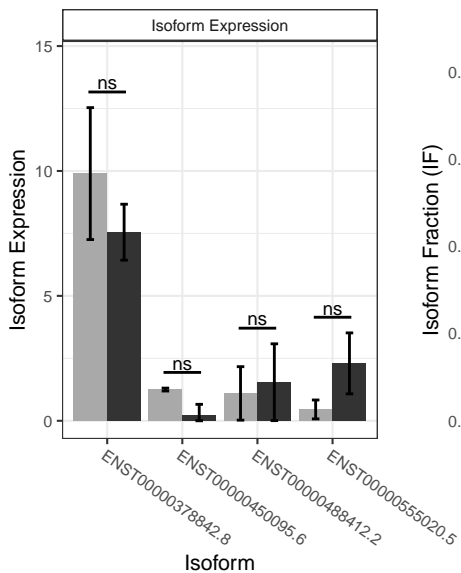
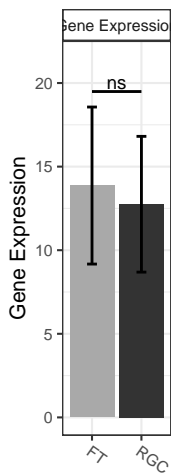
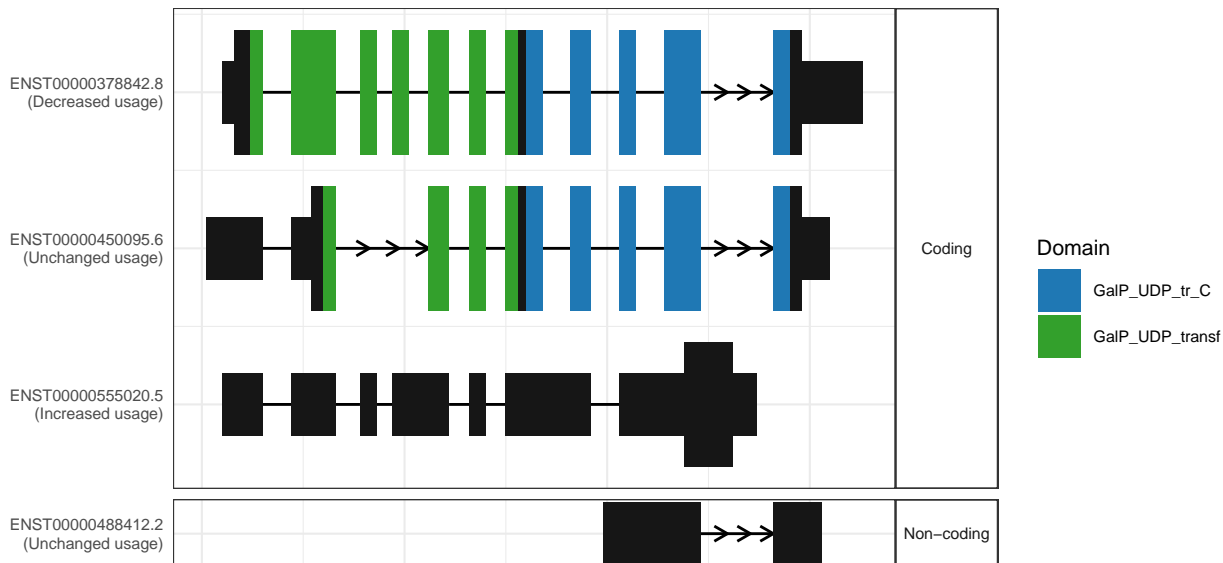
The isoform switch in ITSN2 (FT vs RGC)



Domain



The isoform switch in GALT (FT vs RGC)



Condition

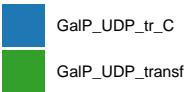
Isoform

Isoform

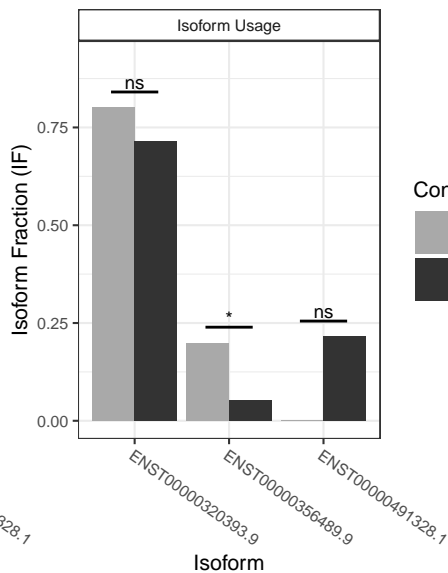
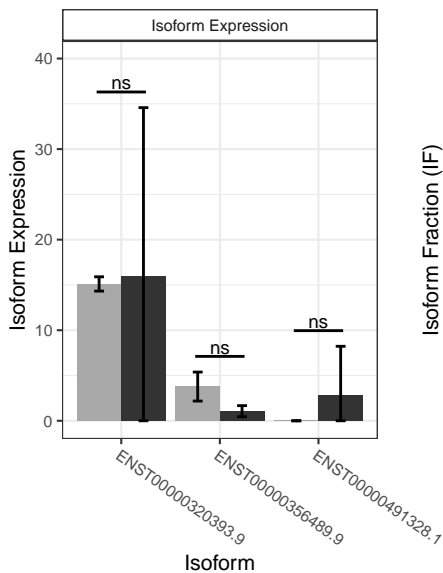
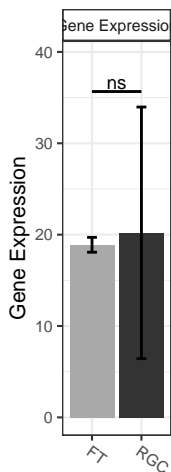
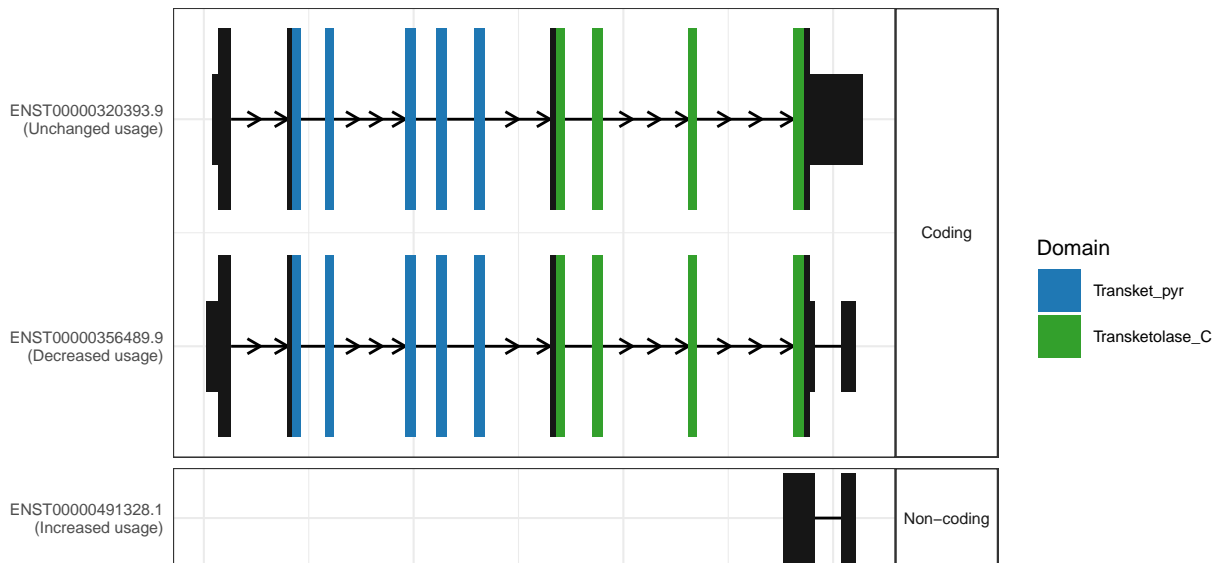
Condition



Domain



The isoform switch in BCKDHB (FT vs RGC)



Domain

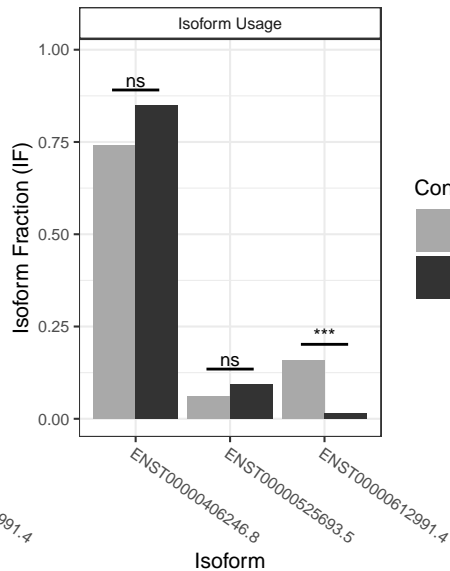
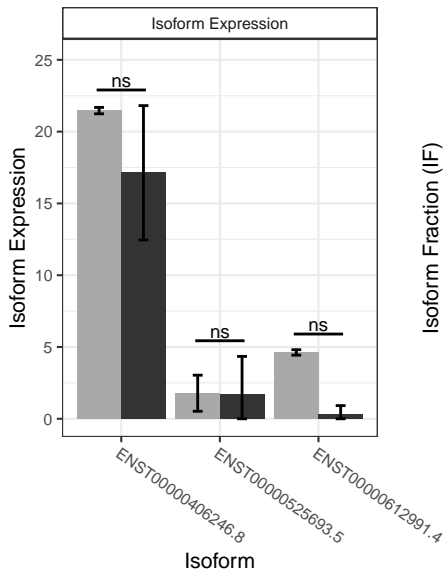
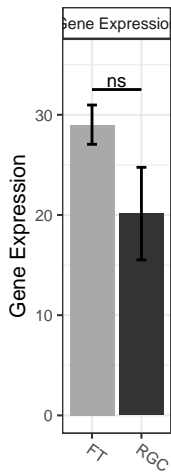
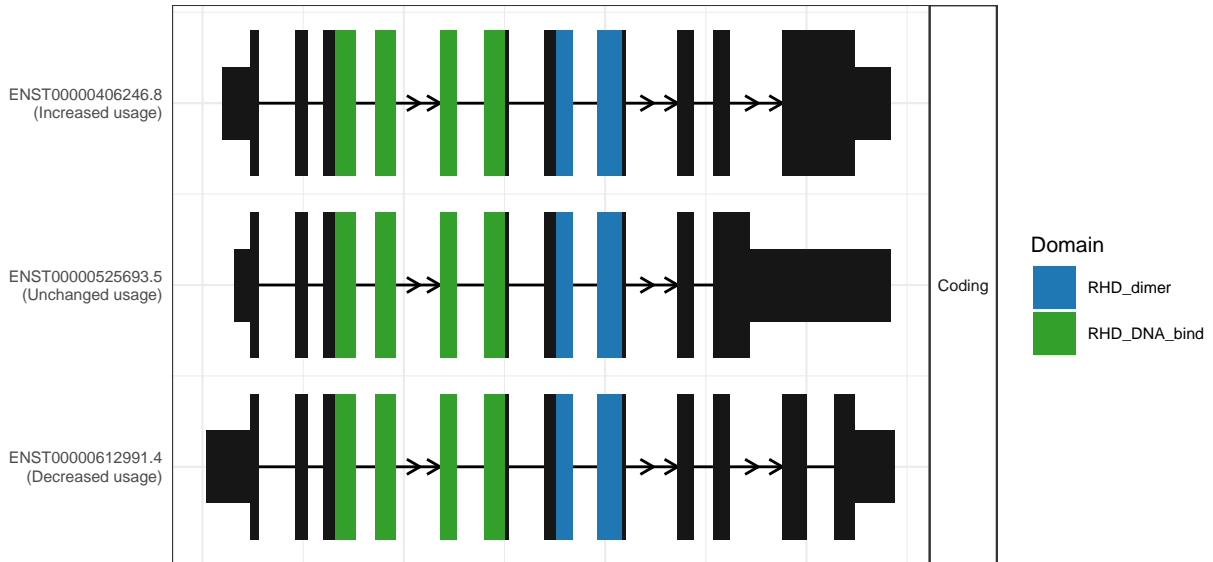


Transket_pyr



Transketolase_C

The isoform switch in RELA (FT vs RGC)



Domain

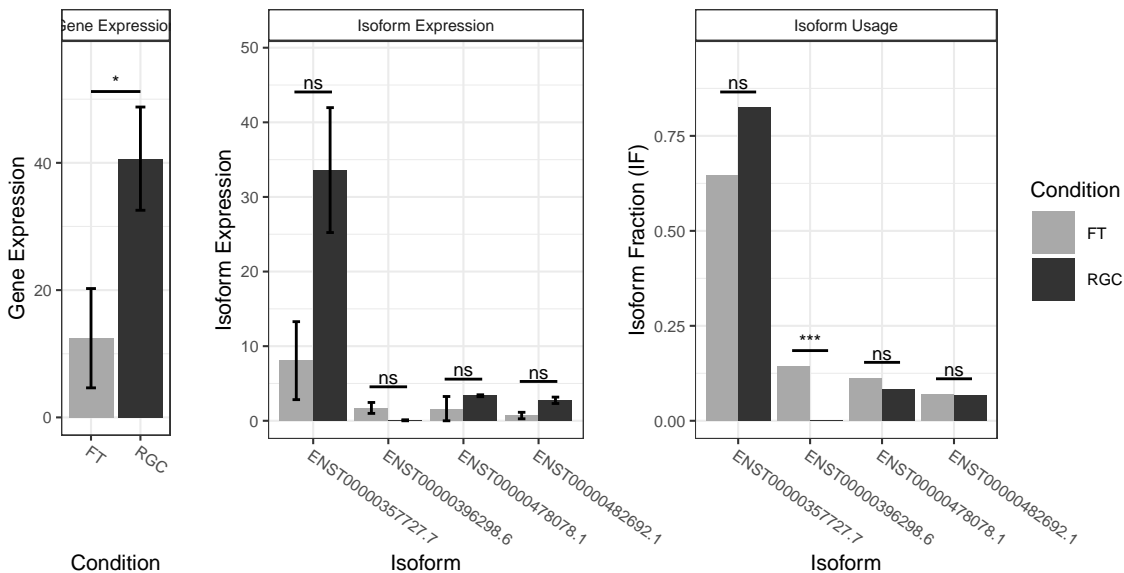
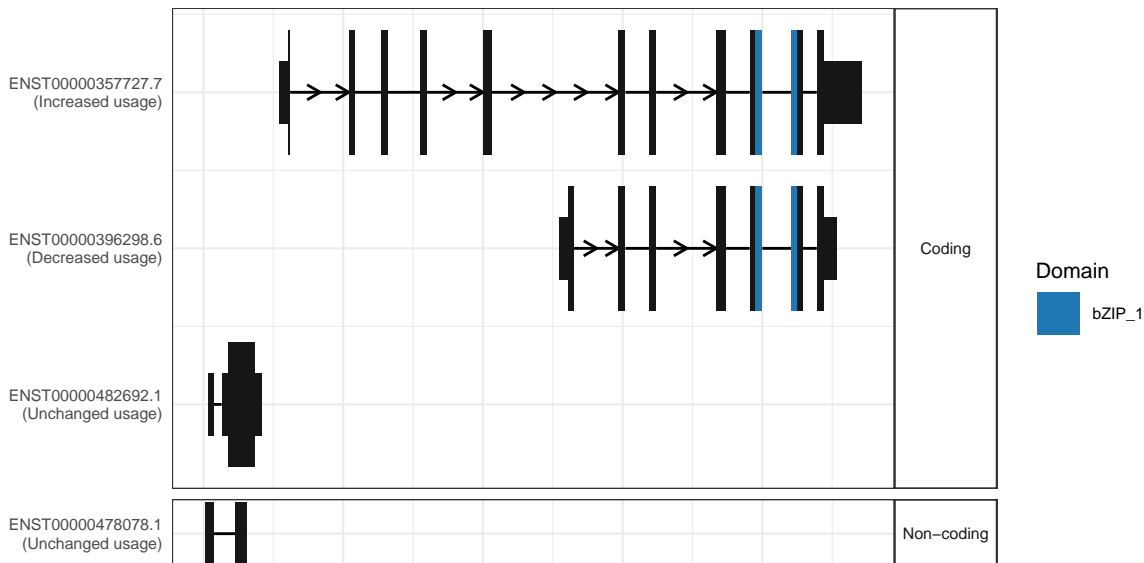


RHD_dimer



RHD_DNA_bind

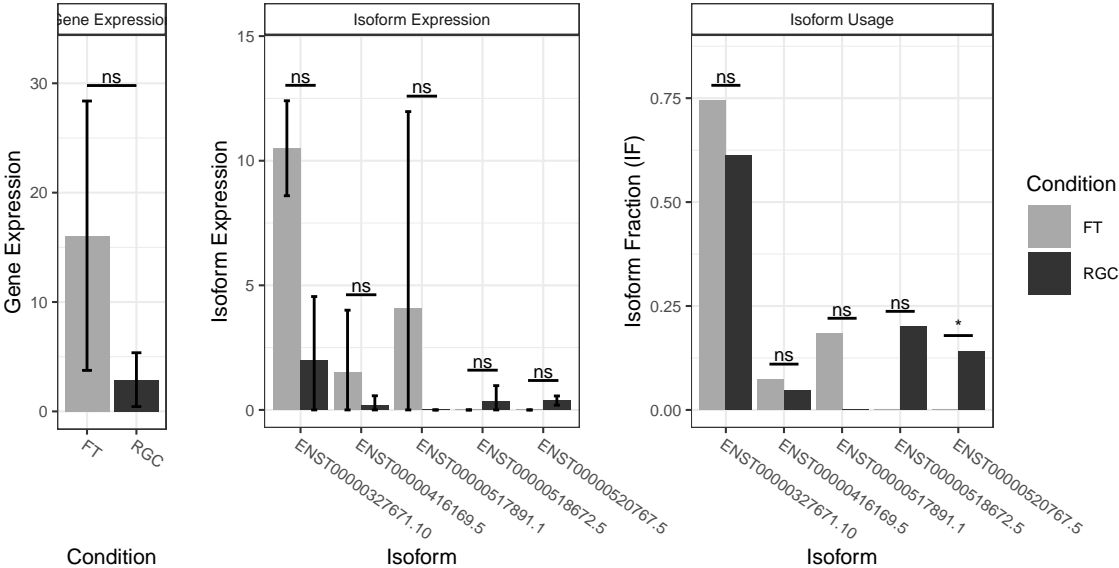
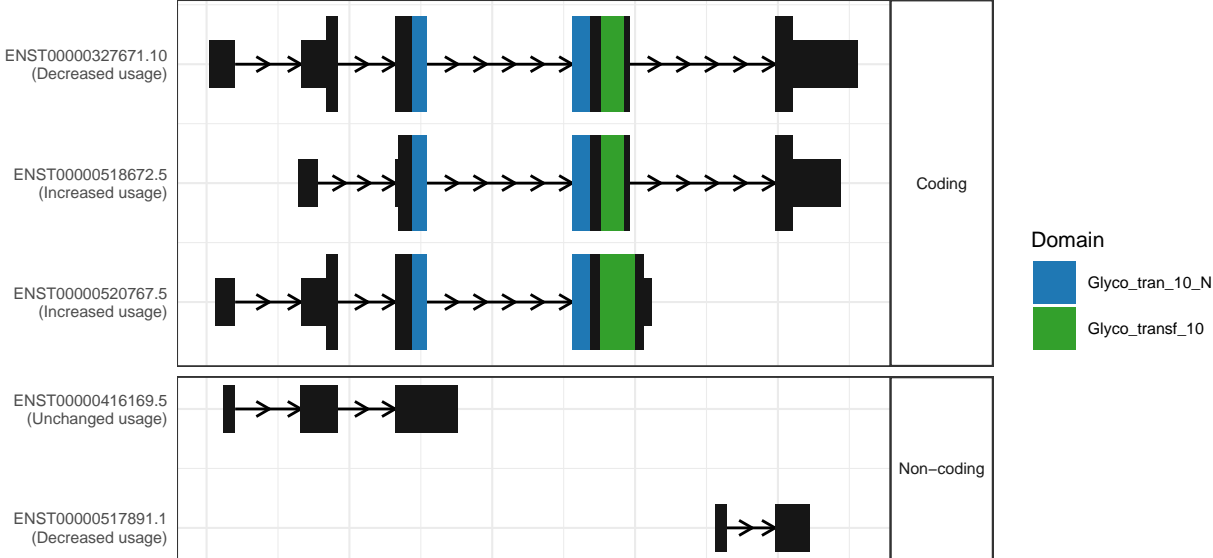
The isoform switch in CREB5 (FT vs RGC)



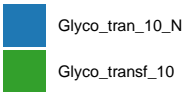
Domain

 bZIP_1

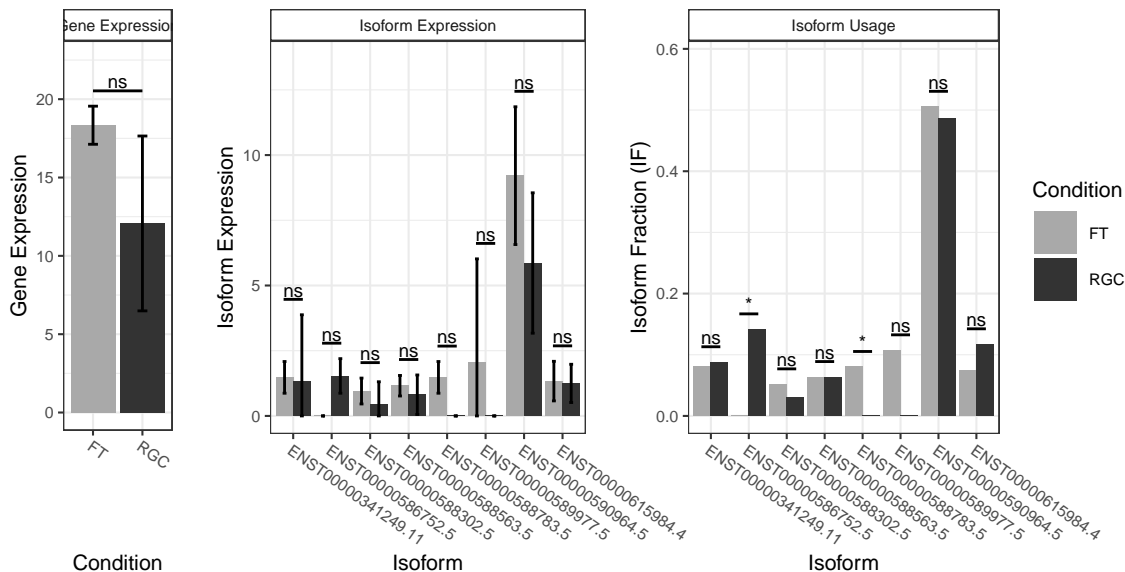
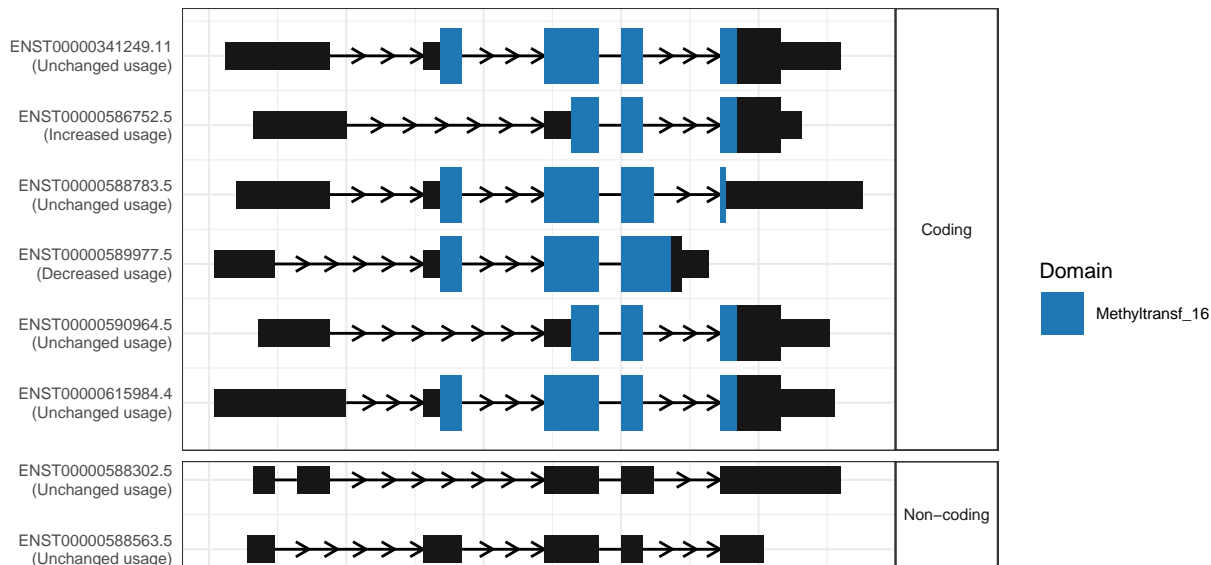
The isoform switch in FUT10 (FT vs RGC)



Domain



The isoform switch in METTL23 (FT vs RGC)

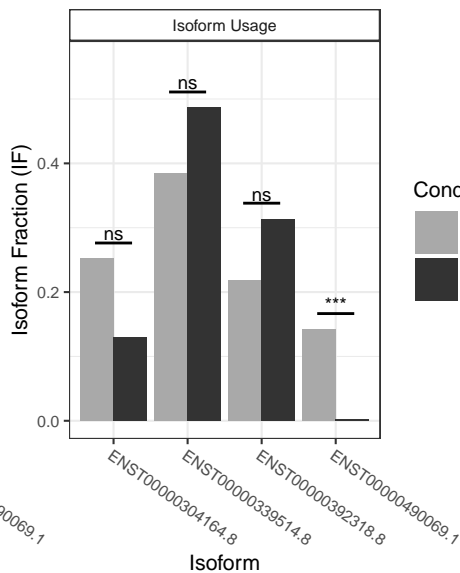
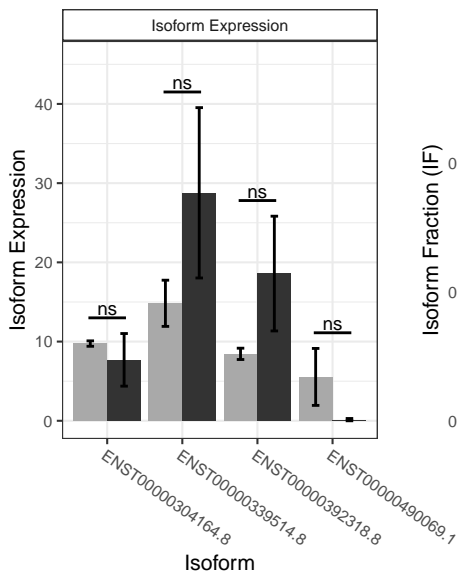
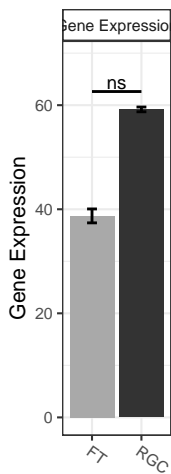
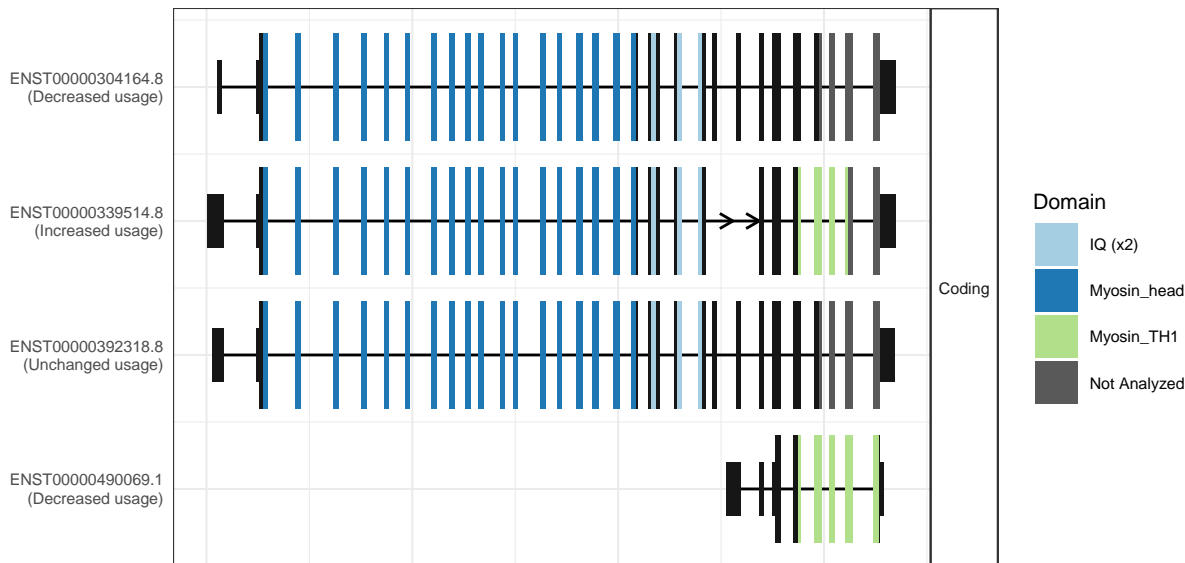


Domain



Methyltransf_16

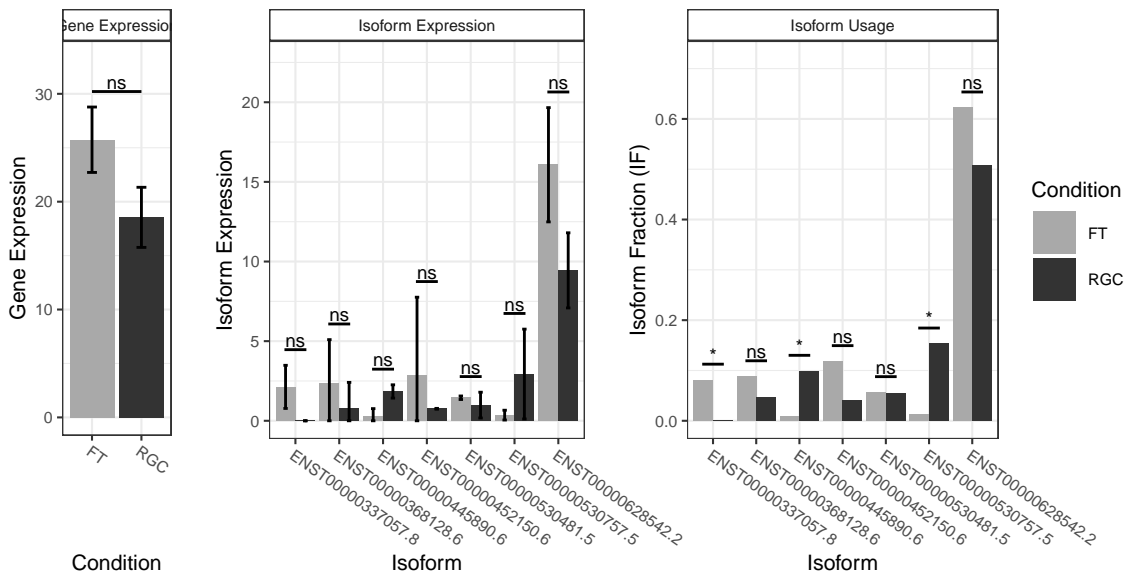
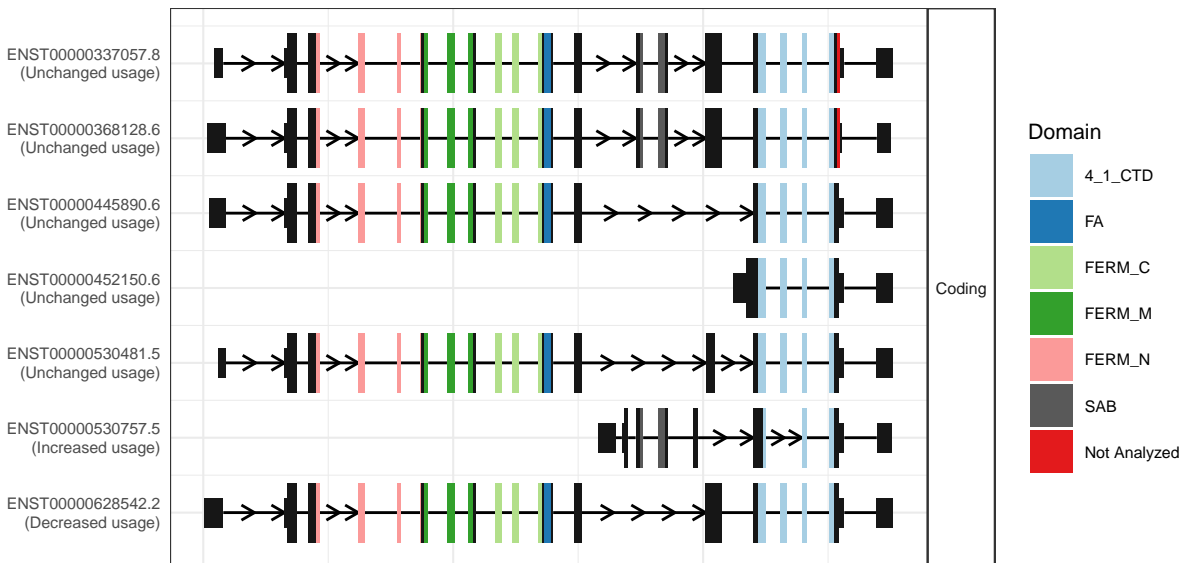
The isoform switch in MYO1B (FT vs RGC)










Domain

IQ (x2)
Myosin_head
Myosin_TH1
Not Analyzed

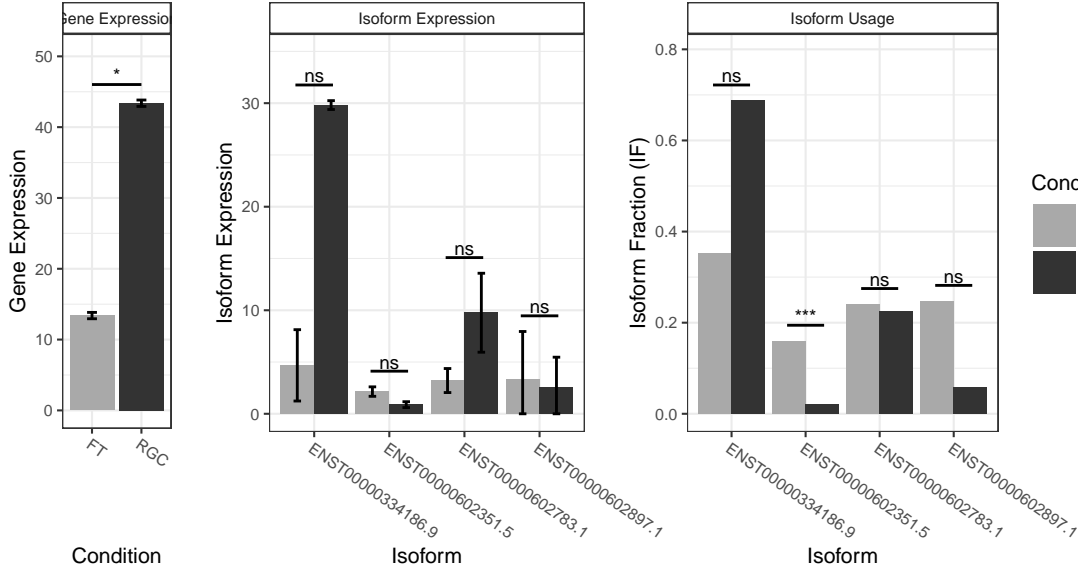
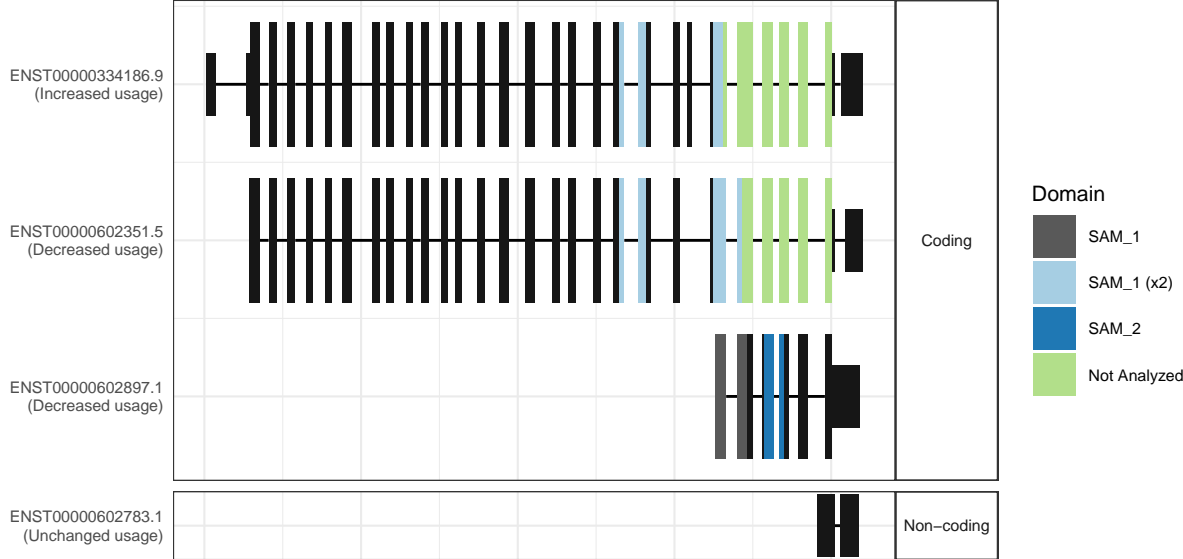
The isoform switch in EPB41L2 (FT vs RGC)



Domain

	4_1_CTD
	FA
	FERM_C
	FERM_M
	FERM_N
	SAB
	Not Analyzed

The isoform switch in PPFIA3 (FT vs RGC)



Domain



SAM_1



SAM_1 (x2)

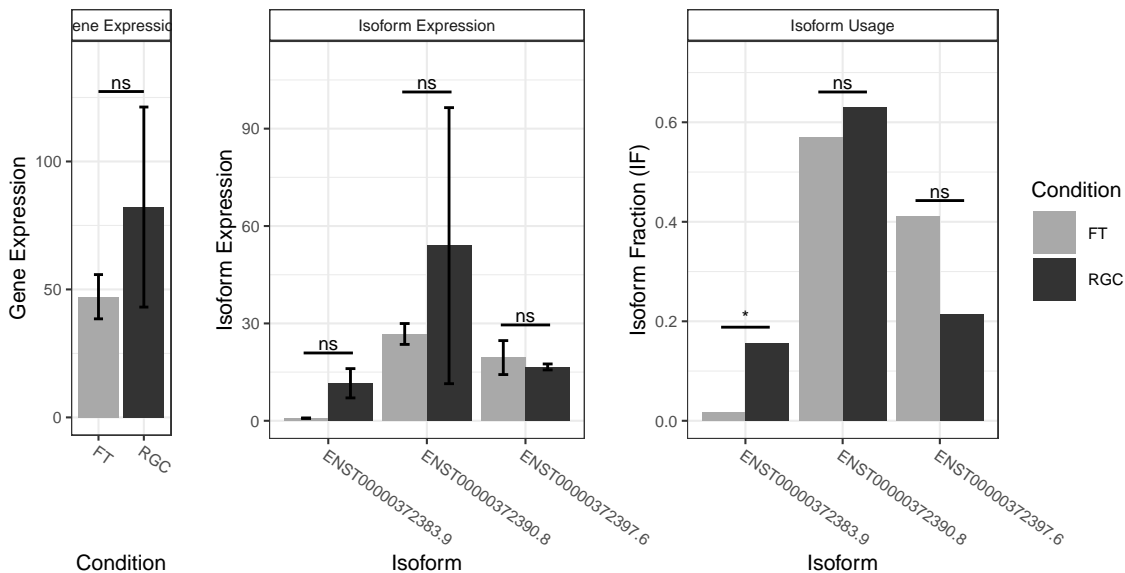
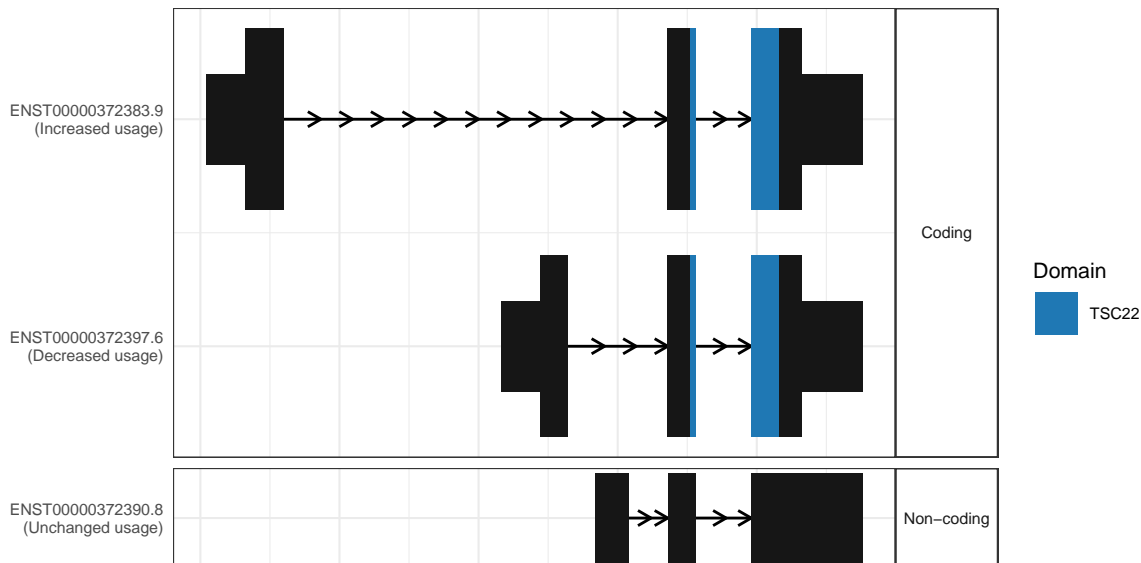


SAM_2



Not Analyzed

The isoform switch in TSC22D3 (FT vs RGC)

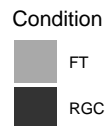
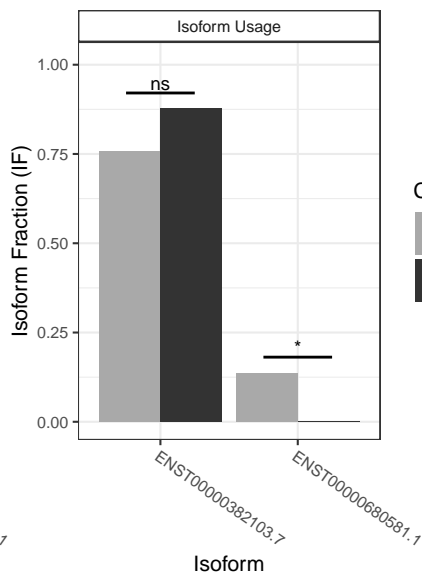
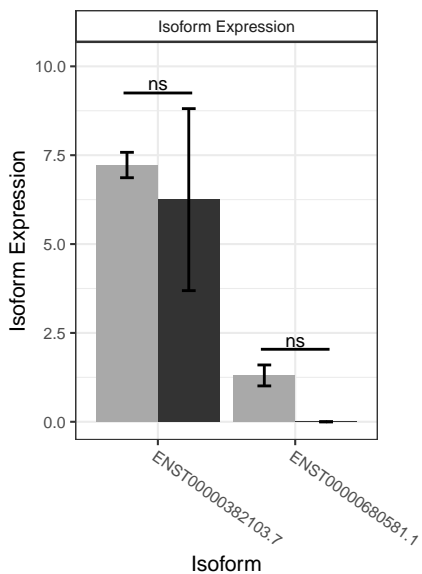
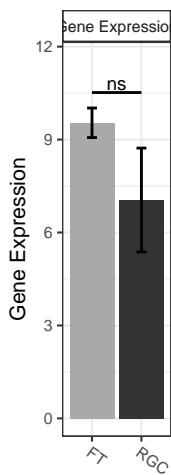
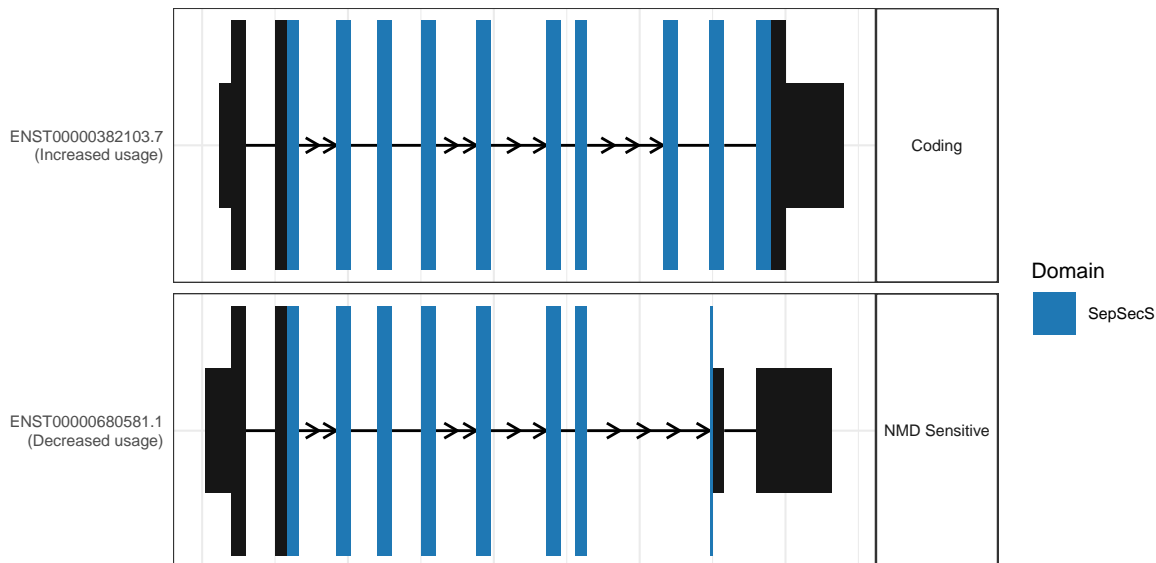


Domain



TSC22

The isoform switch in SEPSECS (FT vs RGC)

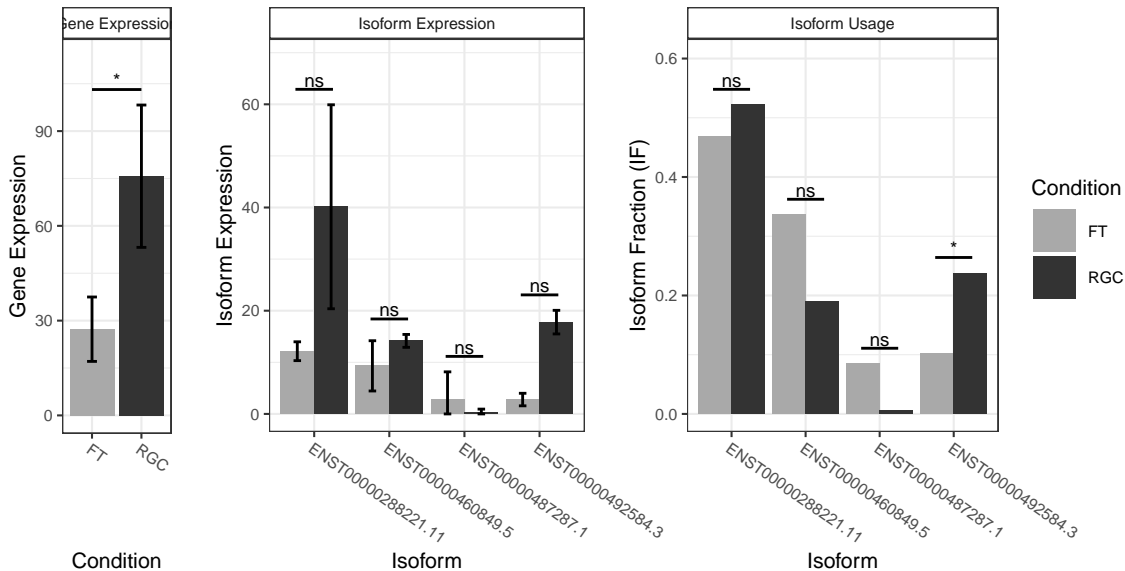
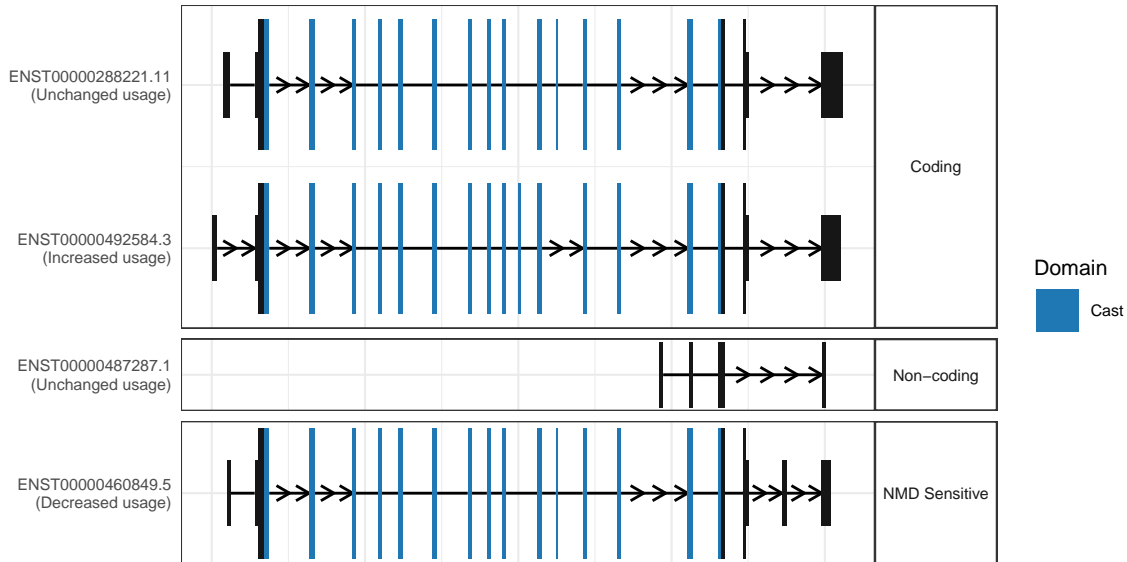


Domain



SepSecS

The isoform switch in ERC2 (FT vs RGC)

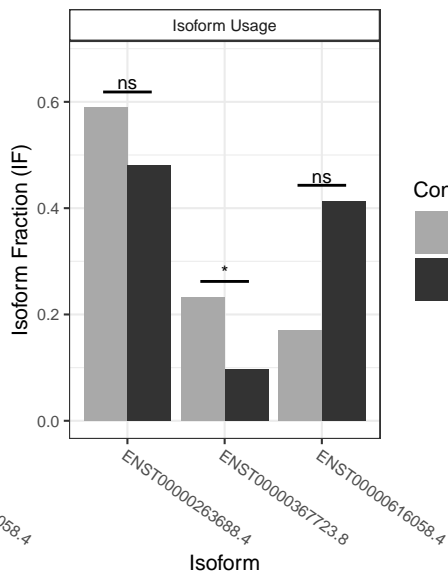
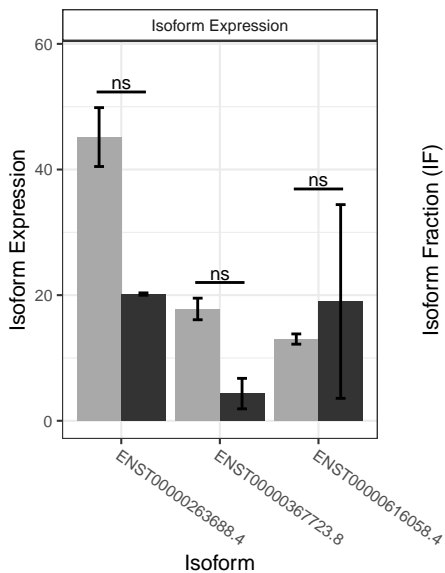
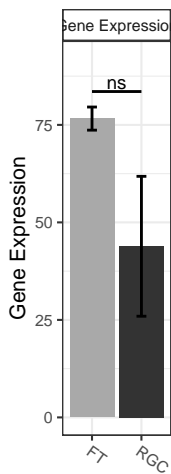
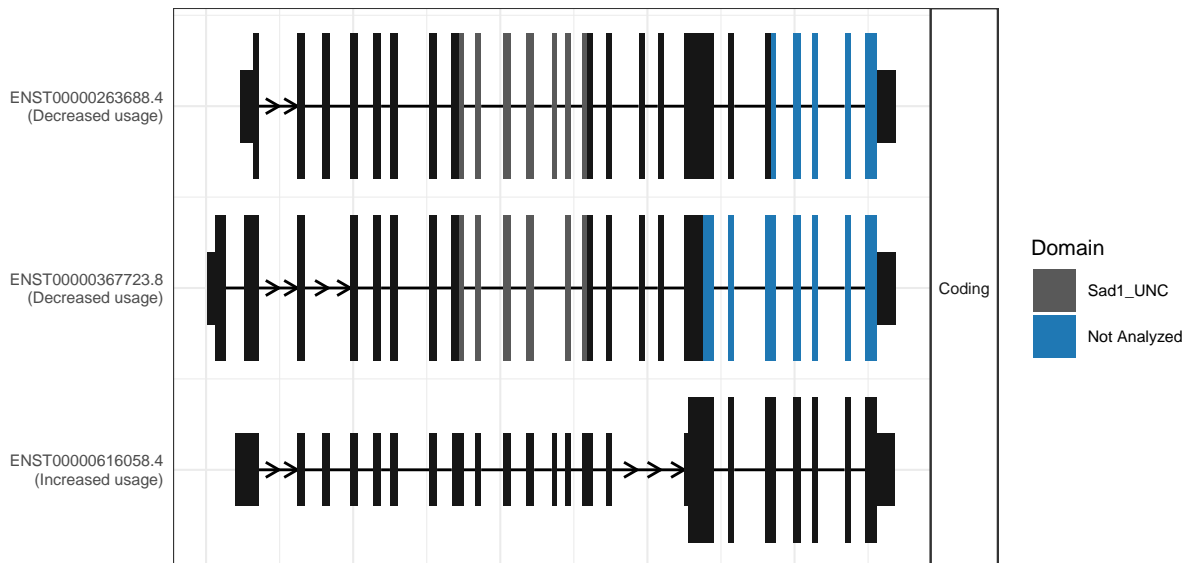


Domain



Cast

The isoform switch in SUCO (FT vs RGC)



Domain

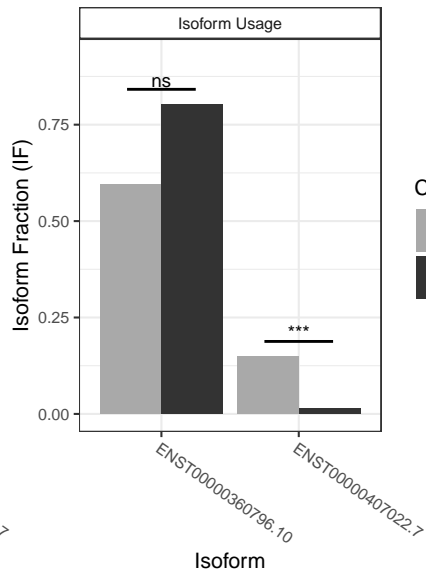
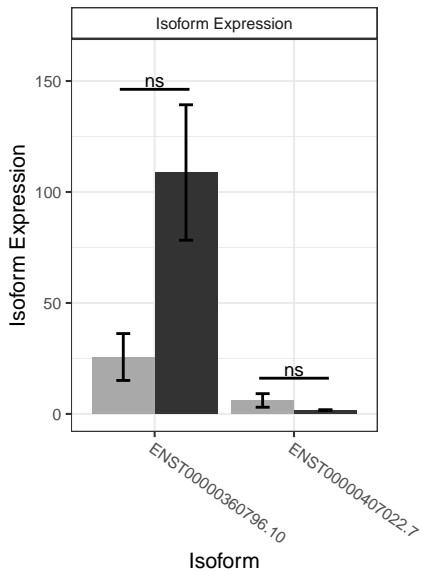
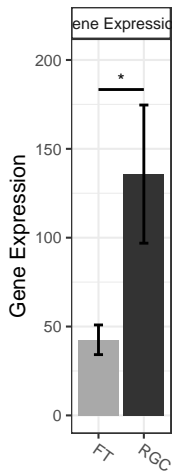
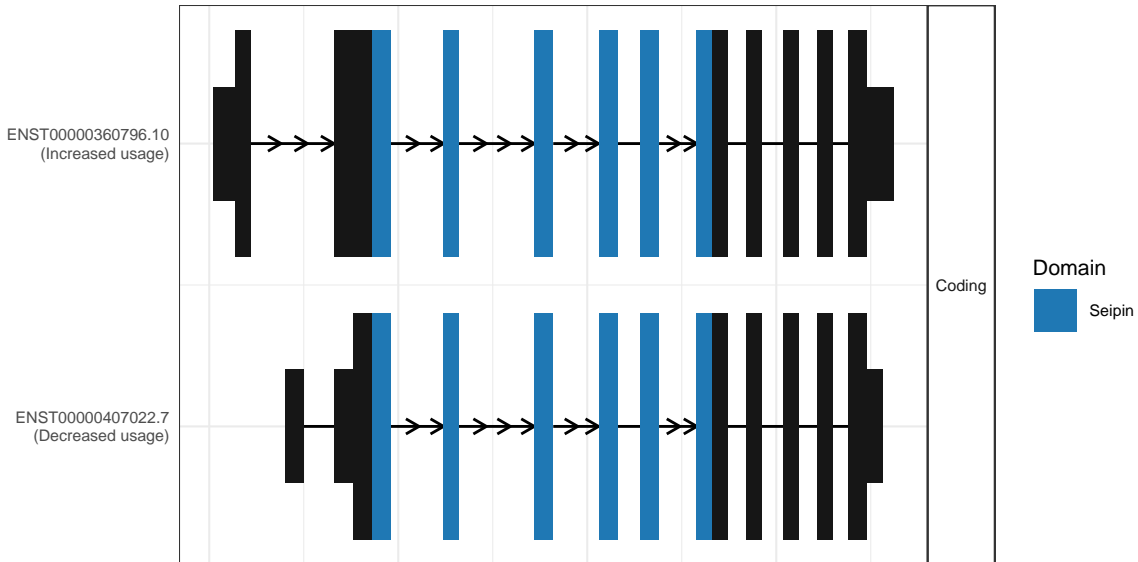


Sad1_UNC



Not Analyzed

The isoform switch in BSCL2 (FT vs RGC)

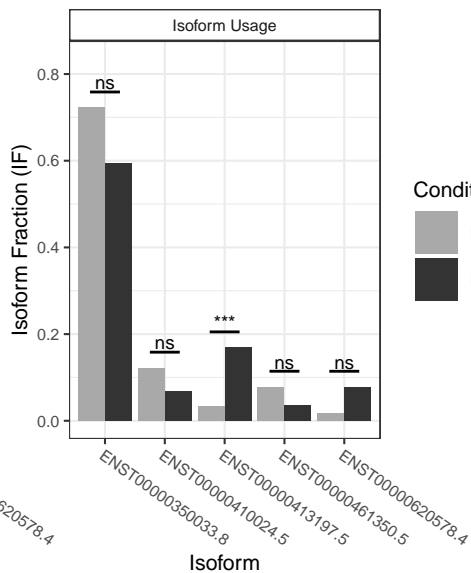
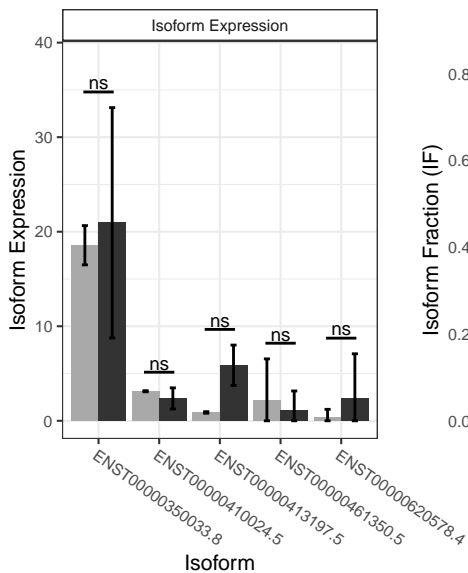
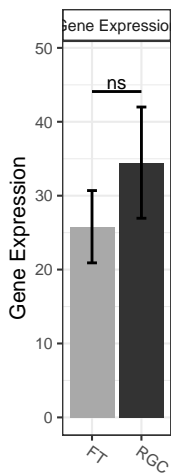
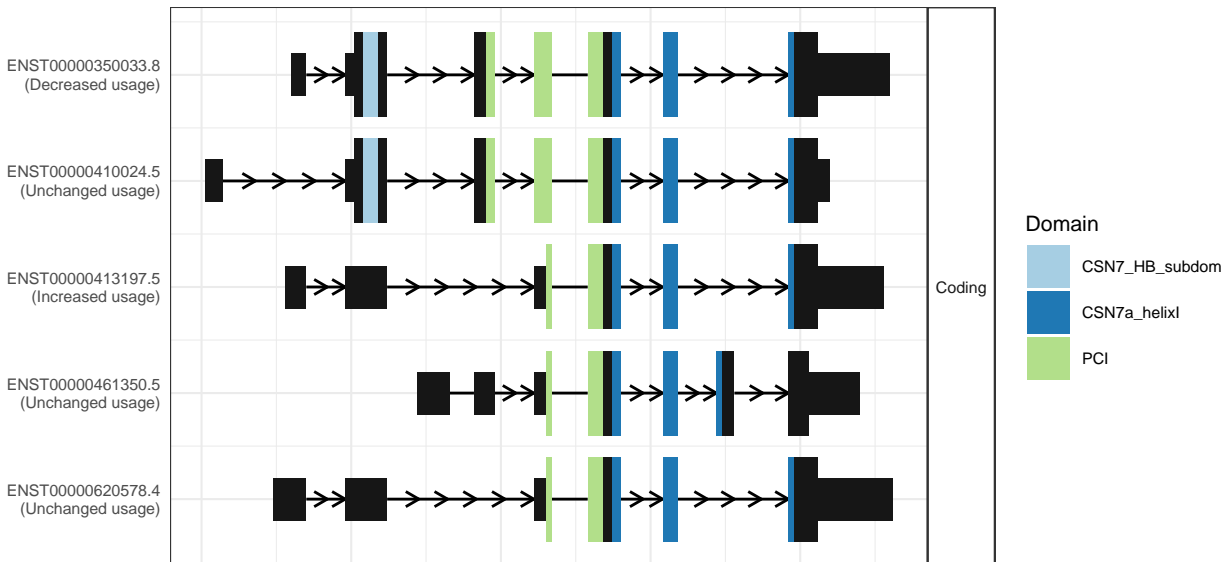


Domain






Seipin

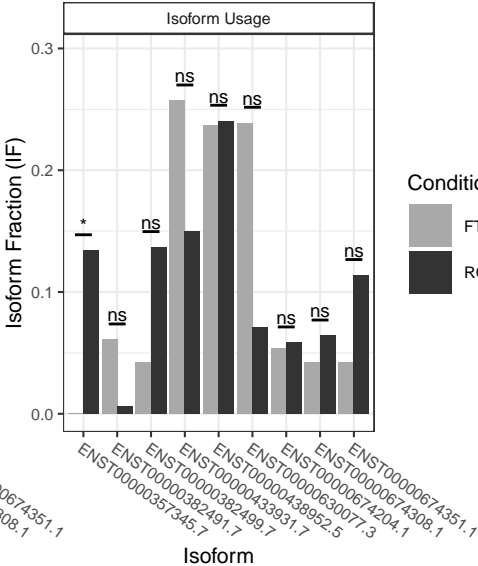
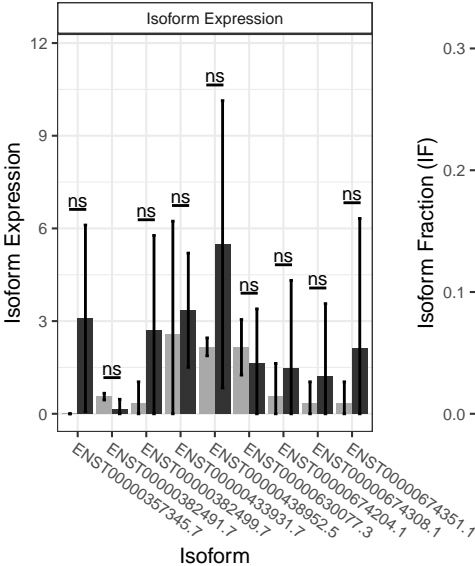
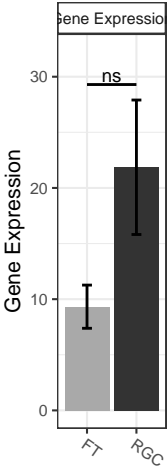
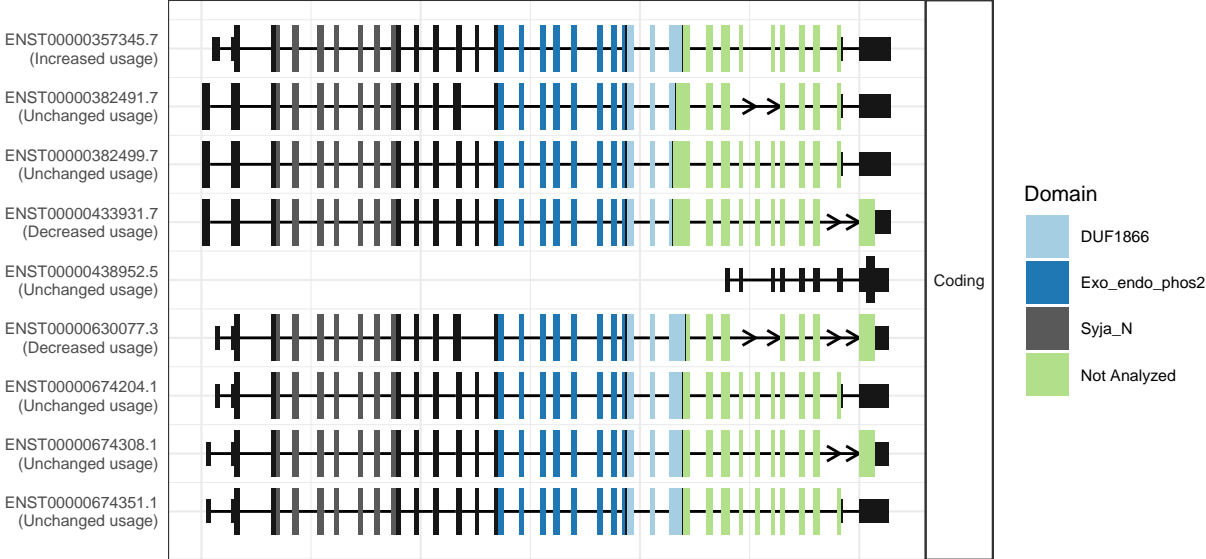
The isoform switch in COPS7B (FT vs RGC)



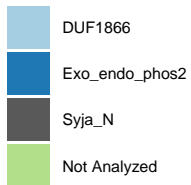
Domain

	CSN7_HB_subdom
	CSN7a_helixI
	PCI

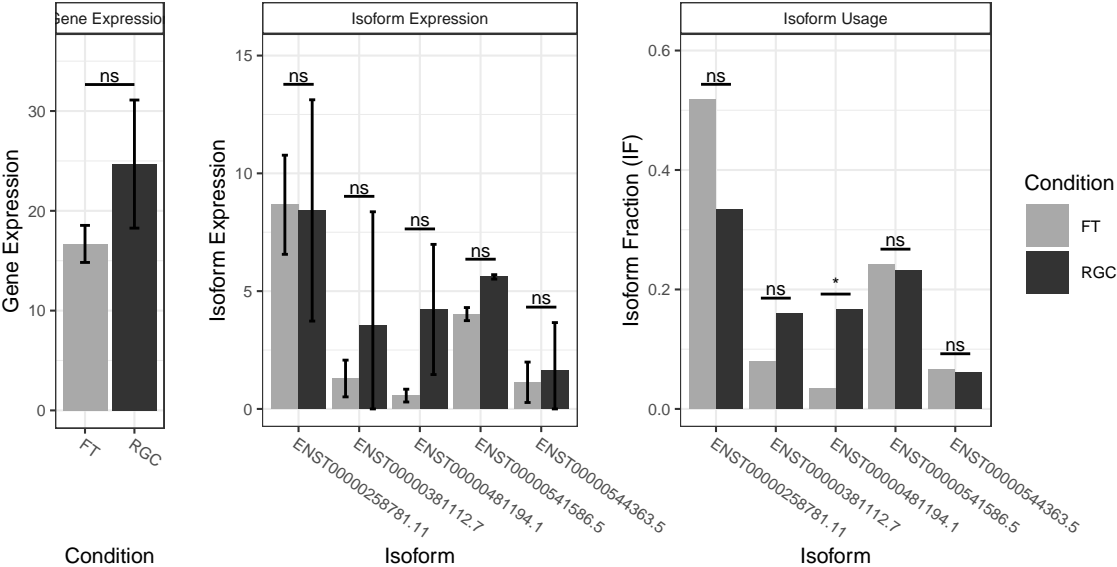
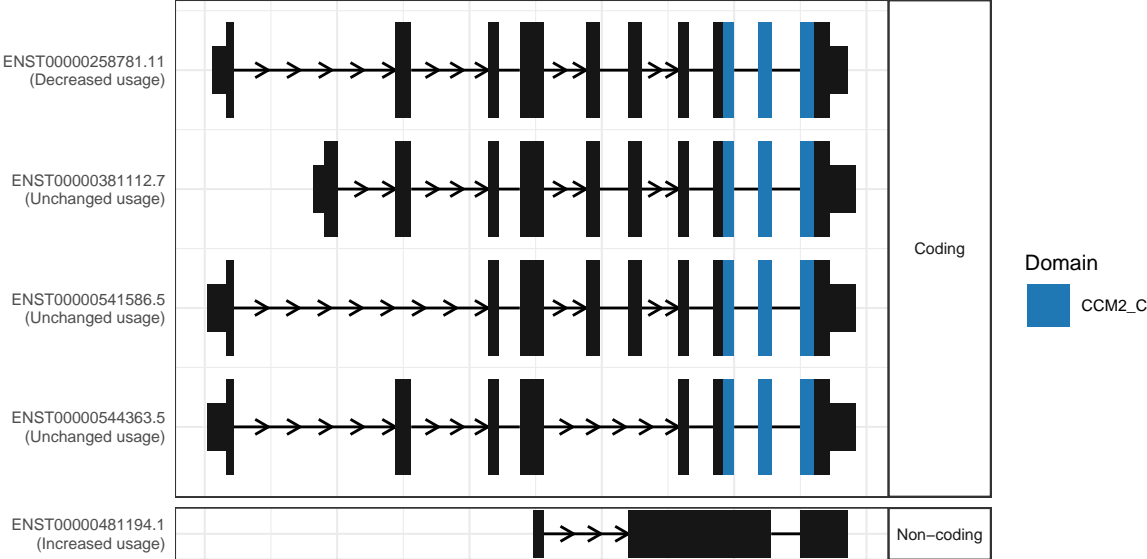
The isoform switch in SYNJ1 (FT vs RGC)



Domain



The isoform switch in CCM2 (FT vs RGC)

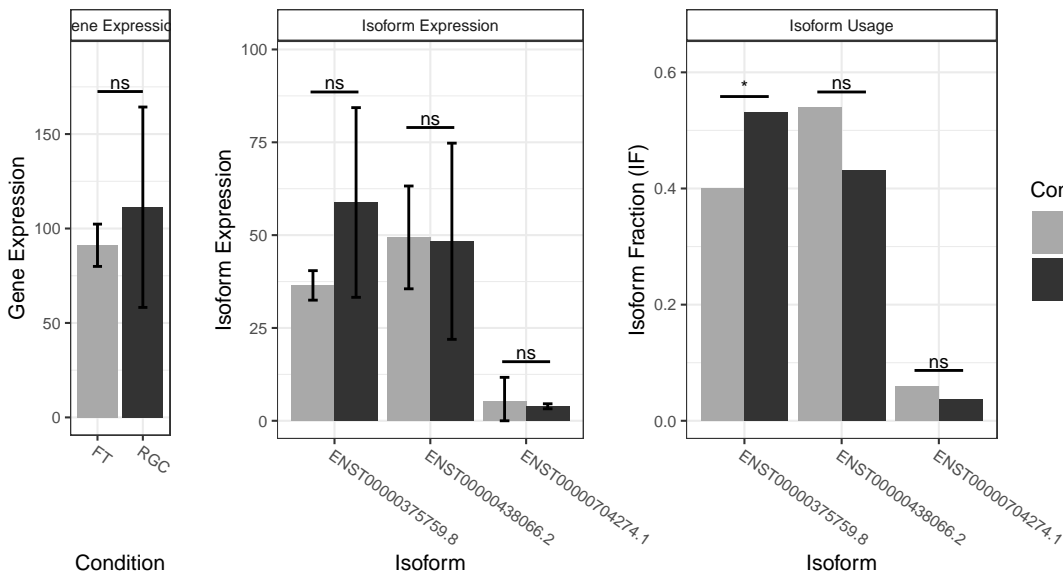
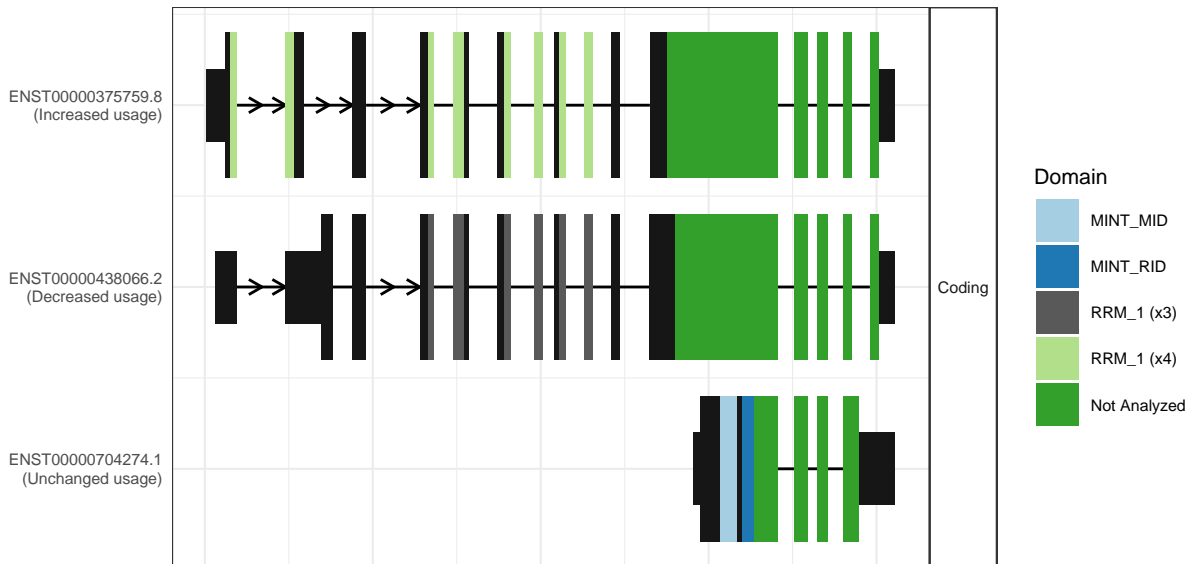


Domain

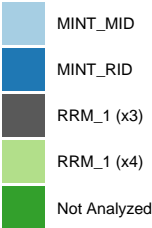


CCM2_C

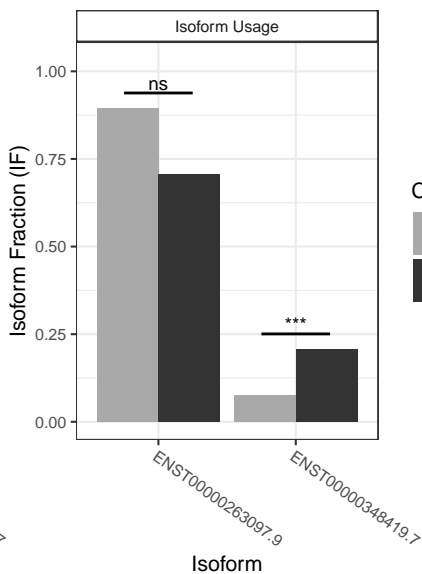
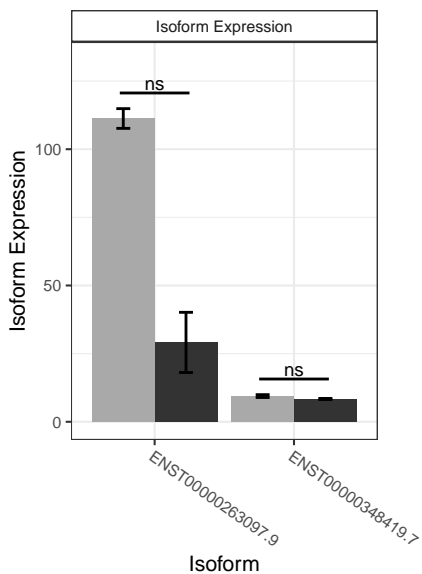
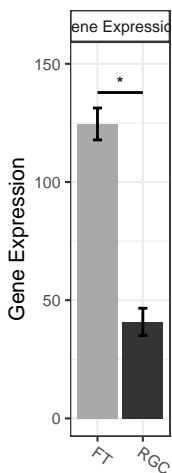
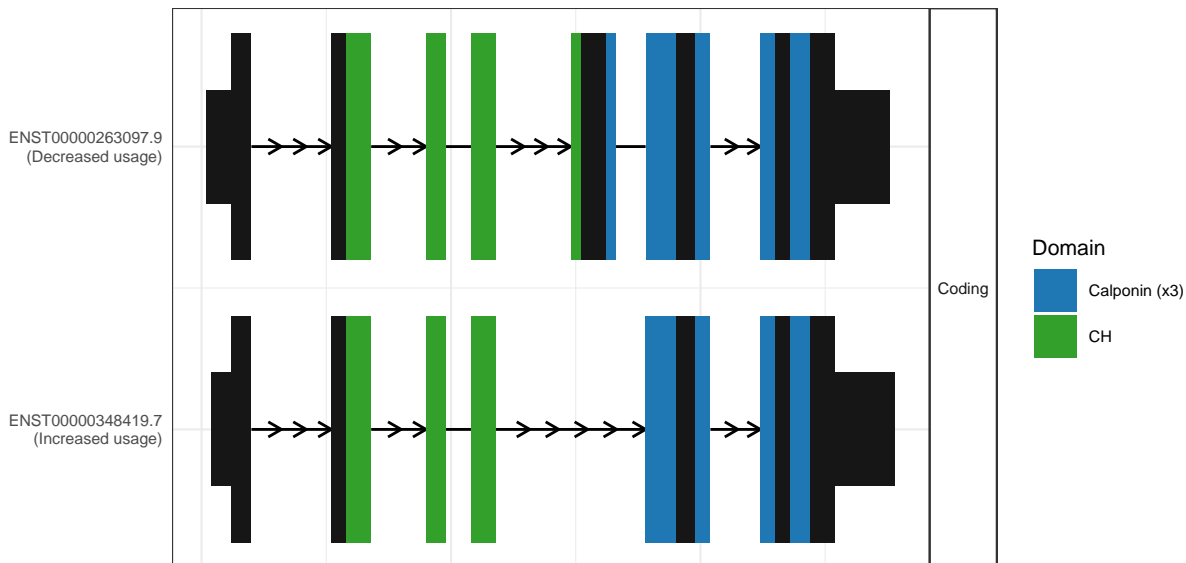
The isoform switch in SPEN (FT vs RGC)



Domain



The isoform switch in CNN2 (FT vs RGC)



Domain

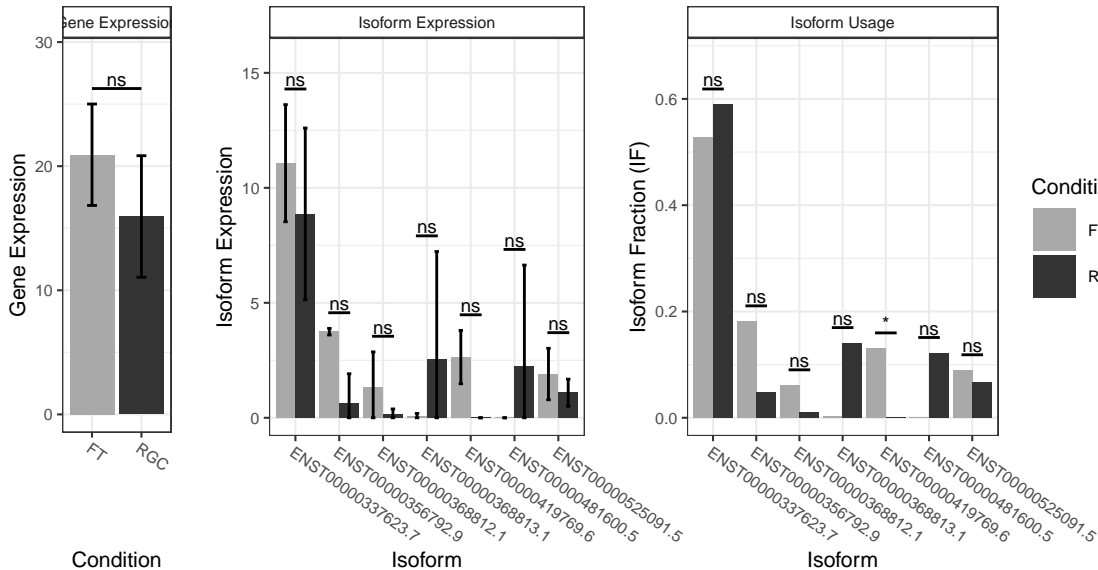
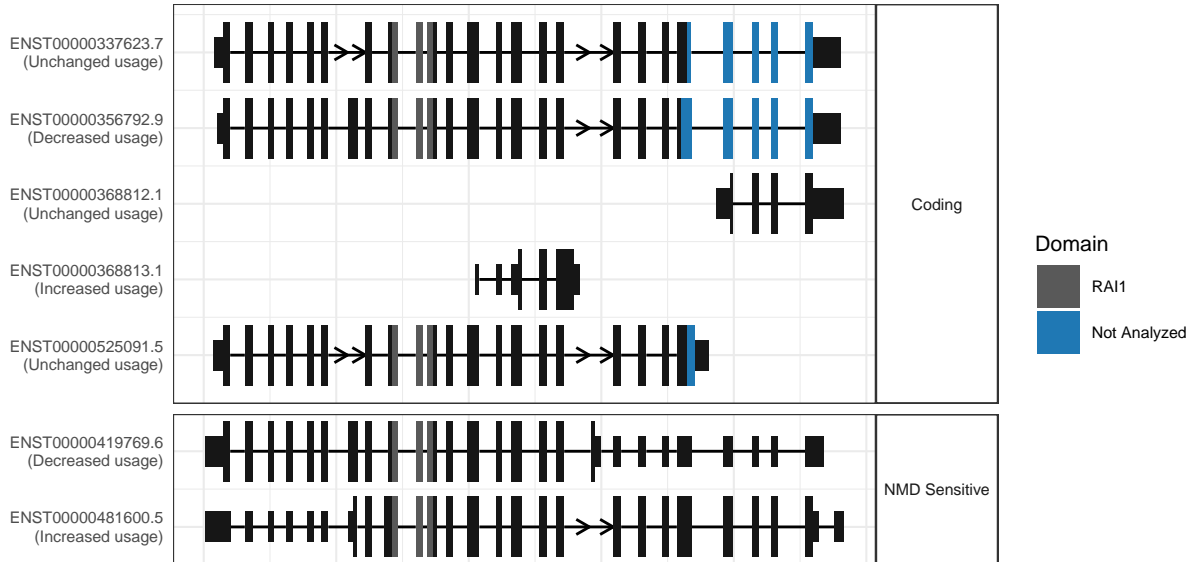


Calponin (x3)



CH

The isoform switch in EDRF1 (FT vs RGC)



Domain

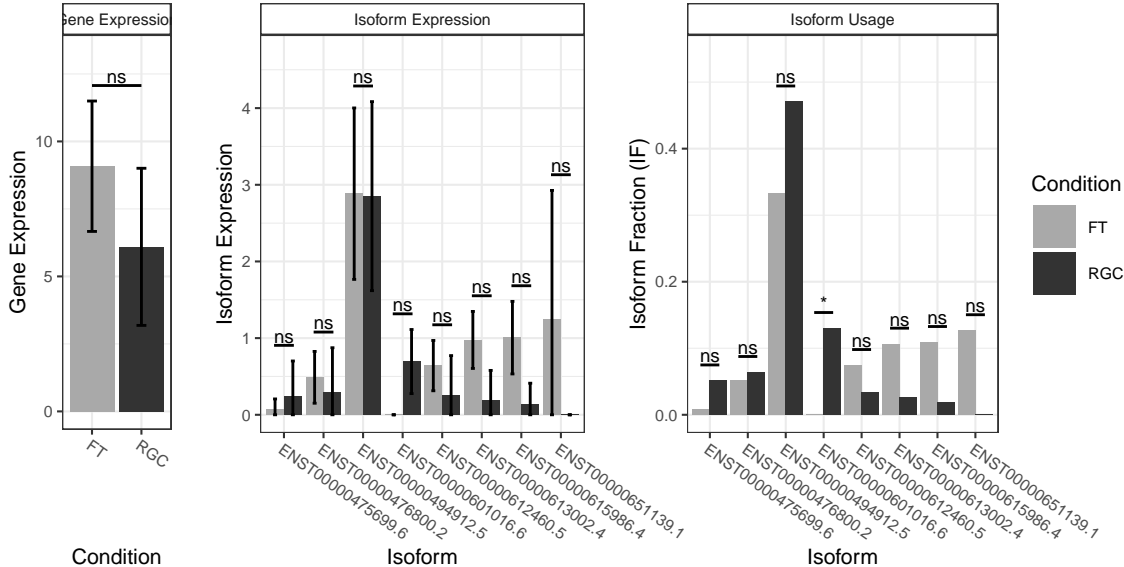
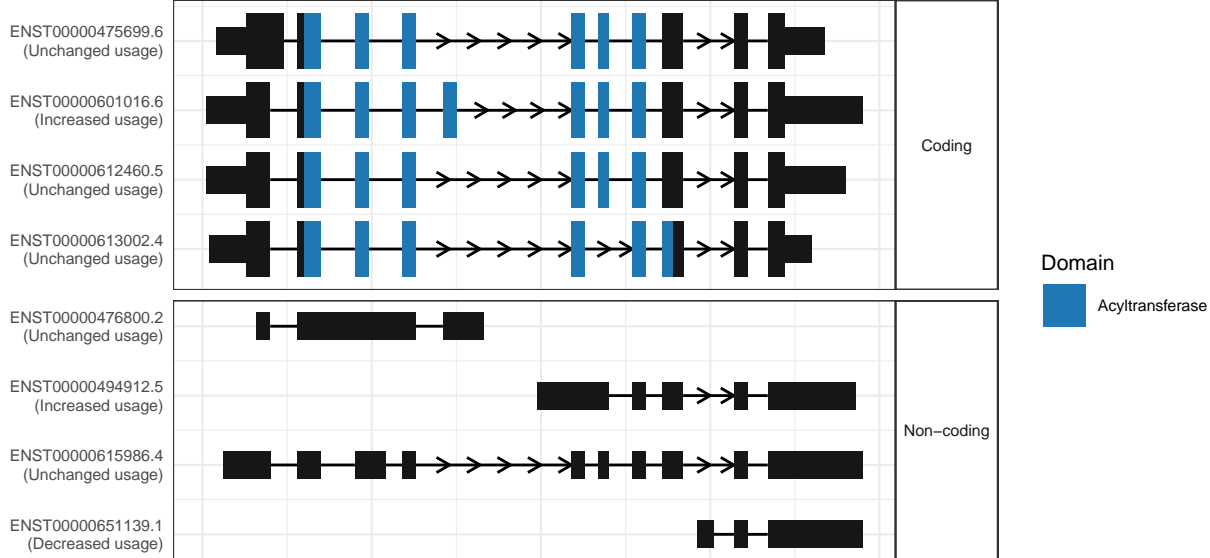


RAI1



Not Analyzed

The isoform switch in TAFAZZIN (FT vs RGC)

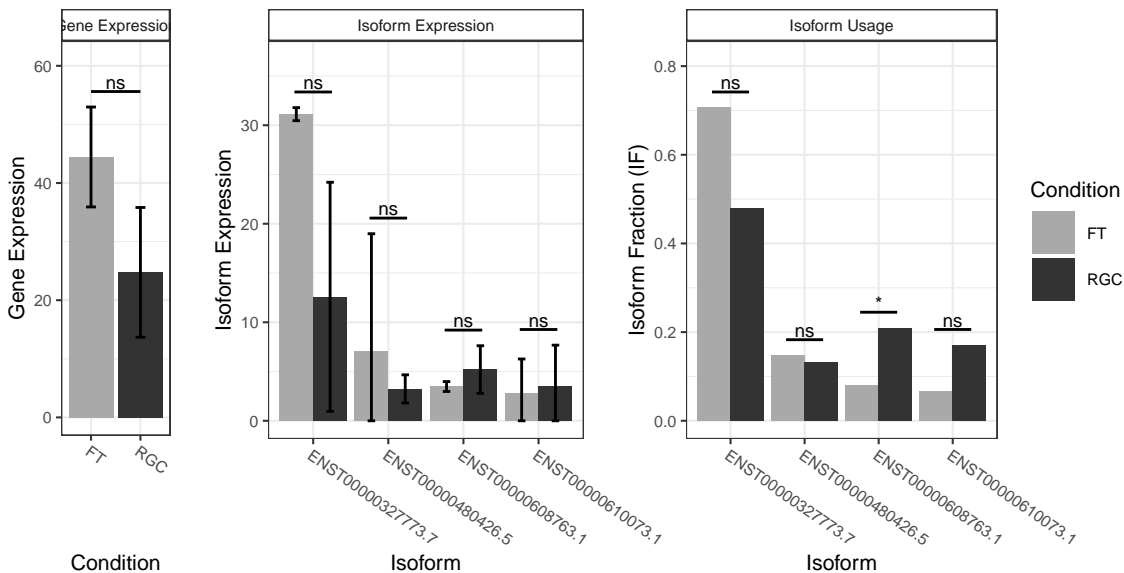
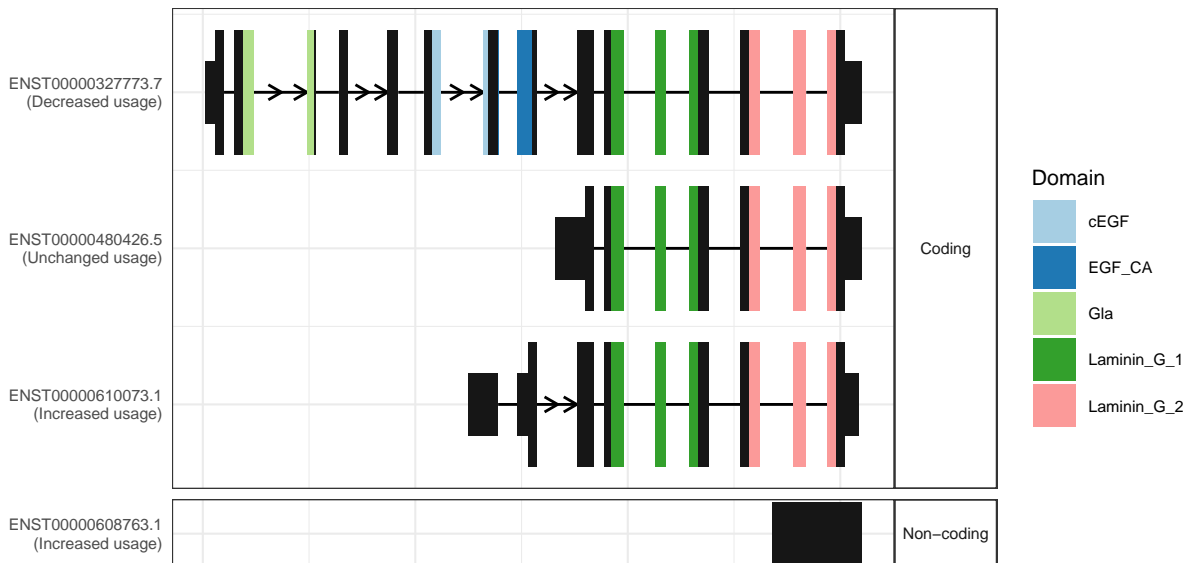


Domain








Acyltransferase

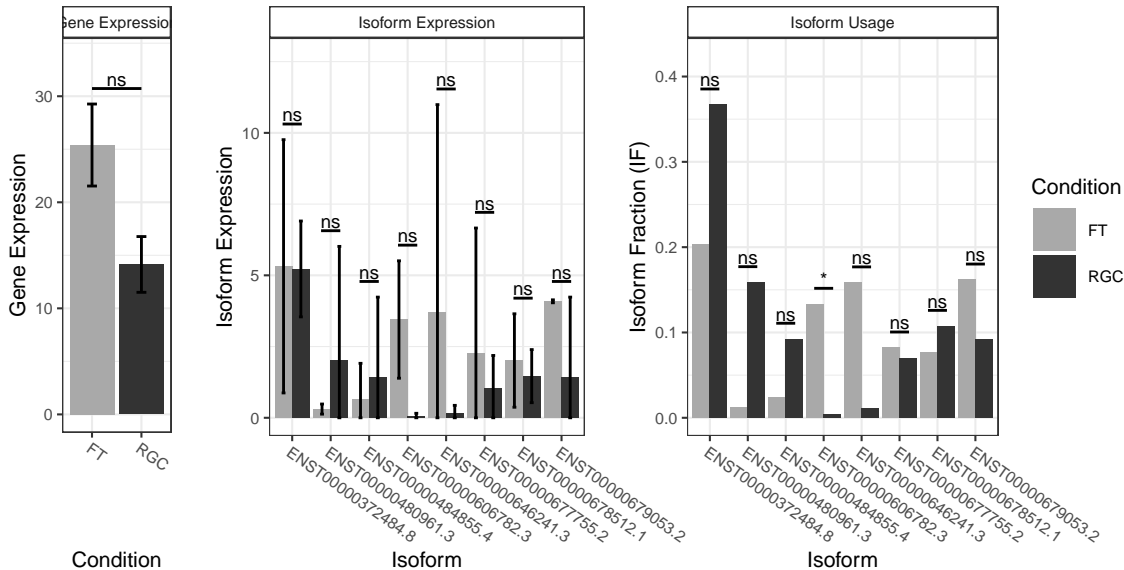
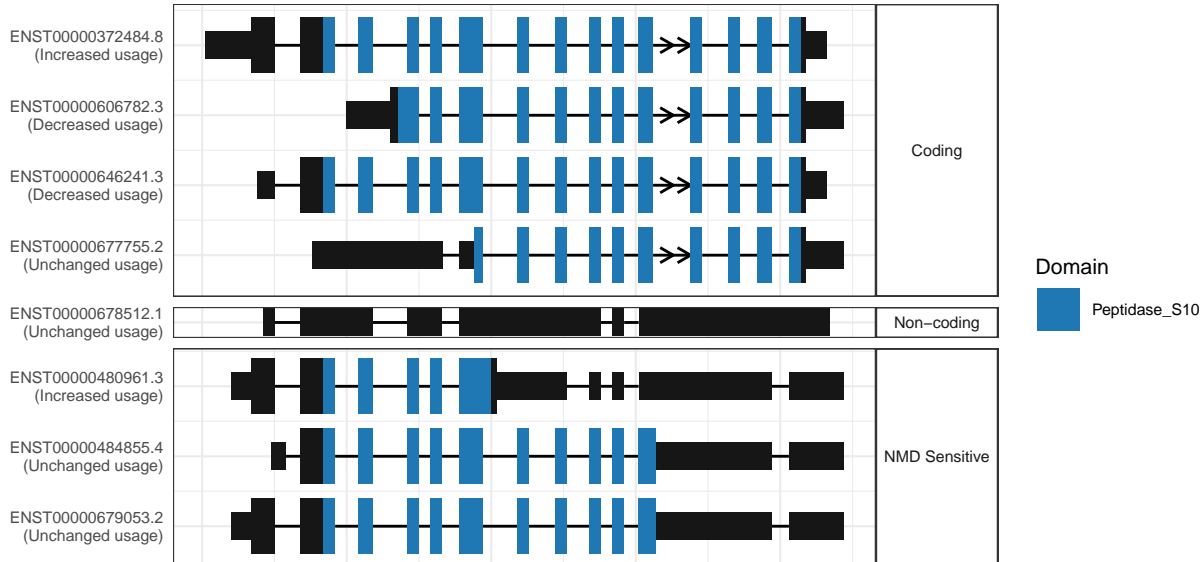
The isoform switch in GAS6 (FT vs RGC)



Domain

	cEGF
	EGF_CA
	Gla
	Laminin_G_1
	Laminin_G_2

The isoform switch in CTSA (FT vs RGC)

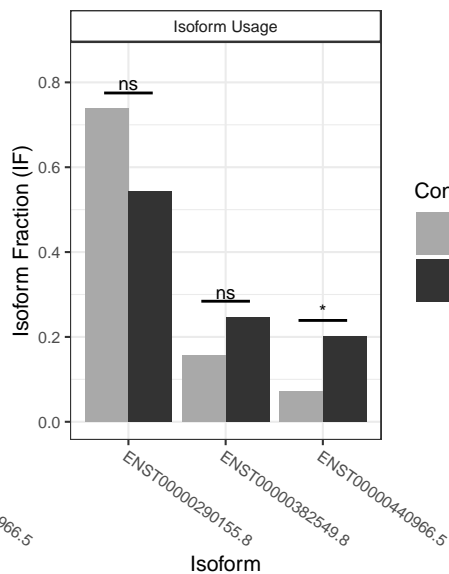
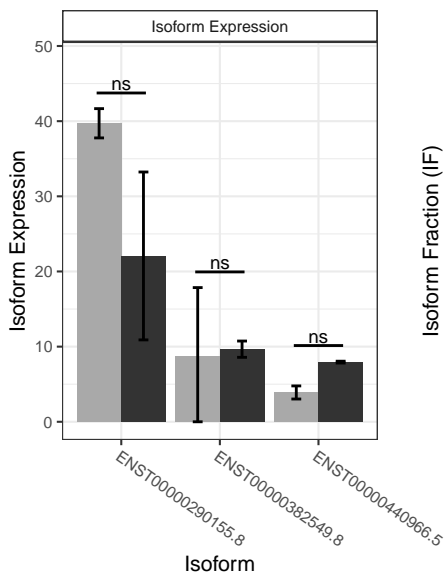
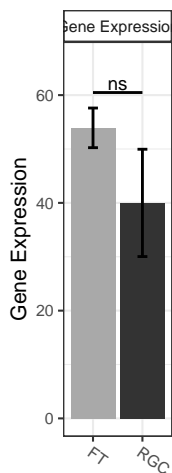
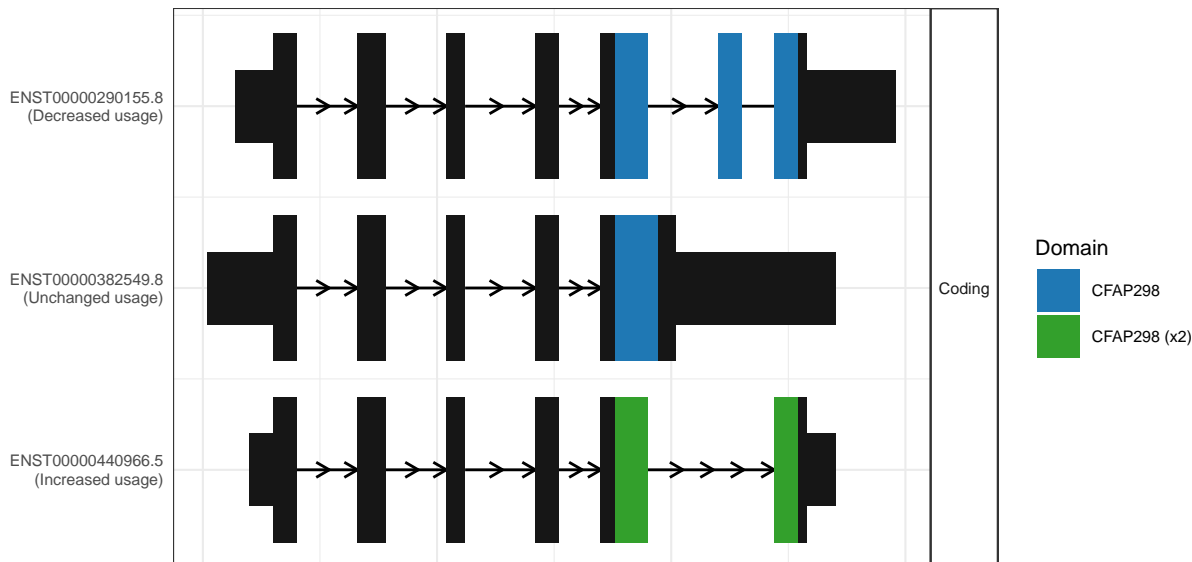


Domain



Peptidase_S10

The isoform switch in CFAP298 (FT vs RGC)



Domain

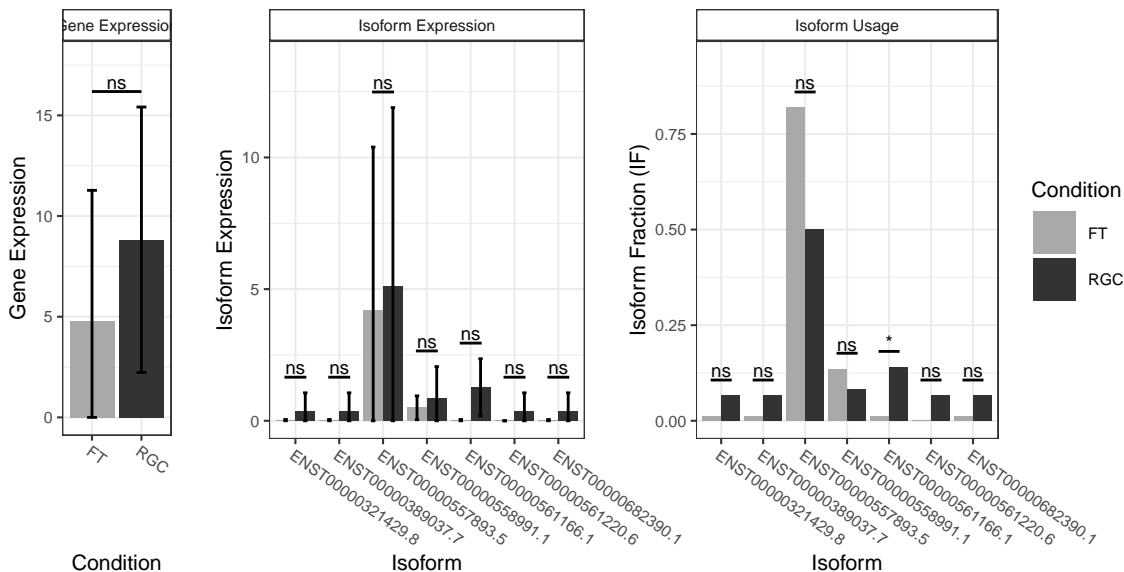
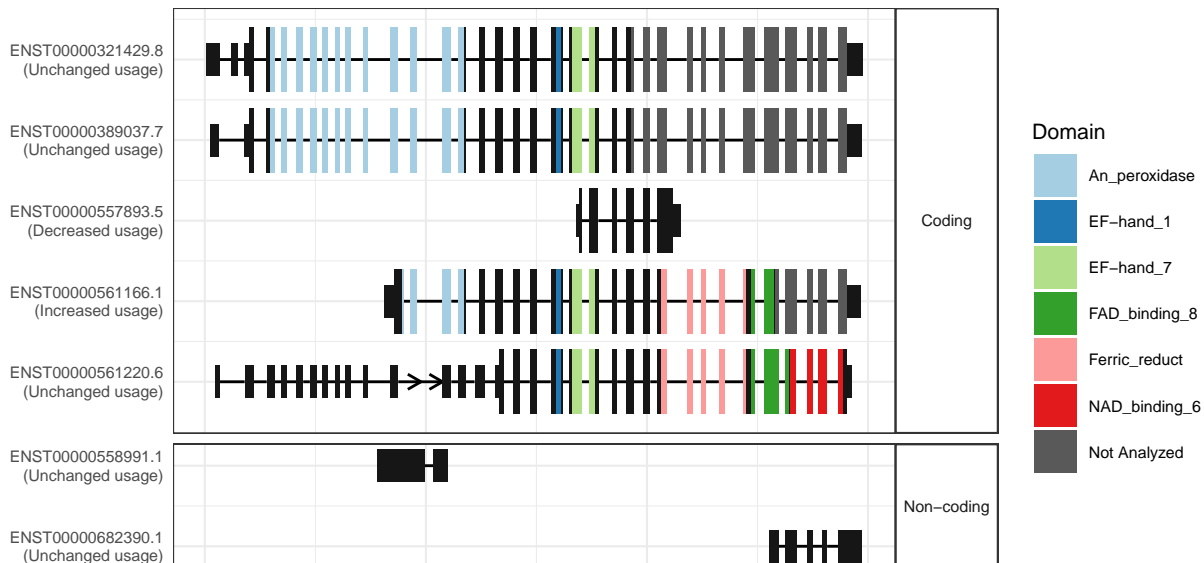


CFAP298



CFAP298 (x2)

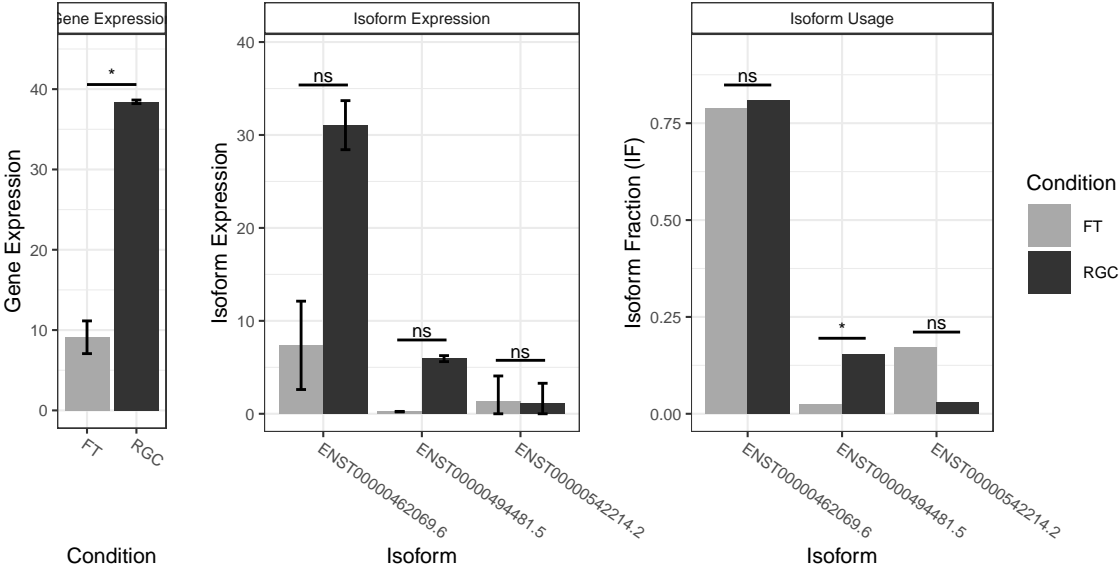
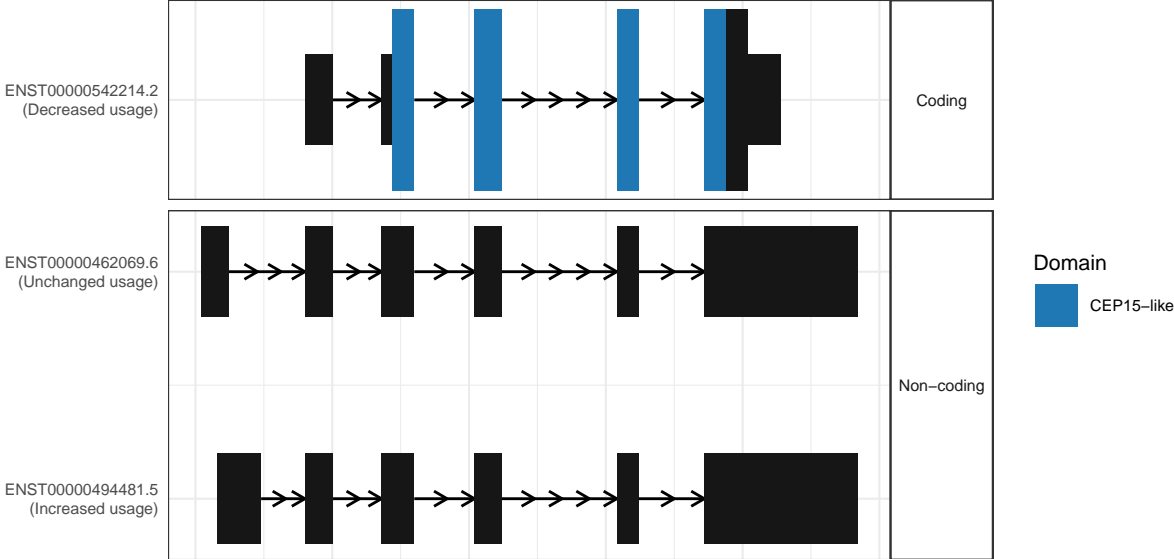
The isoform switch in DUOX1 (FT vs RGC)



Domain

An_peroxidase
EF-hand_1
EF-hand_7
FAD_binding_8
Ferric_reduct
NAD_binding_6
Not Analyzed

The isoform switch in CEP15 (FT vs RGC)

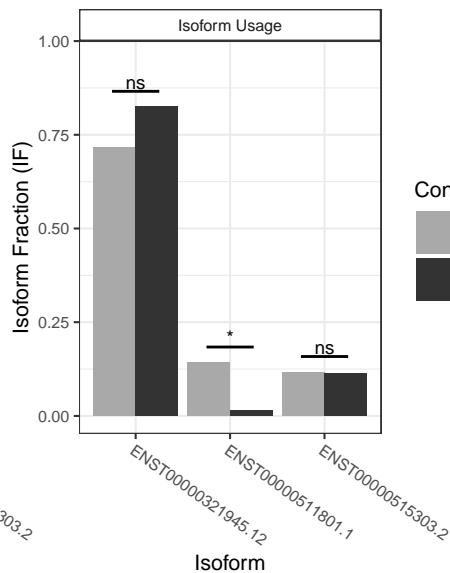
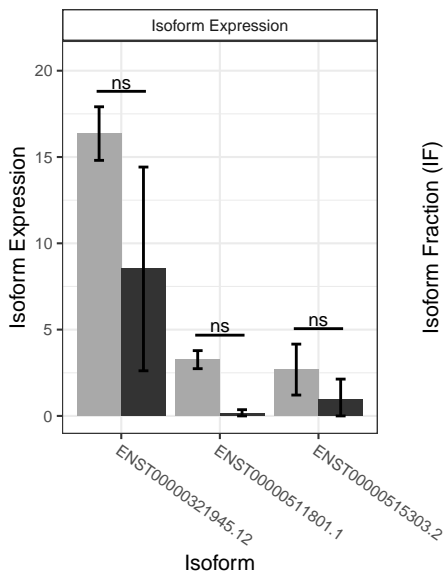
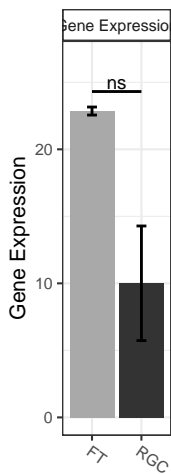
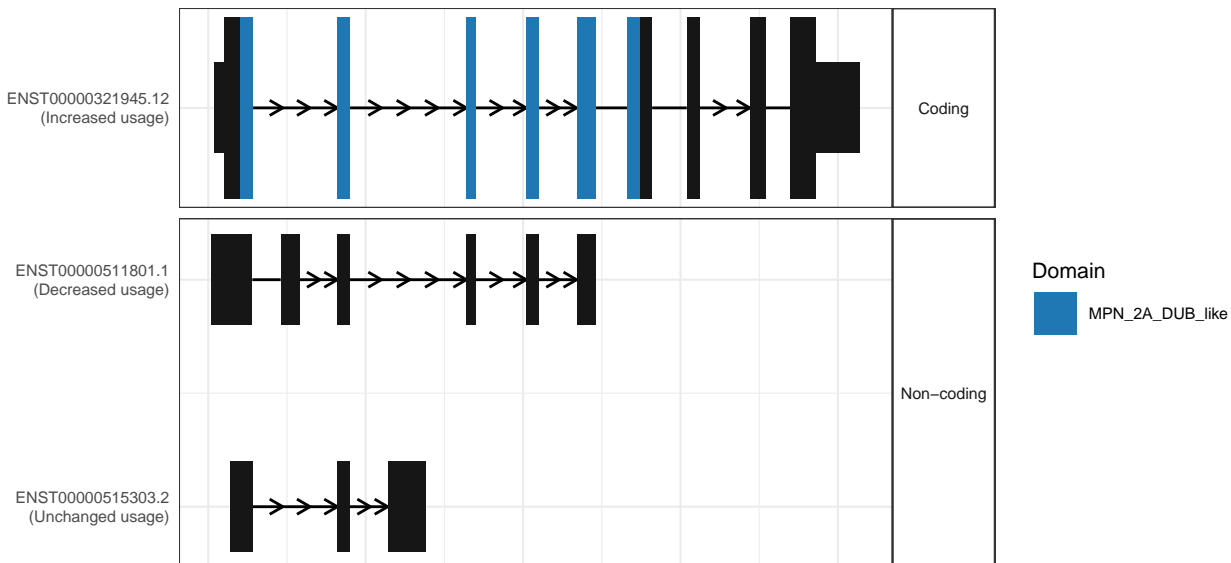


Domain



CEP15-like

The isoform switch in ABRAXAS1 (FT vs RGC)

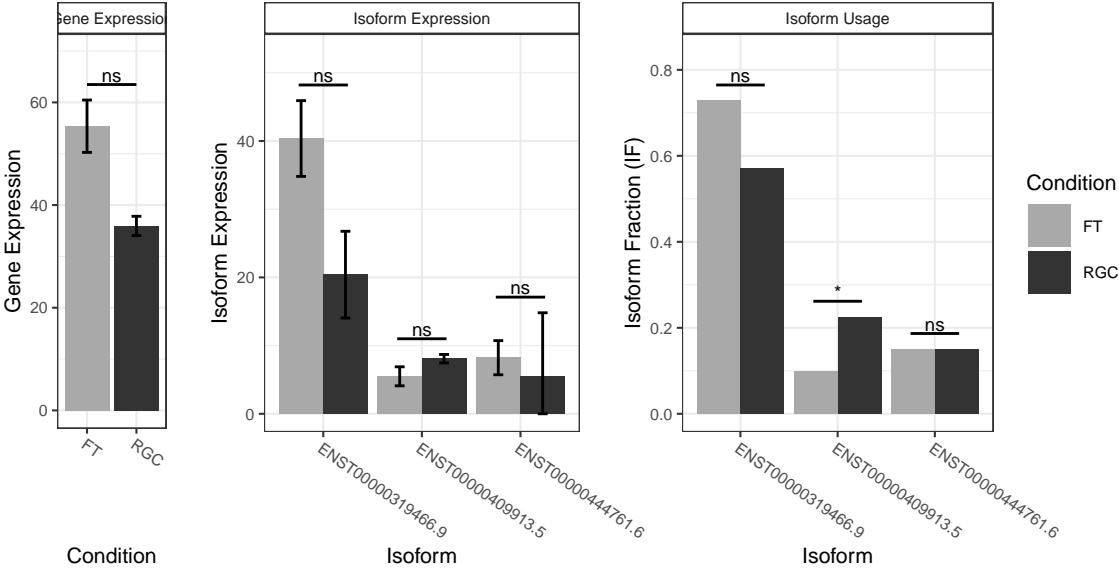
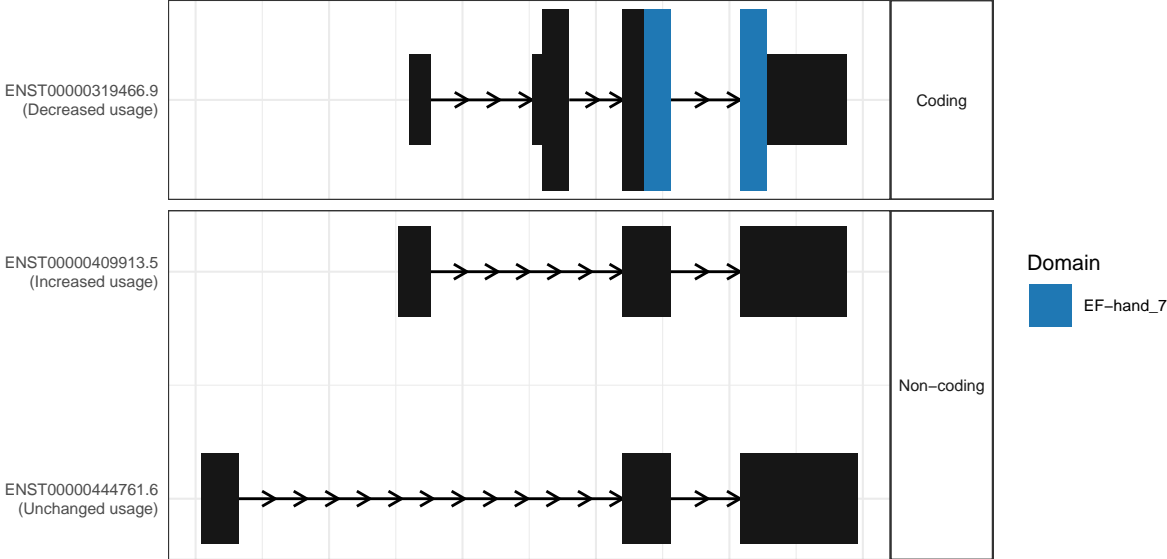


Domain



MPN_2A_DUB_like

The isoform switch in MCFD2 (FT vs RGC)

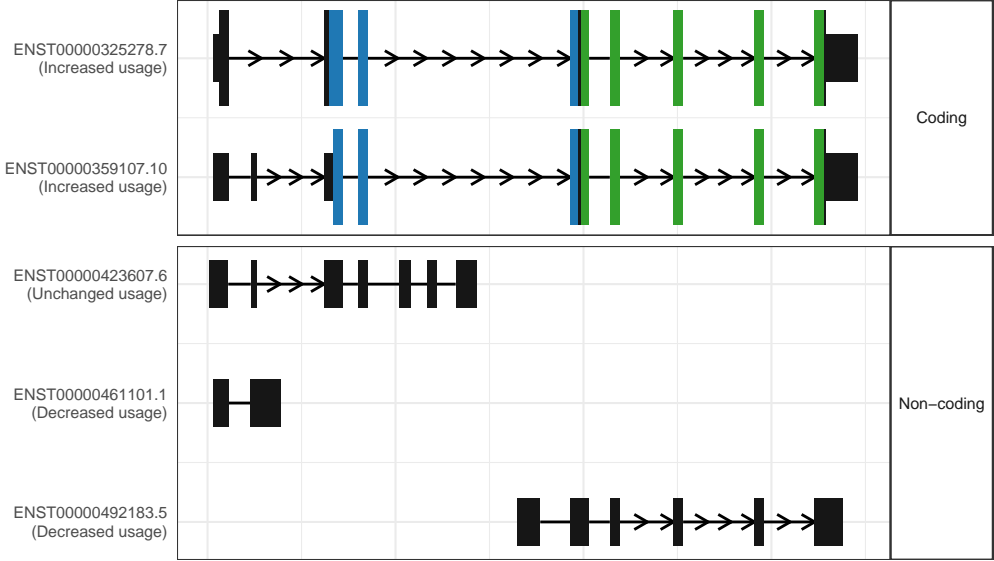


Domain

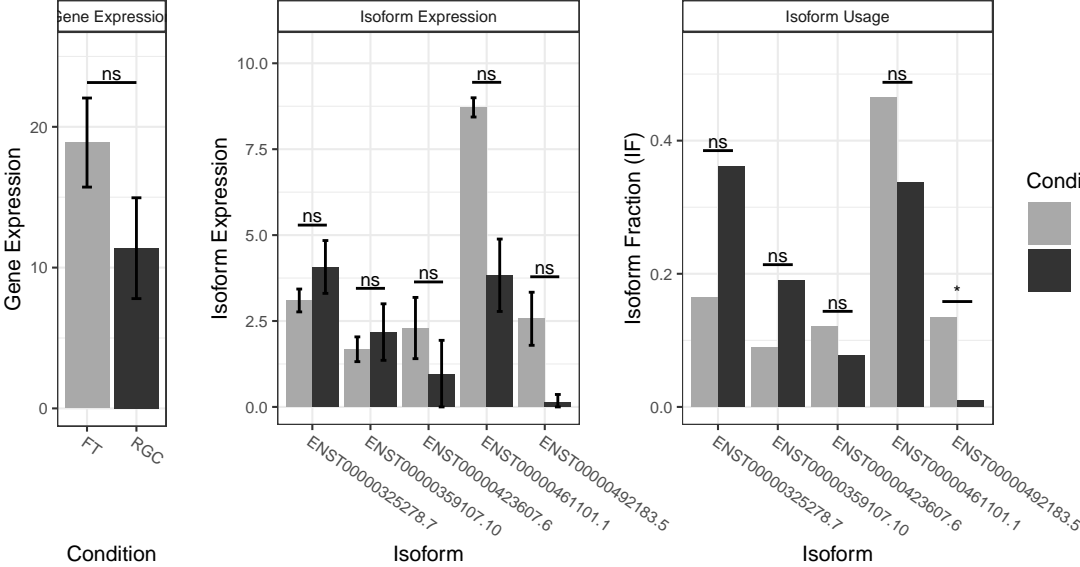
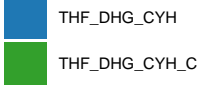


EF-hand_7

The isoform switch in MTHFD2L (FT vs RGC)



Domain



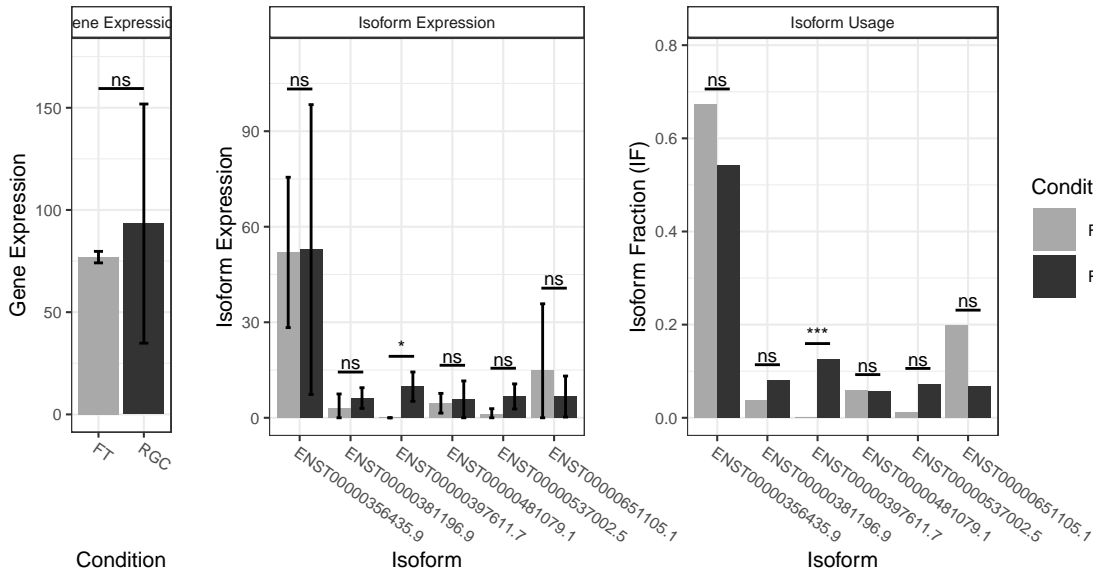
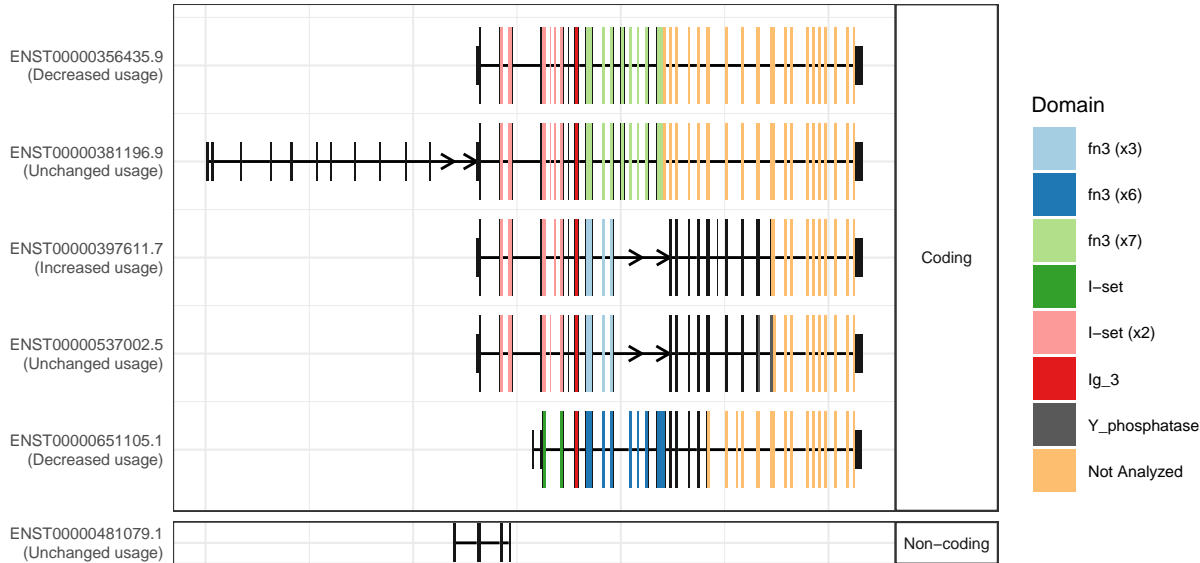
Domain



THF_DHG_CYH

THF_DHG_CYH_C

The isoform switch in PTPRD (FT vs RGC)



Domain



fn3 (x3)



fn3 (x6)



fn3 (x7)



I-set



I-set (x2)



Ig_3

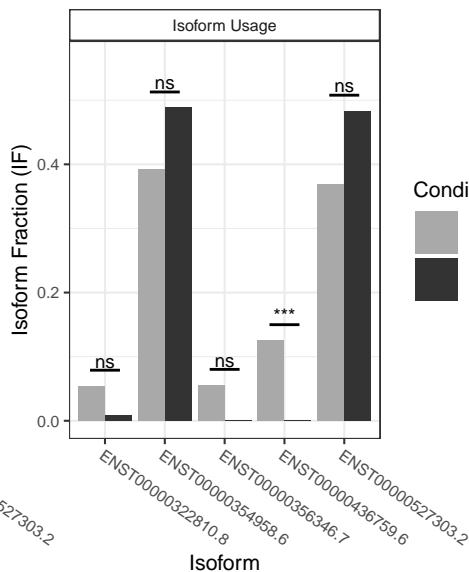
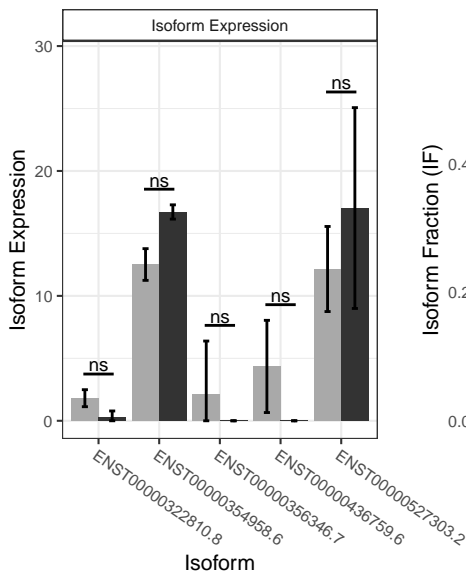
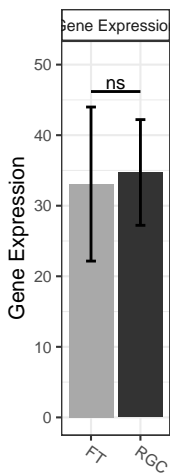
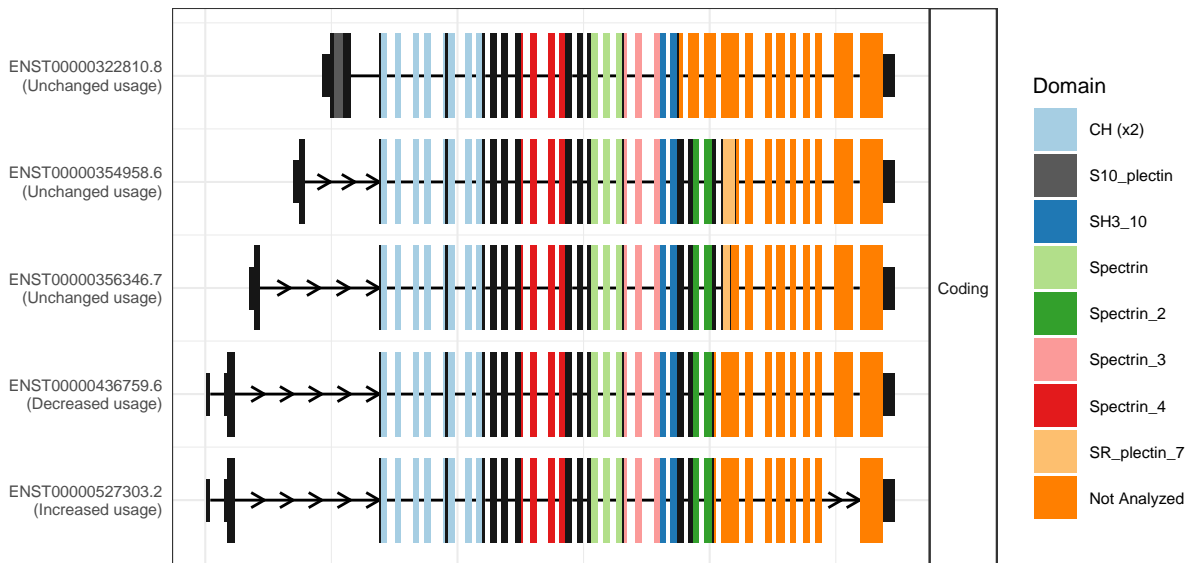


Y_phosphatase





Not Analyzed

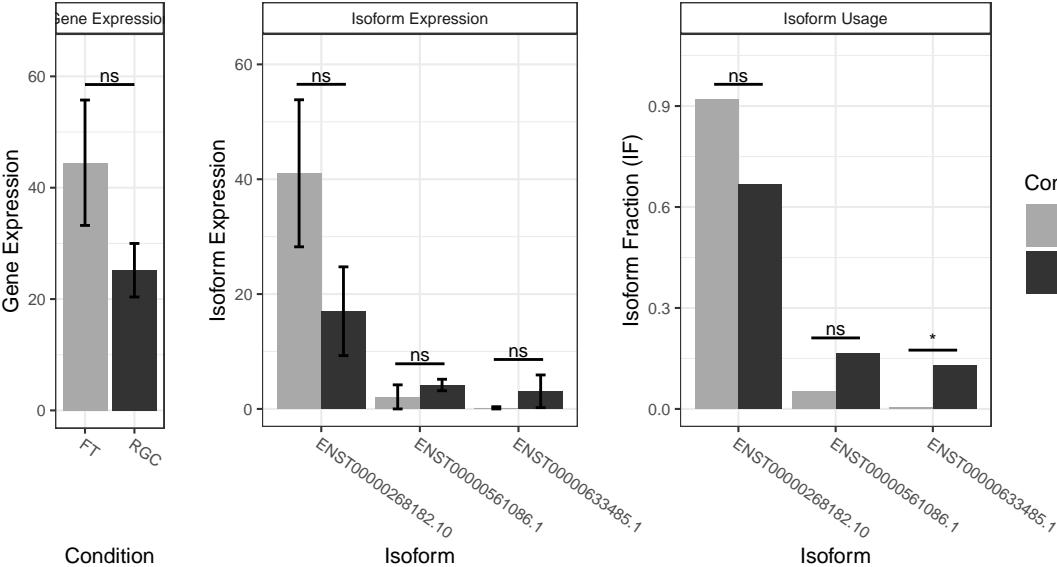
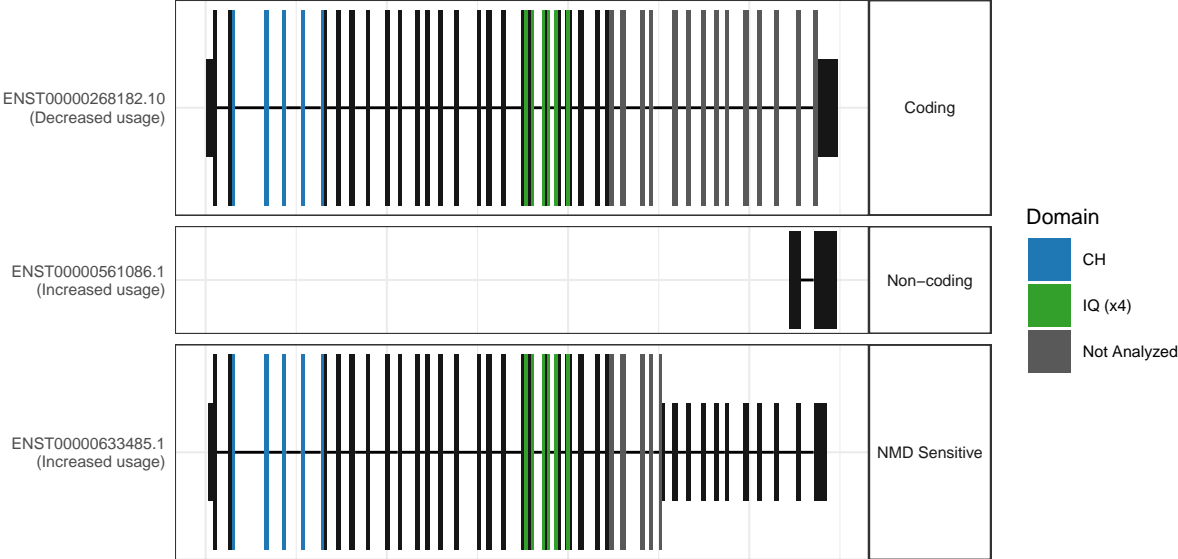
The isoform switch in PLEC (FT vs RGC)



Domain

	CH (x2)
	S10_plectin
	SH3_10
	Spectrin
	Spectrin_2
	Spectrin_3
	Spectrin_4
	SR_plectin_7
	Not Analyzed

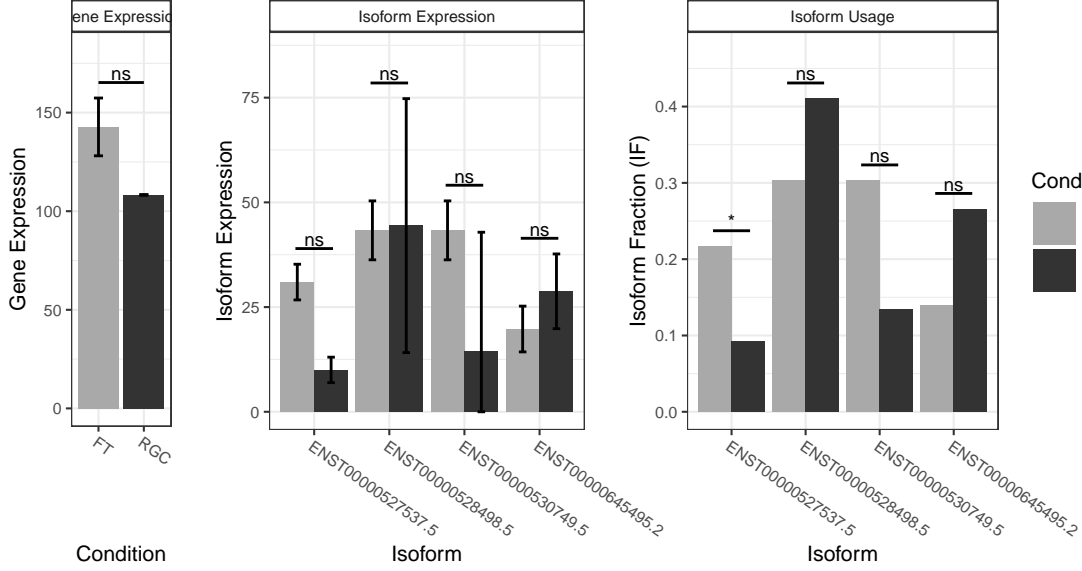
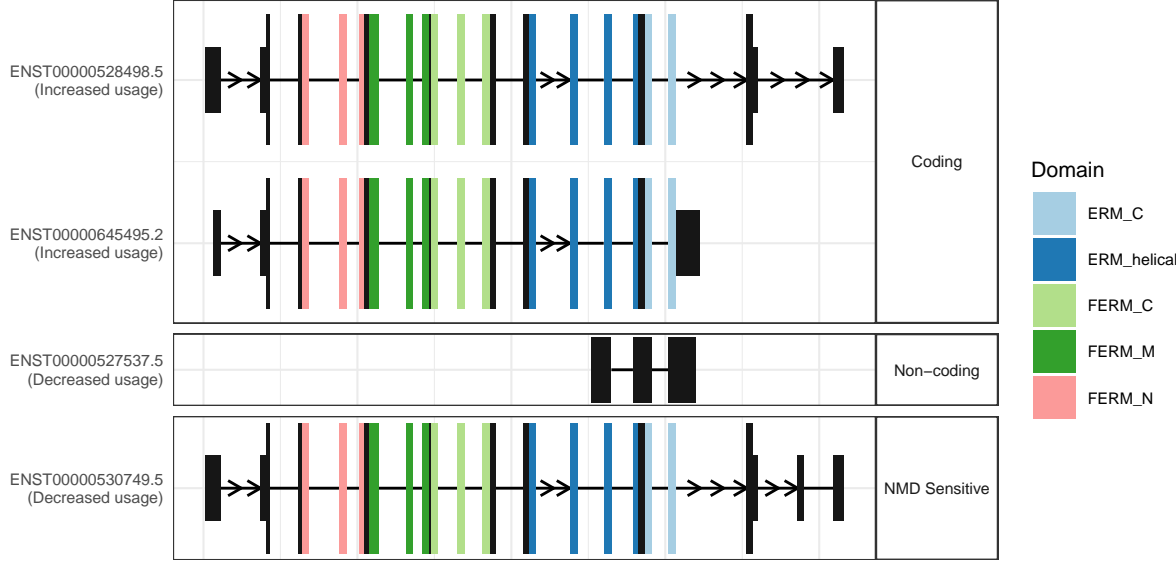
The isoform switch in IQGAP1 (FT vs RGC)








Domain

	CH
	IQ (x4)
	Not Analyzed

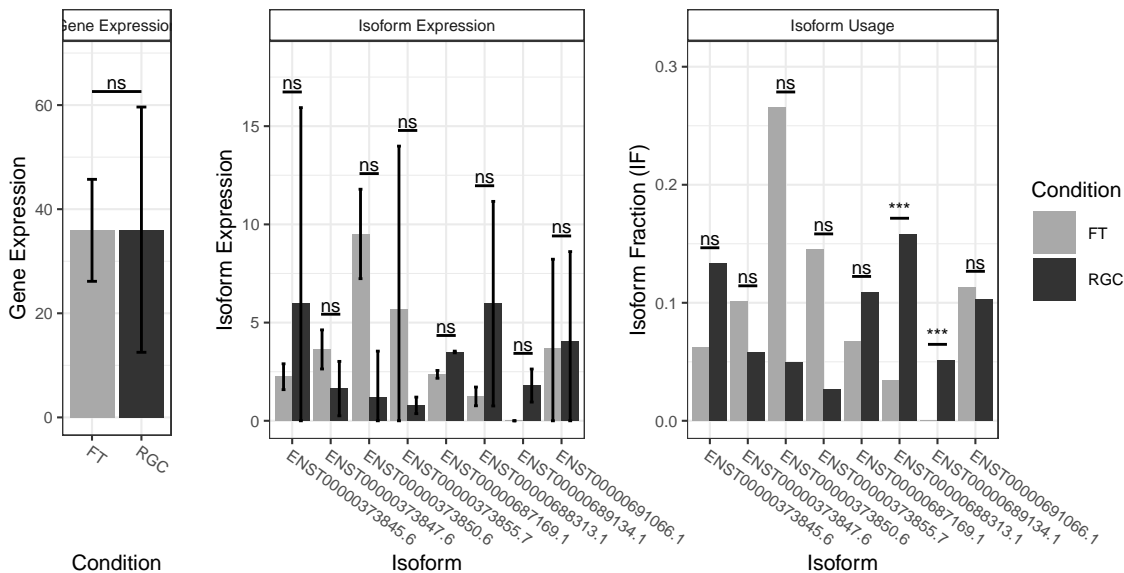
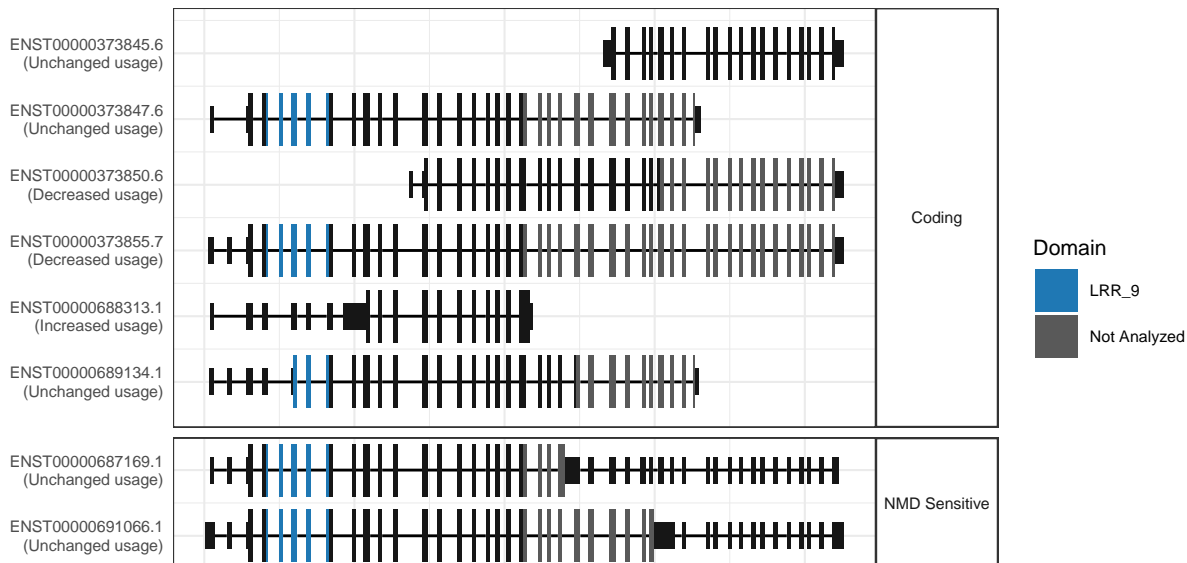
The isoform switch in RDX (FT vs RGC)



Domain

	ERM_C
	ERM_helical
	FERM_C
	FERM_M
	FERM_N

The isoform switch in CNTRL (FT vs RGC)



Domain

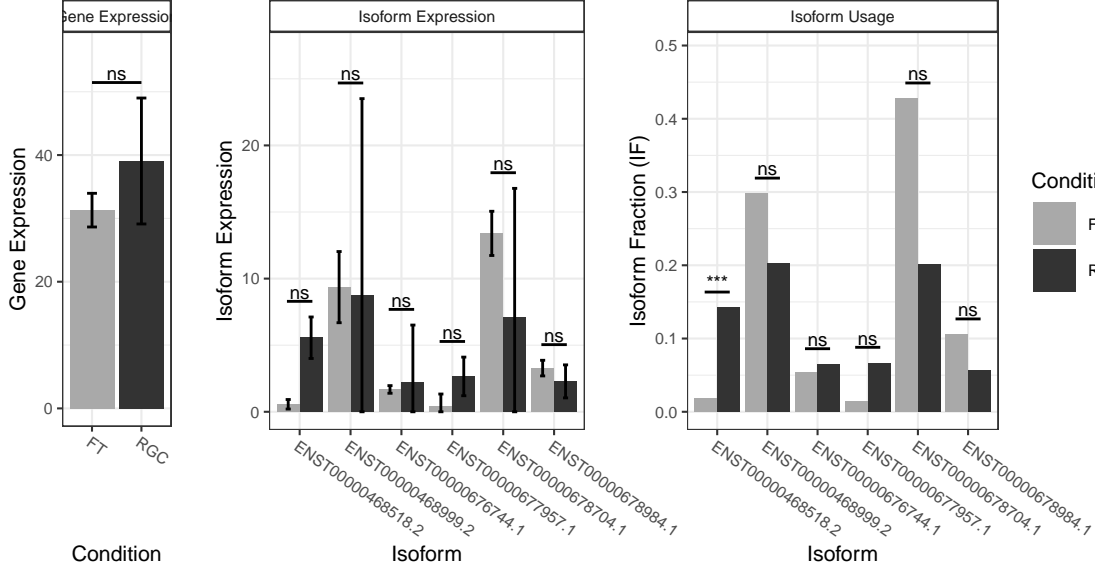
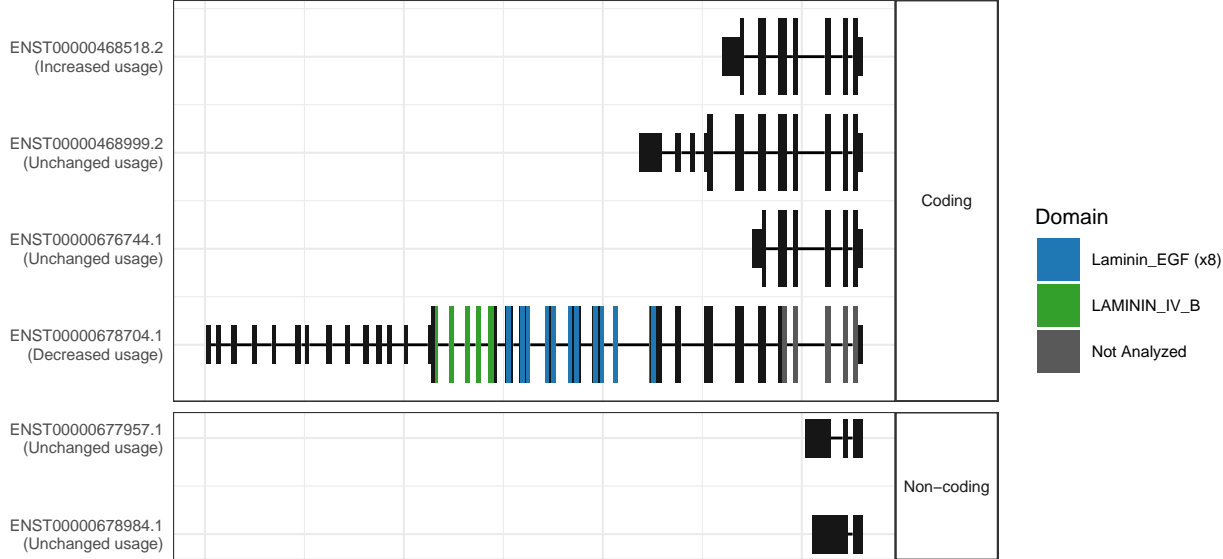


LRR_9






Not Analyzed

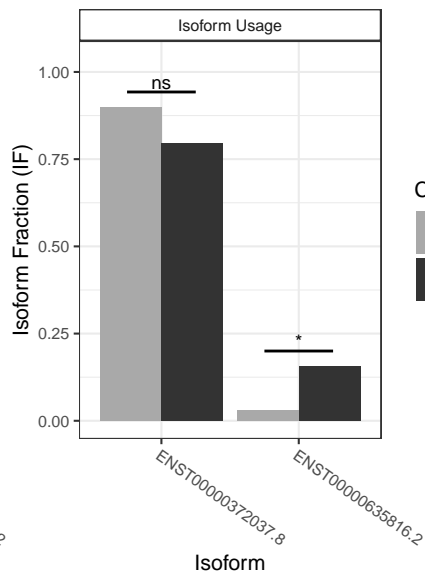
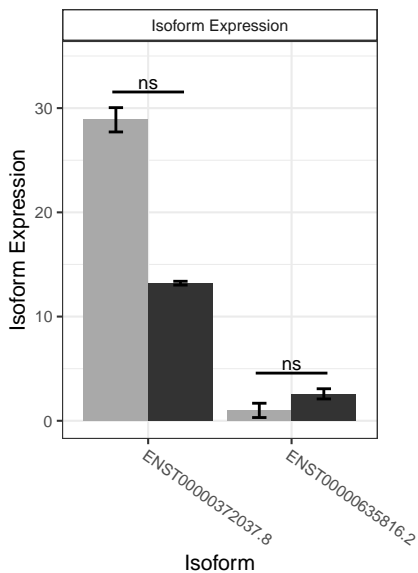
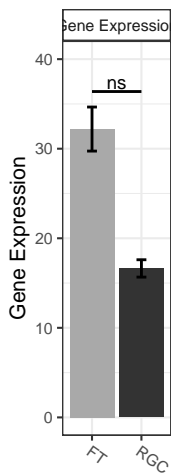
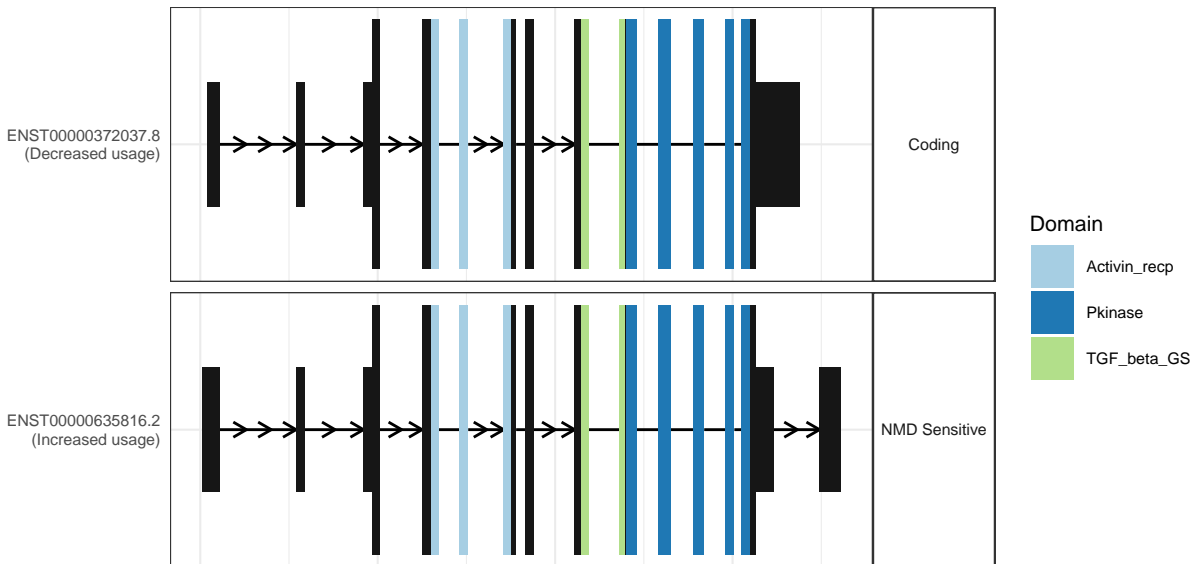
The isoform switch in LAMB1 (FT vs RGC)



Domain

	Laminin_EGF (x8)
	LAMININ_IV_B
	Not Analyzed

The isoform switch in BMPR1A (FT vs RGC)



Domain



Activin_recp

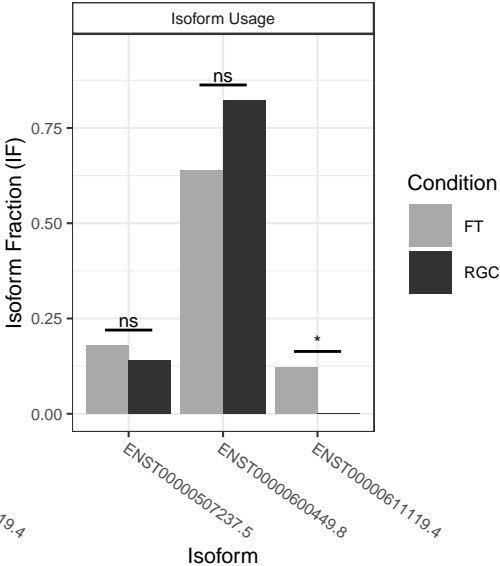
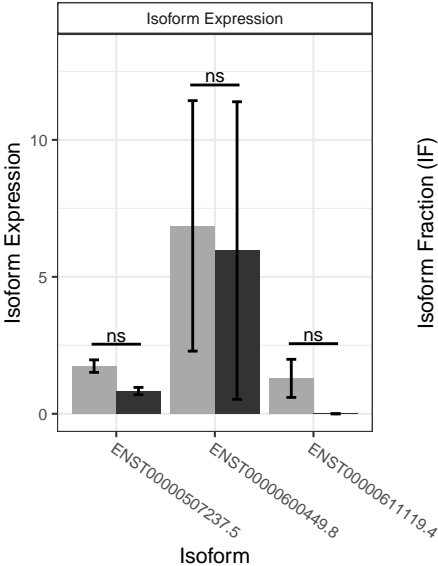
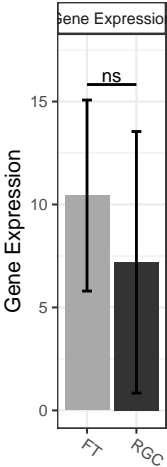
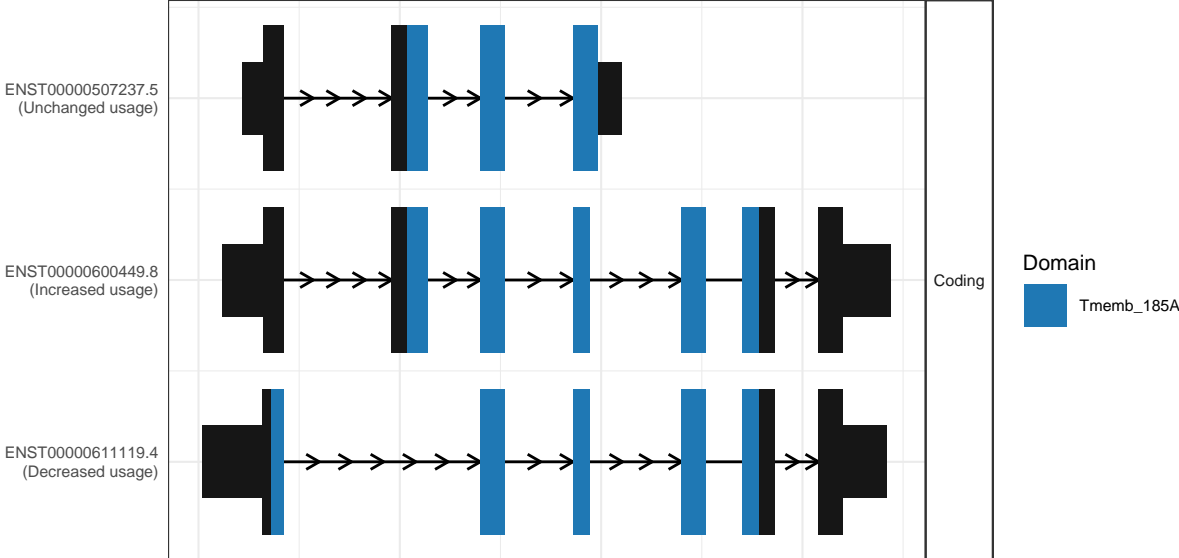


Pkinase



TGF_beta_GS

The isoform switch in TMEM185A (FT vs RGC)

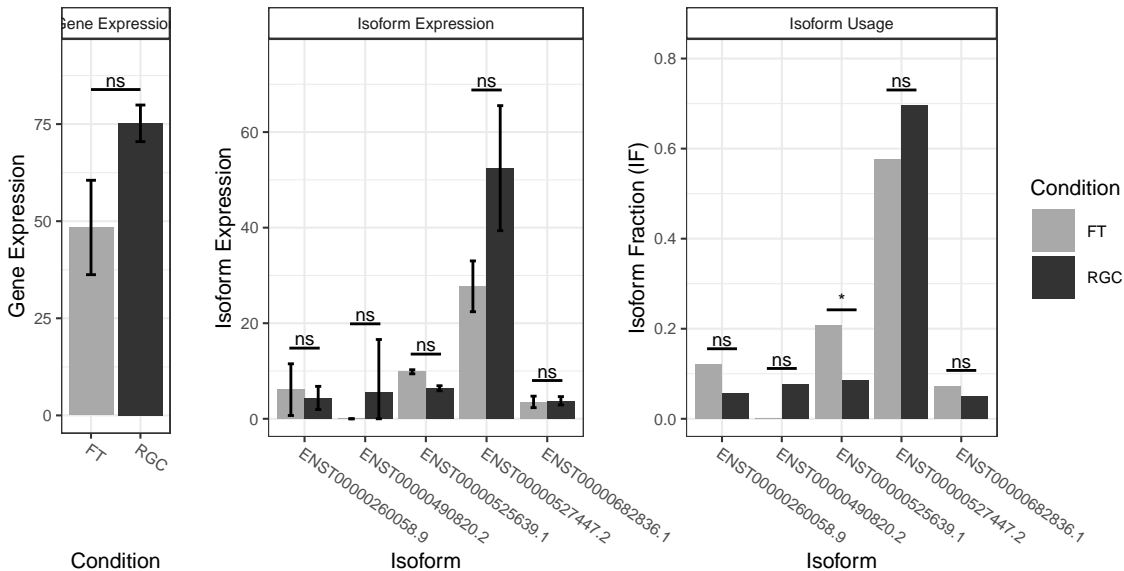
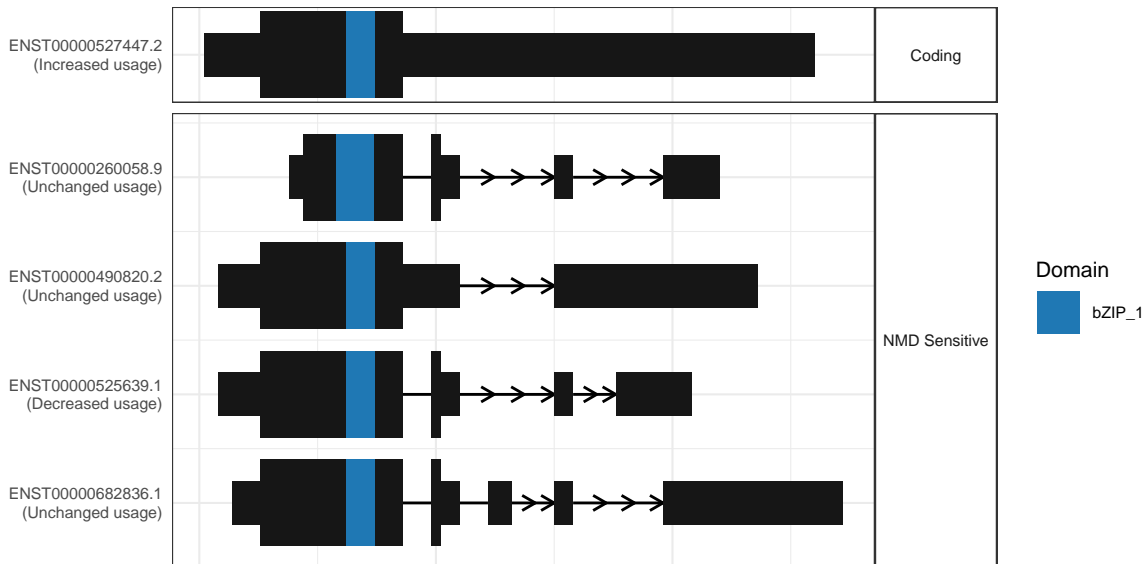


Domain



Tmemb_185A

The isoform switch in CREBZF (FT vs RGC)

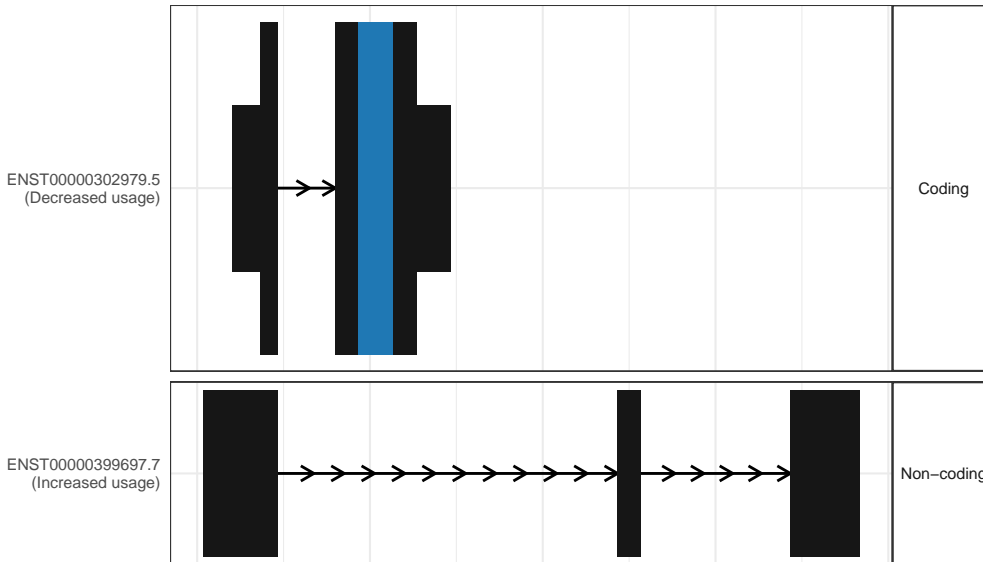


Domain



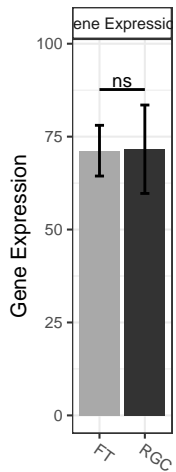
bZIP_1

The isoform switch in POLR1D (FT vs RGC)

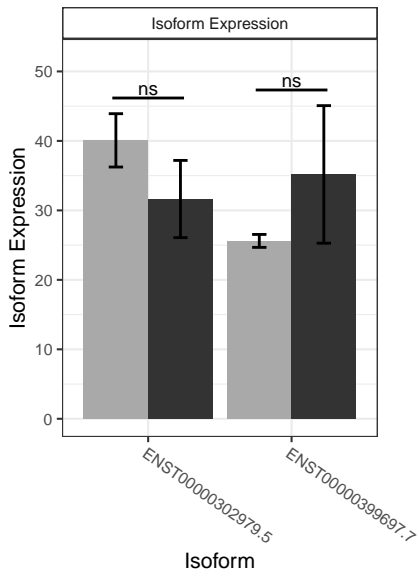


Domain

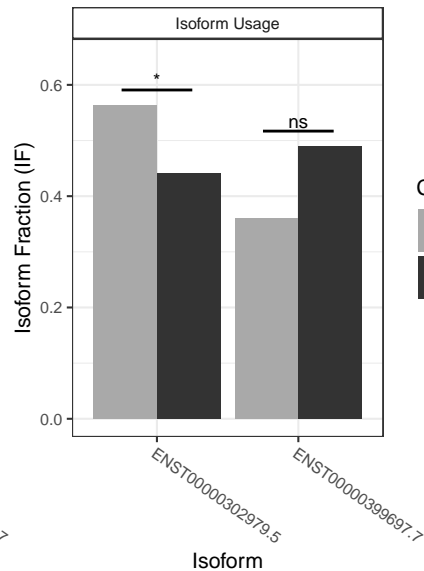
RNA_pol_L_2



Condition



Isoform



Isoform

Condition

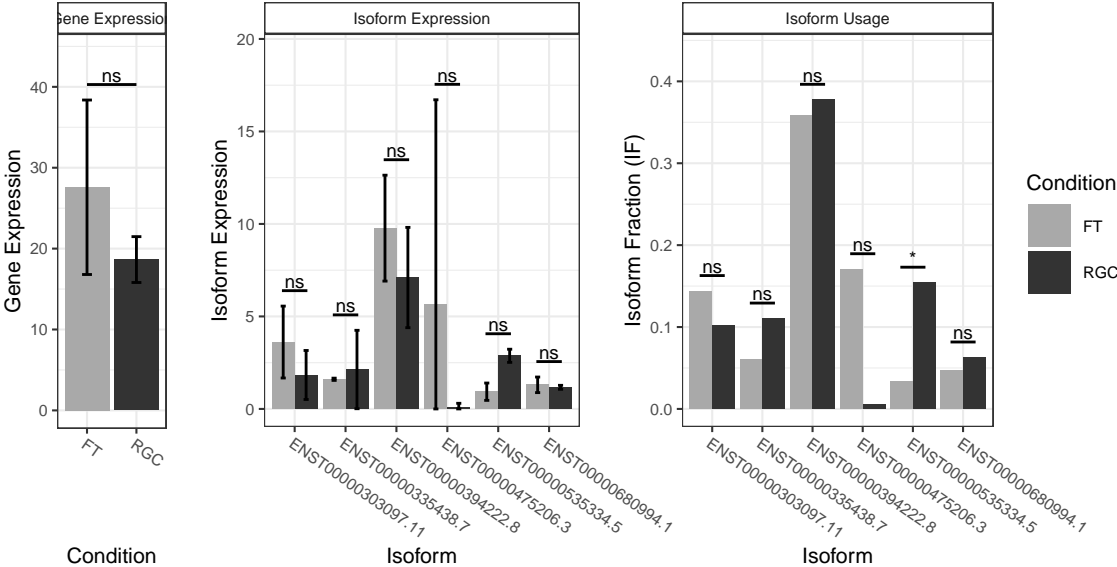
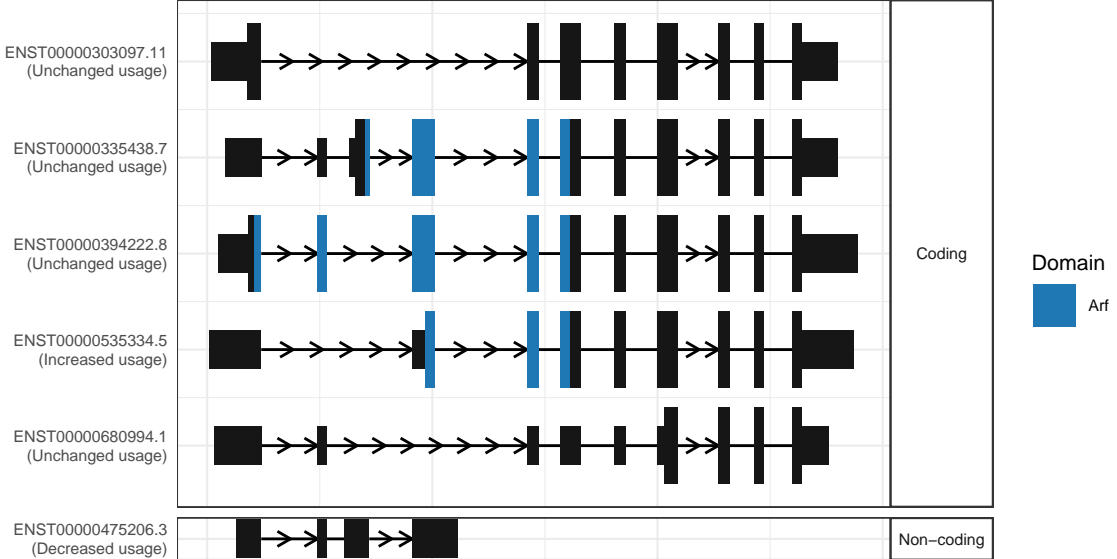
FT
RGC

Domain



RNA_pol_L_2

The isoform switch in ARL13B (FT vs RGC)

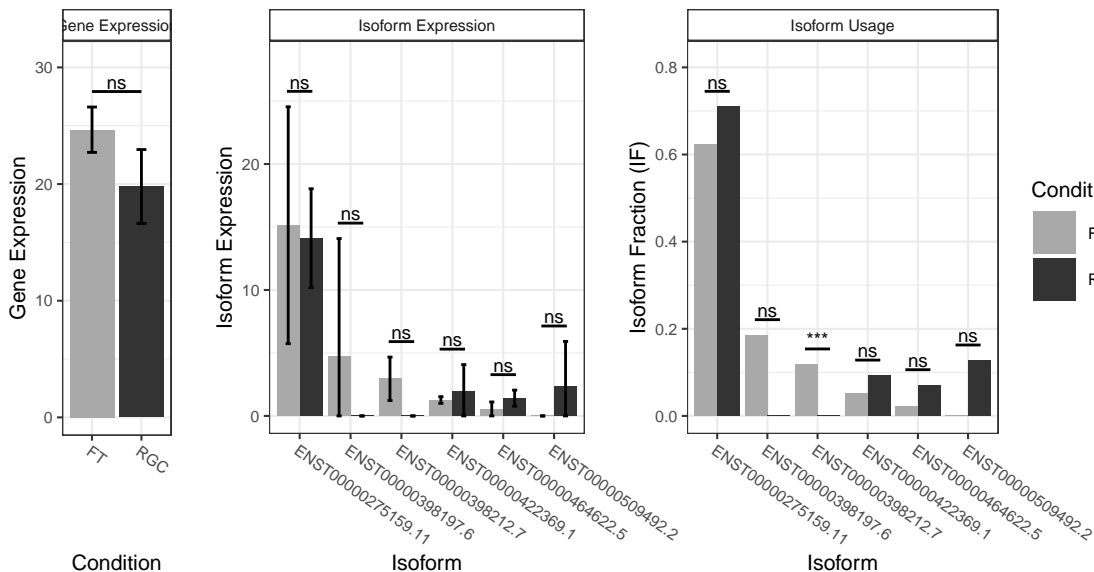
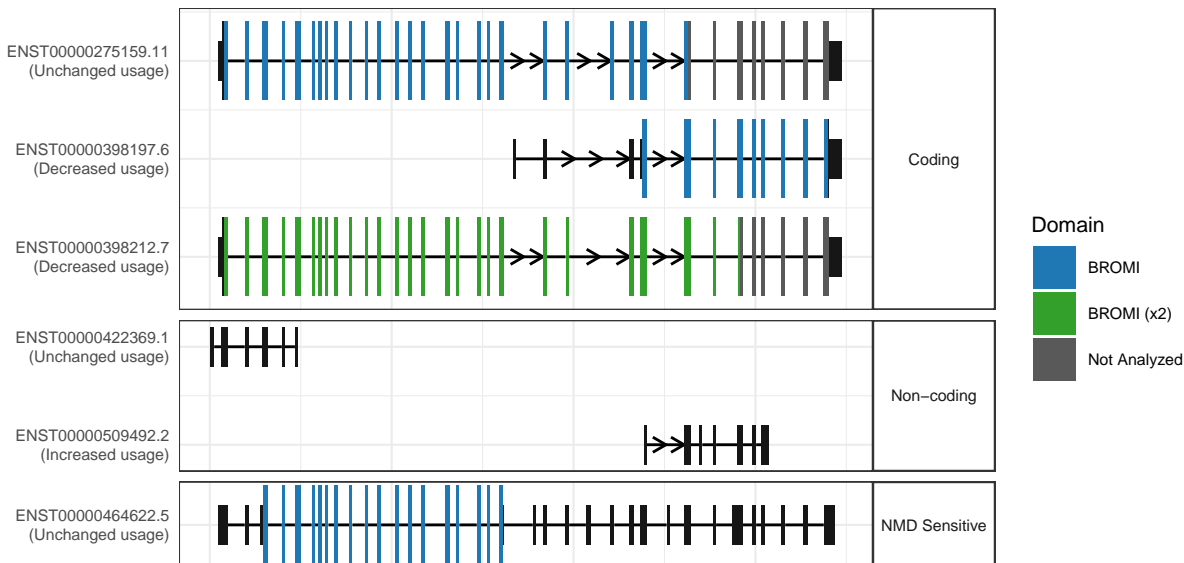


Domain

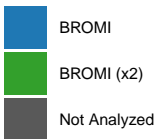


Arf

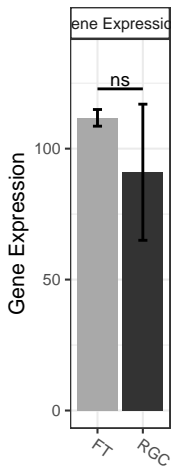
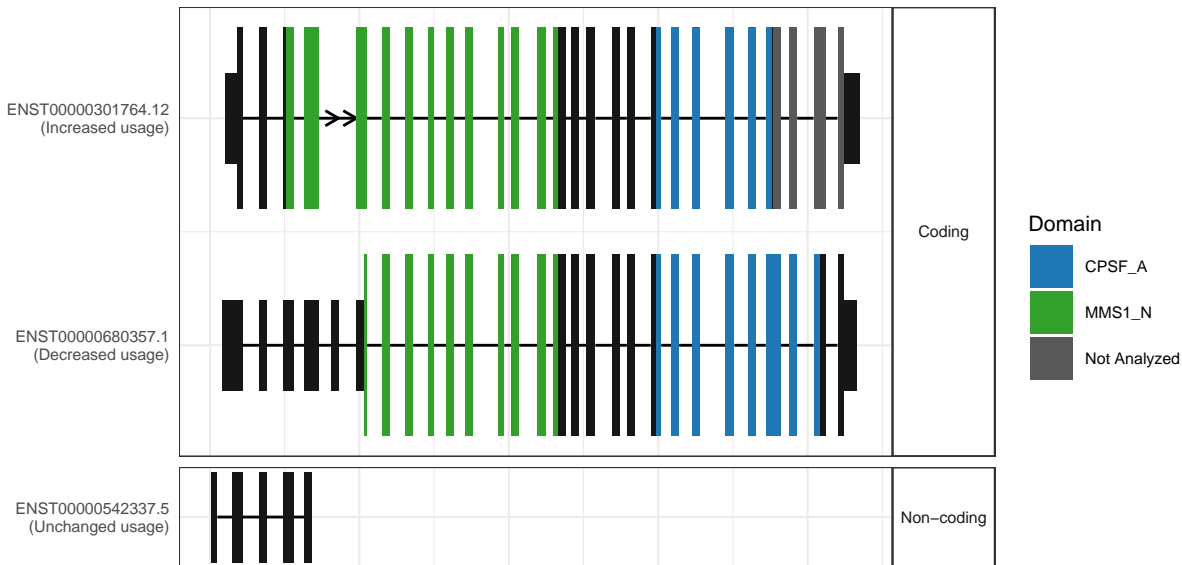
The isoform switch in TBC1D32 (FT vs RGC)



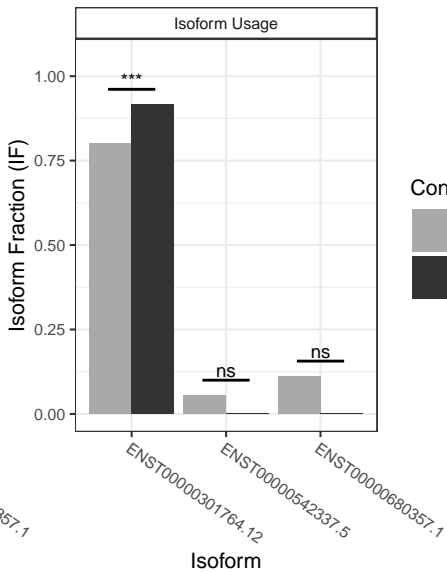
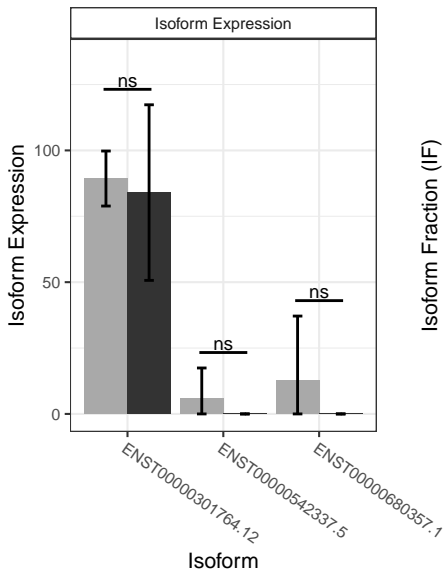
Domain



The isoform switch in DDB1 (FT vs RGC)



Condition



Domain



CPSF_A

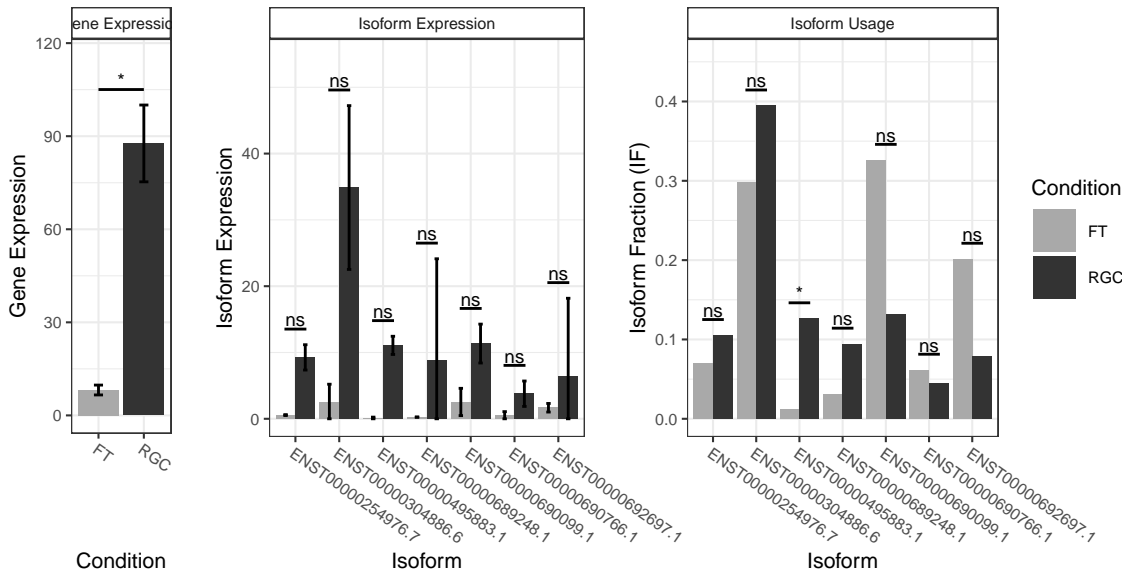
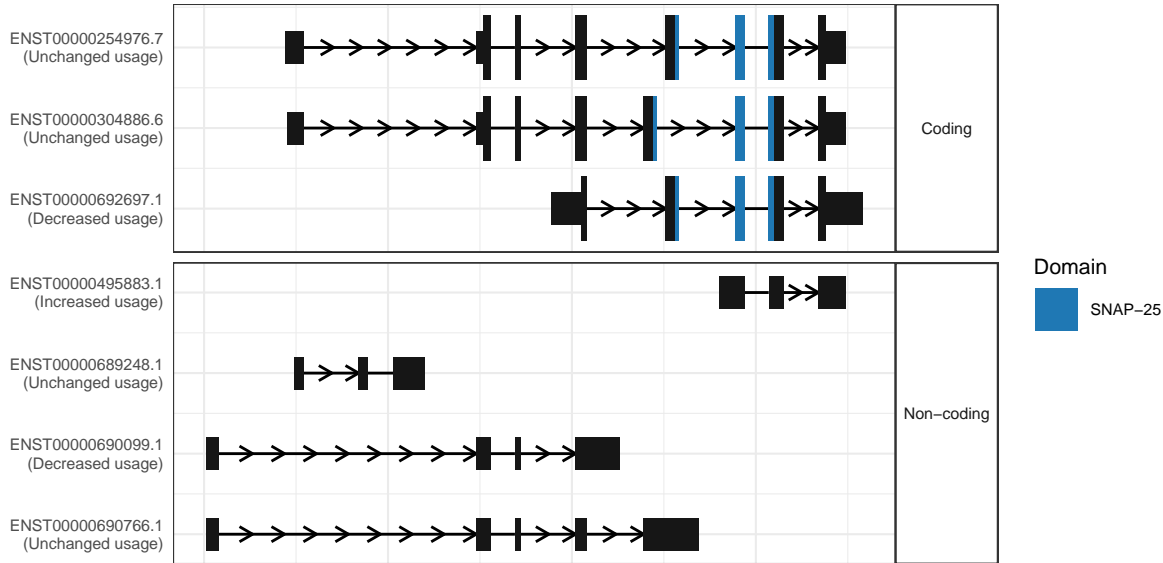


MMS1_N



Not Analyzed

The isoform switch in SNAP25 (FT vs RGC)

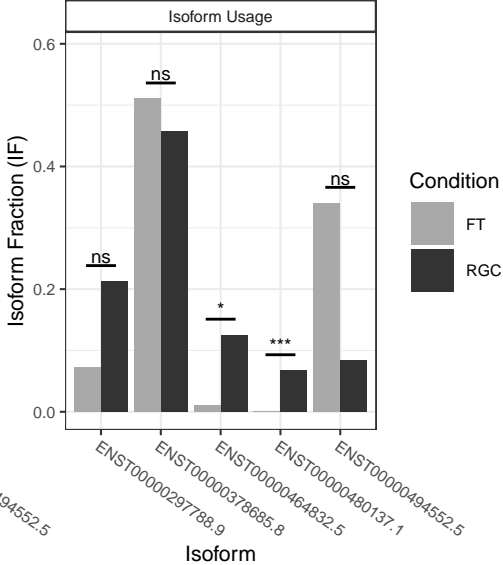
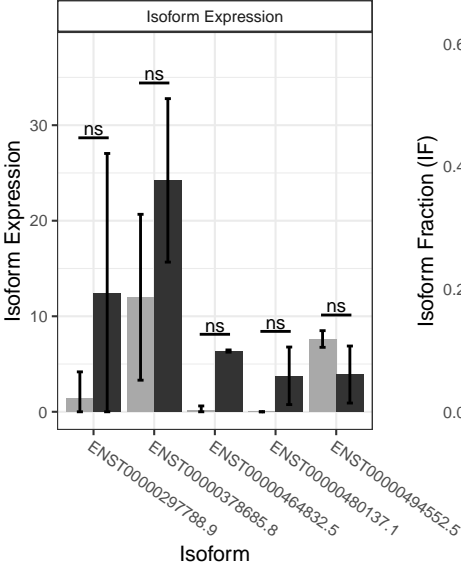
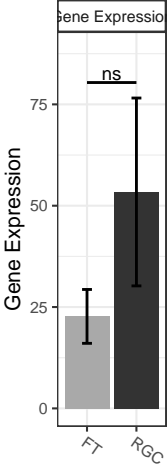
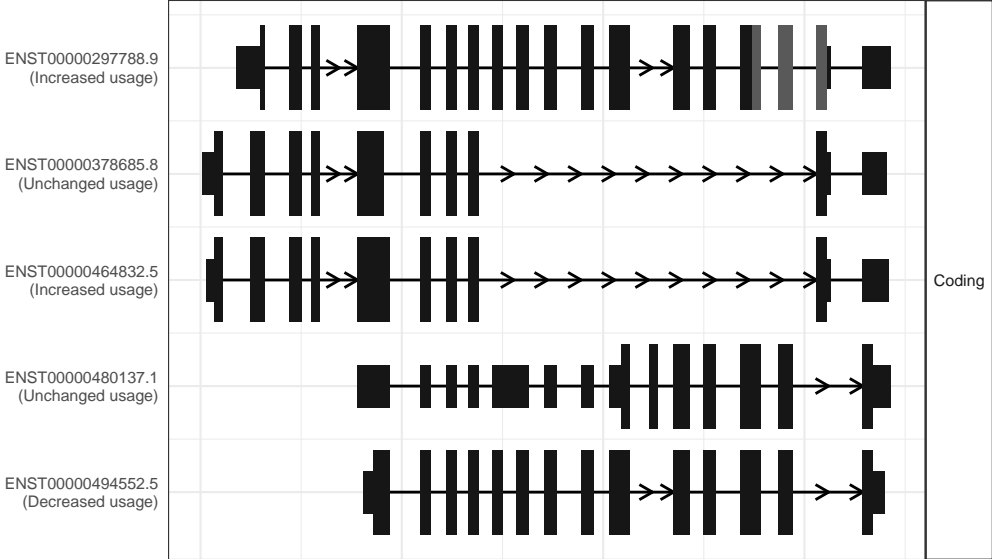


Domain

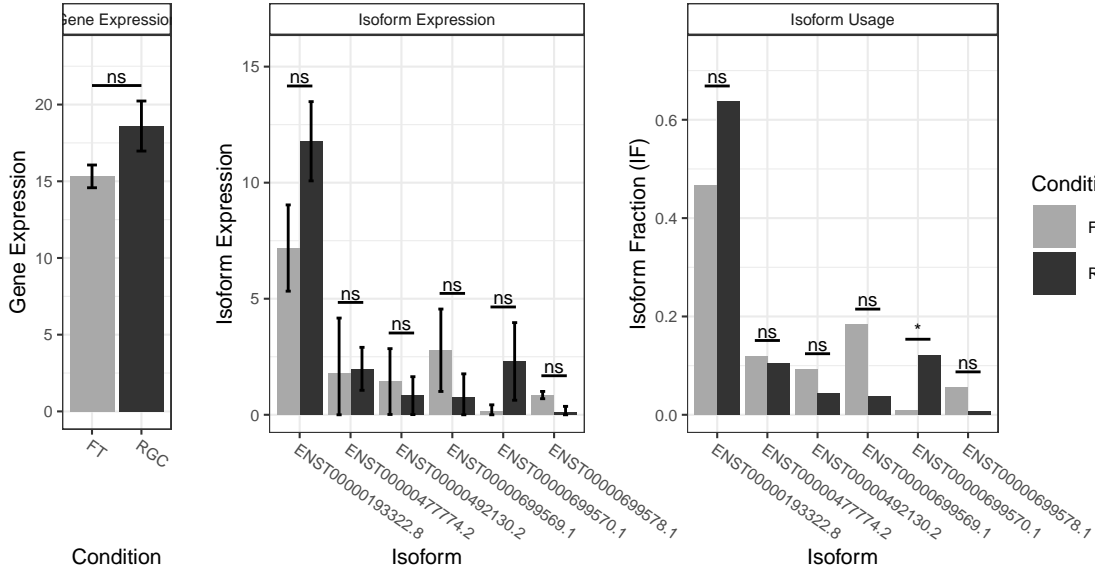
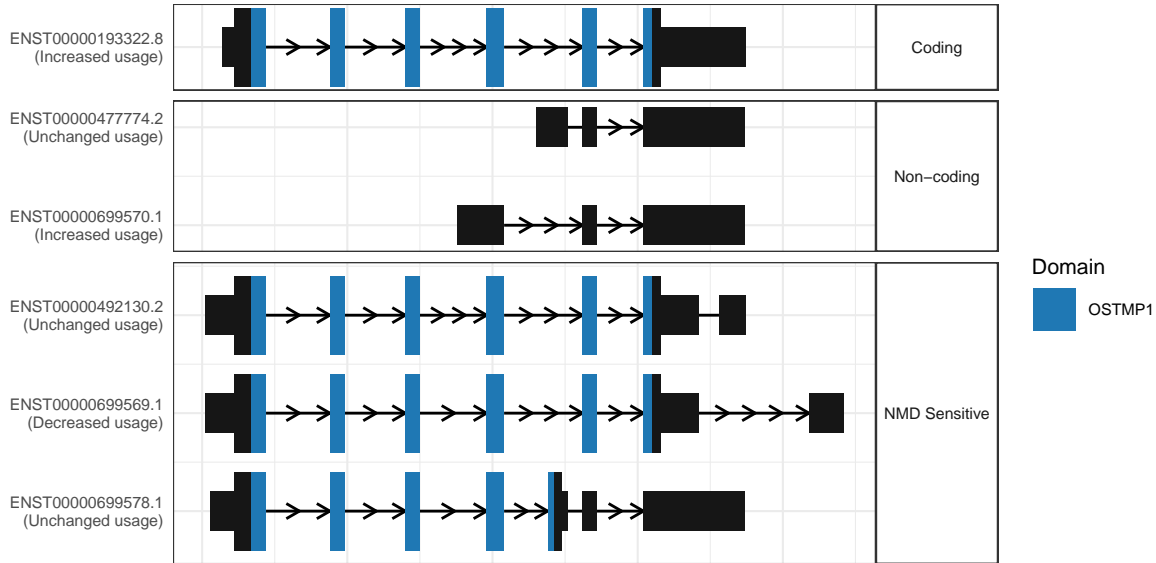


SNAP-25

The isoform switch in CCDC136 (FT vs RGC)



The isoform switch in OSTM1 (FT vs RGC)

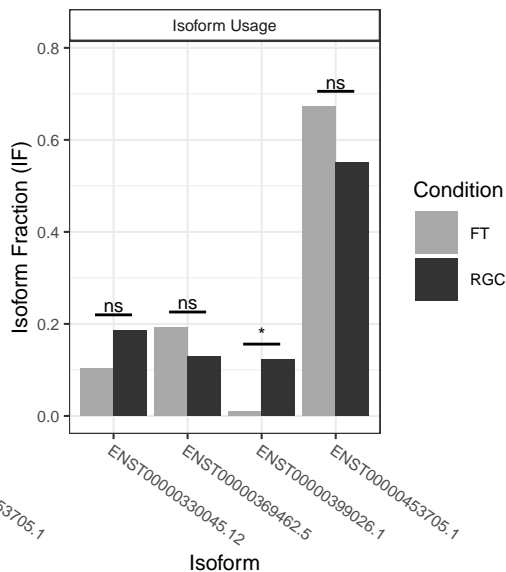
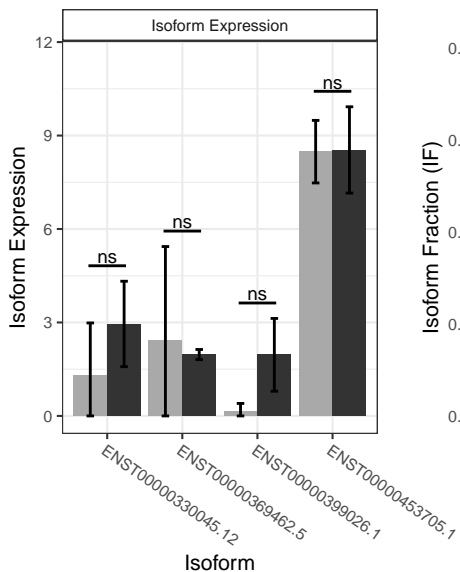
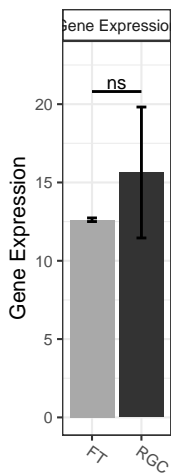
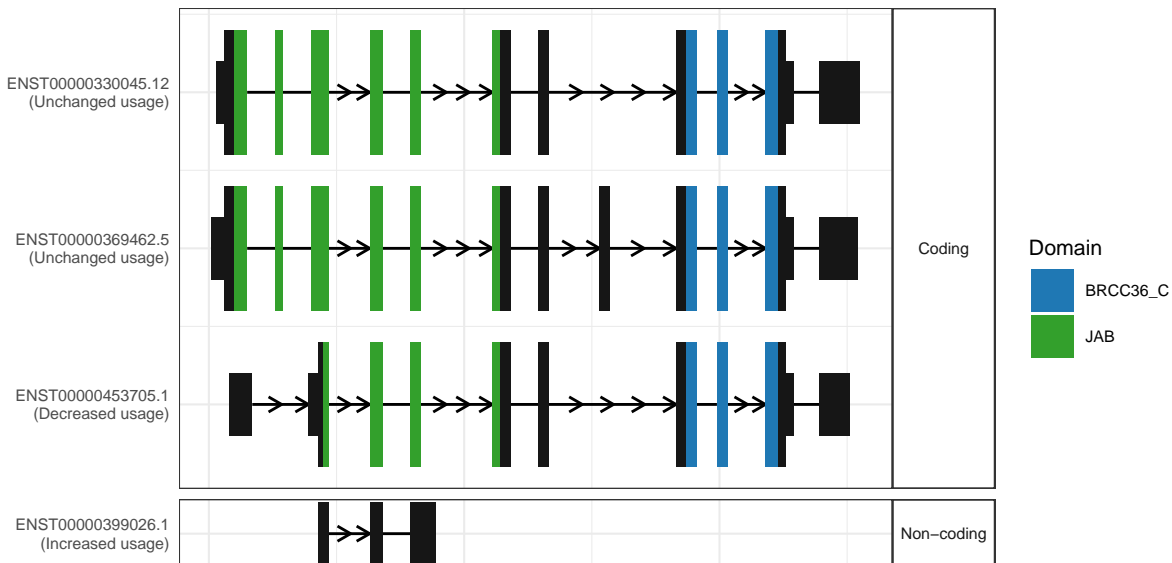


Domain



OSTMP1

The isoform switch in BRCC3 (FT vs RGC)



Condition

Isoform

Isoform

Condition



Domain

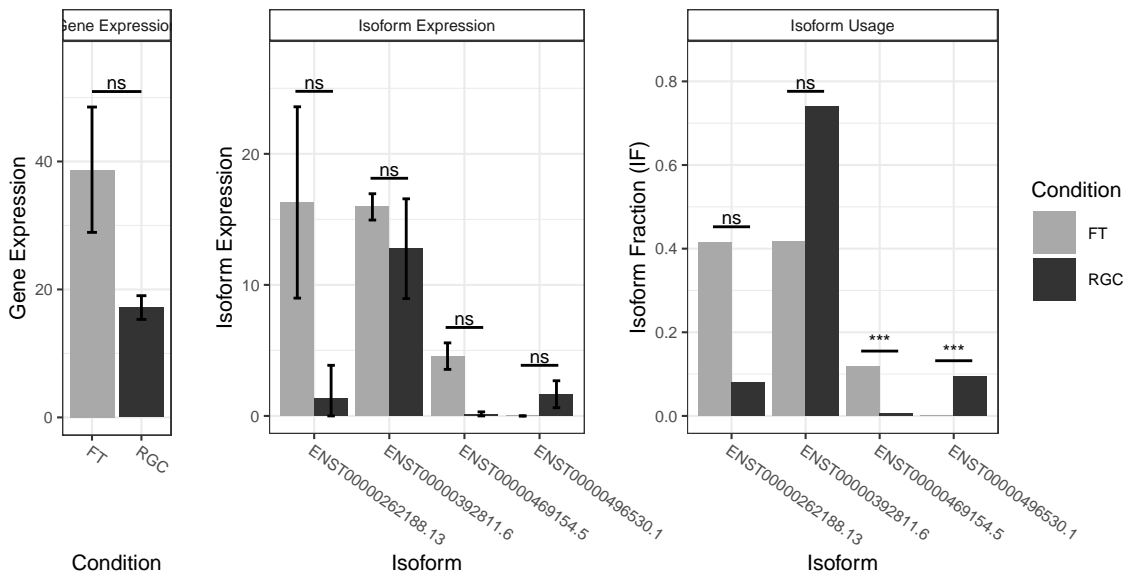
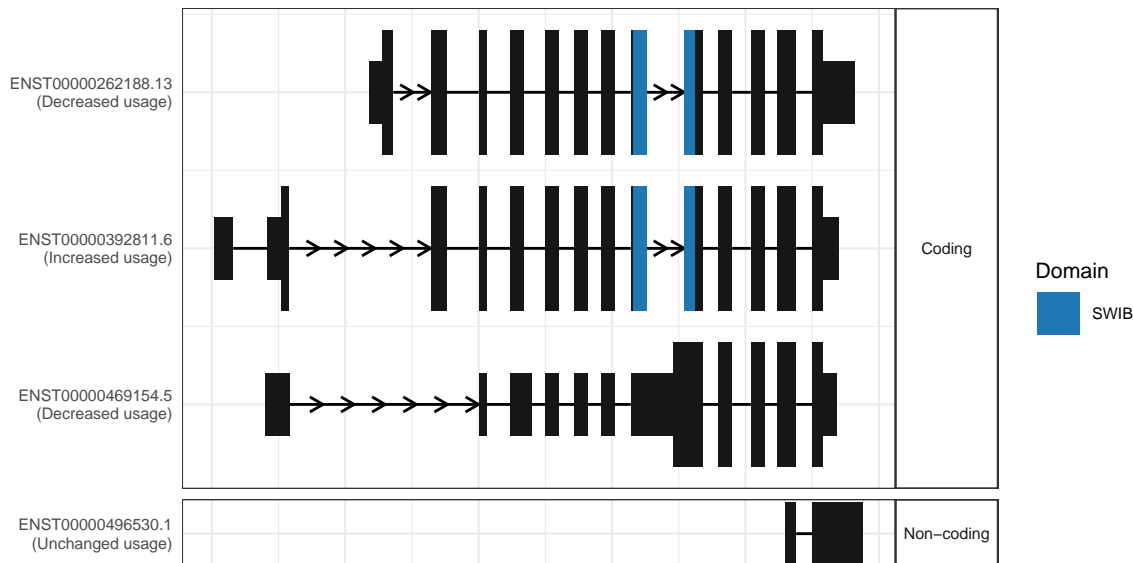


BRCC36_C



JAB

The isoform switch in SMARCD3 (FT vs RGC)

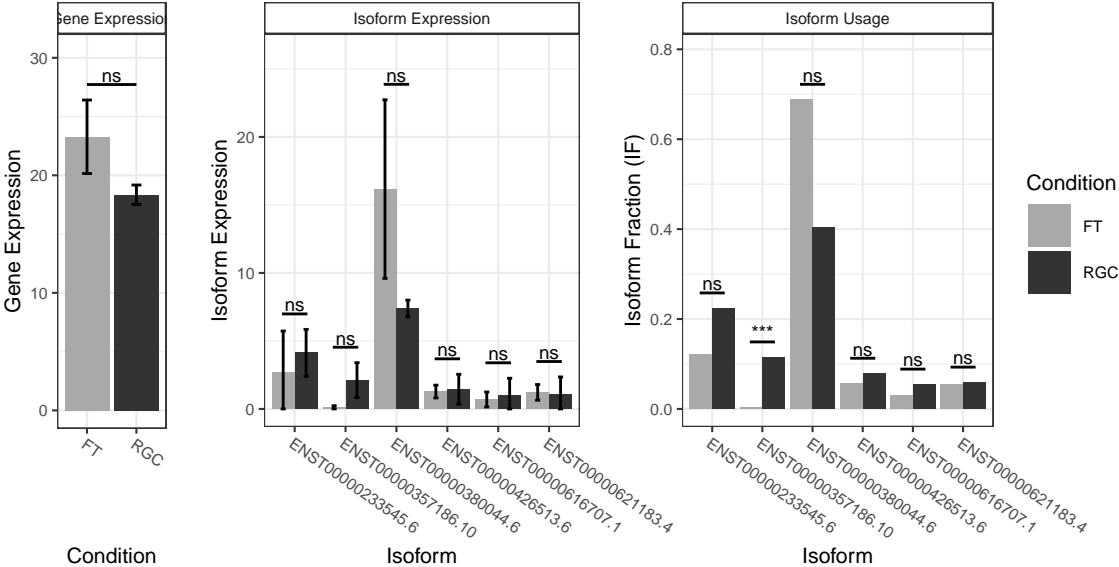
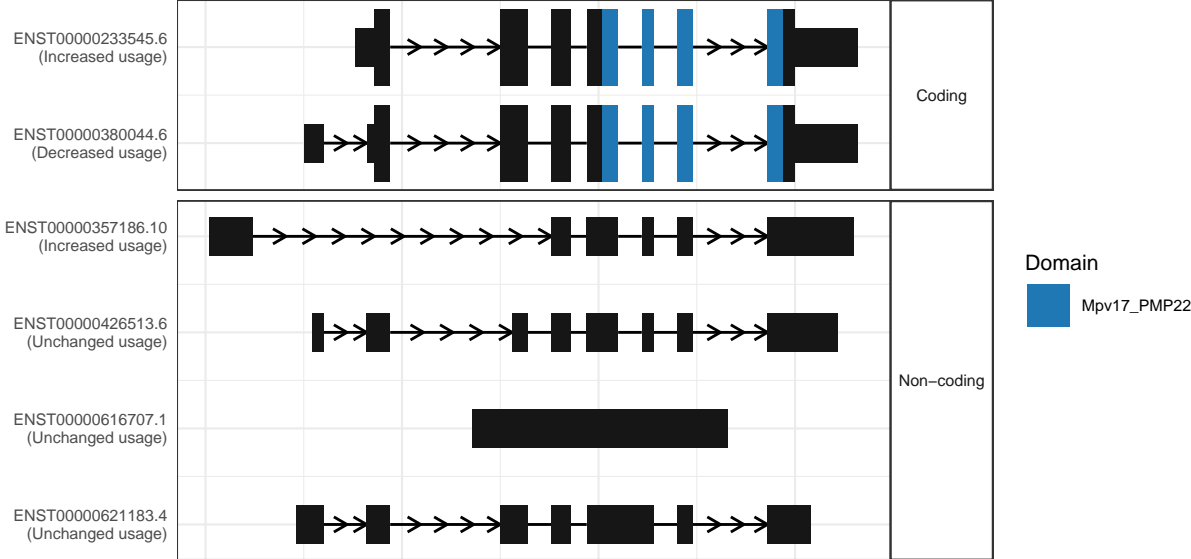


Domain



SWIB

The isoform switch in MPV17 (FT vs RGC)

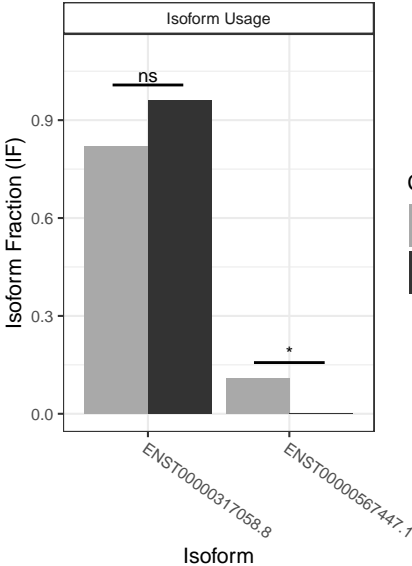
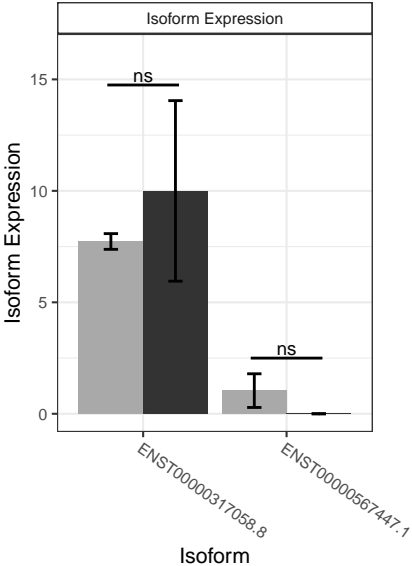
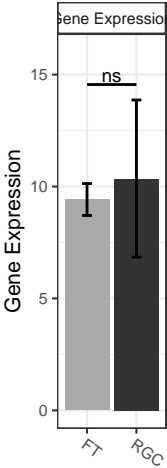
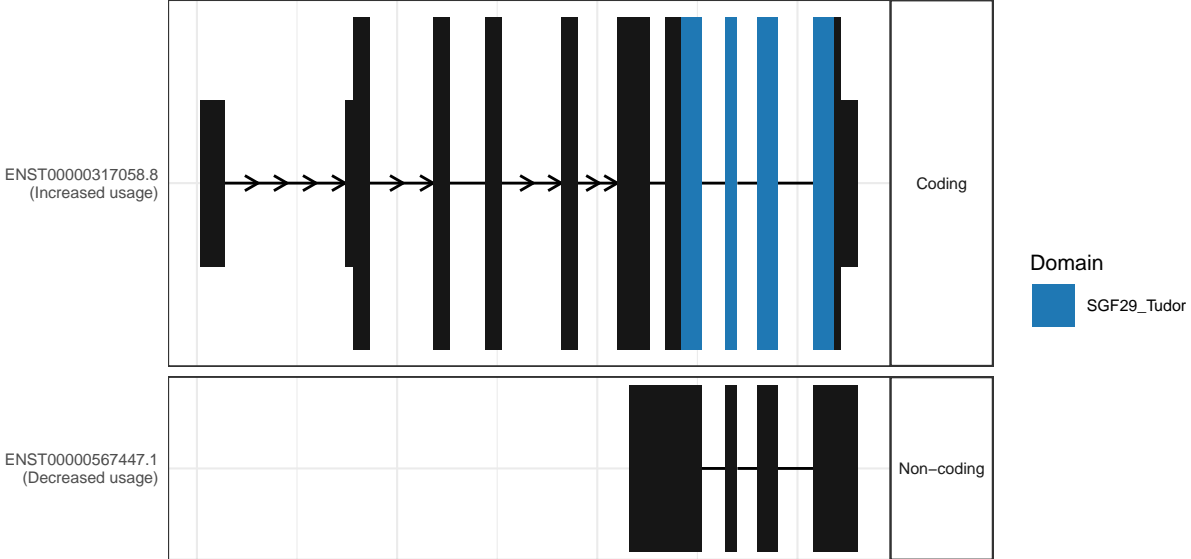


Domain



Mpv17_PMP22

The isoform switch in SGF29 (FT vs RGC)

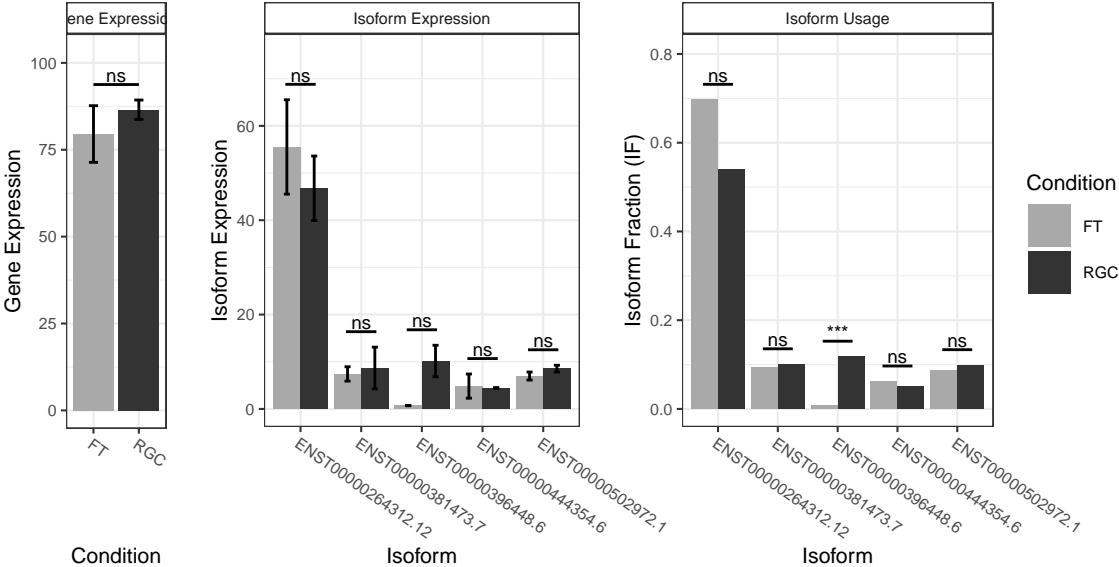
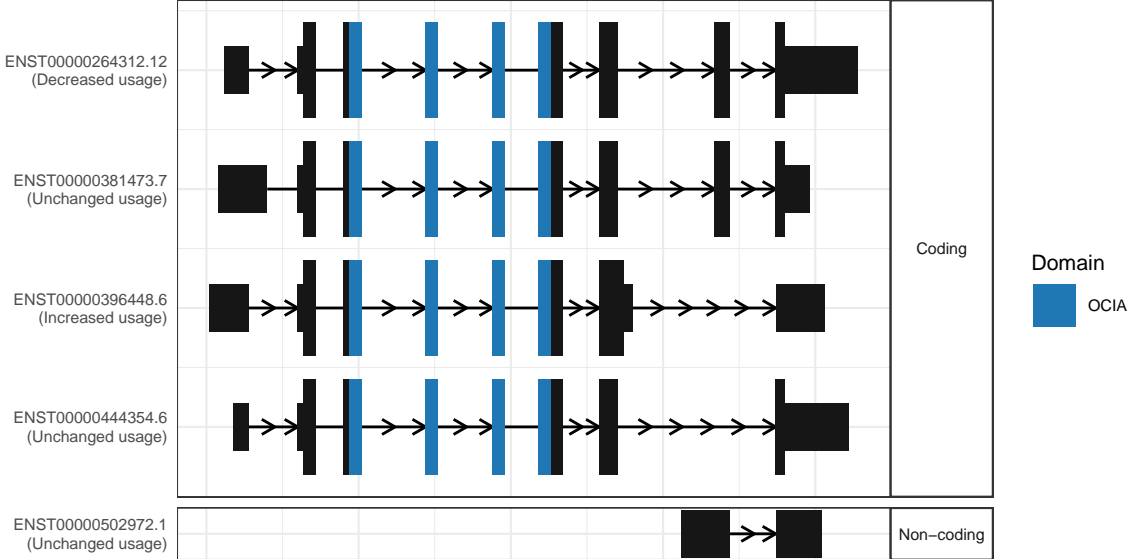


Domain



SGF29_Tudor

The isoform switch in OCIAD1 (FT vs RGC)

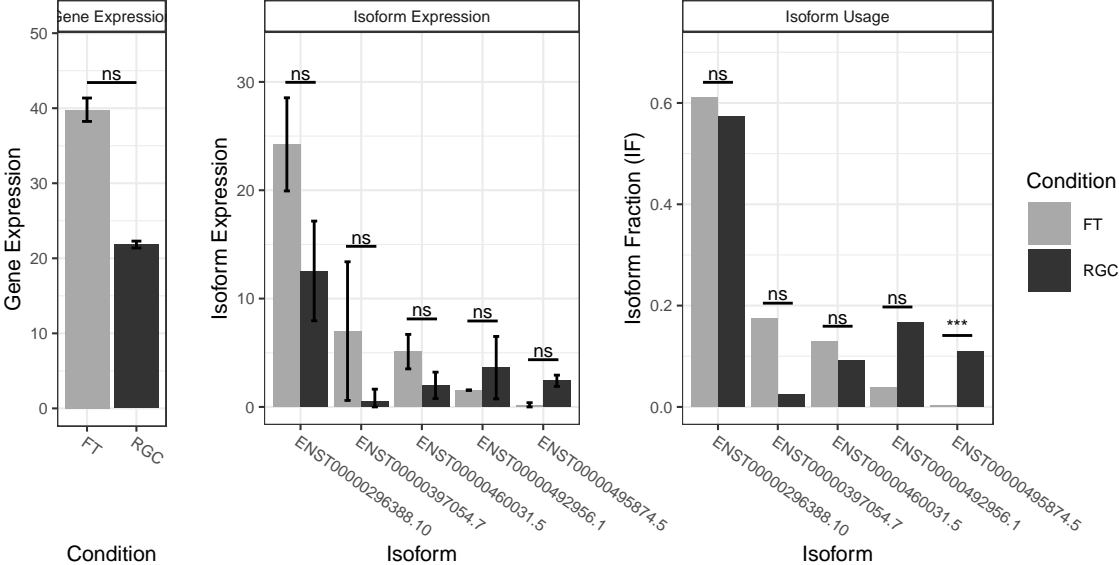
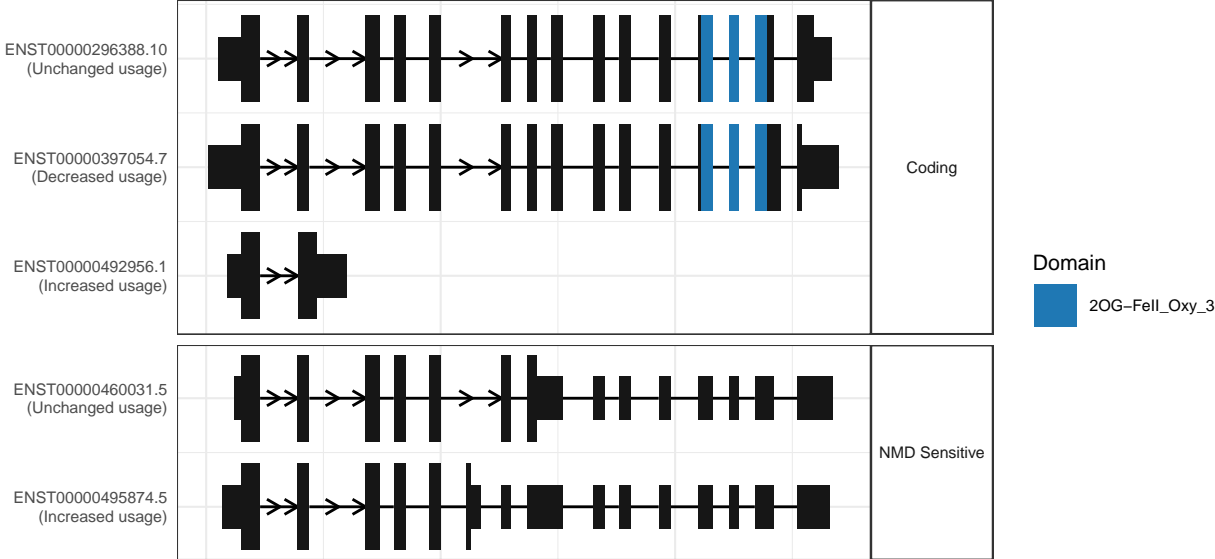


Domain



OCIA

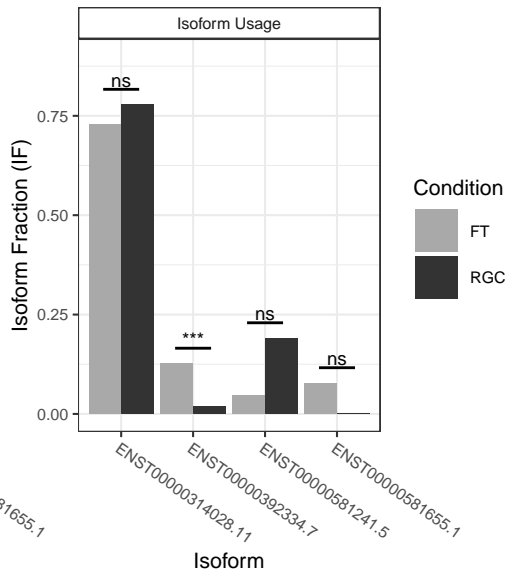
The isoform switch in P3H1 (FT vs RGC)



Domain

 2OG-Fell_Oxy_3

	Coding
	Non-coding
	NMD Sensitive

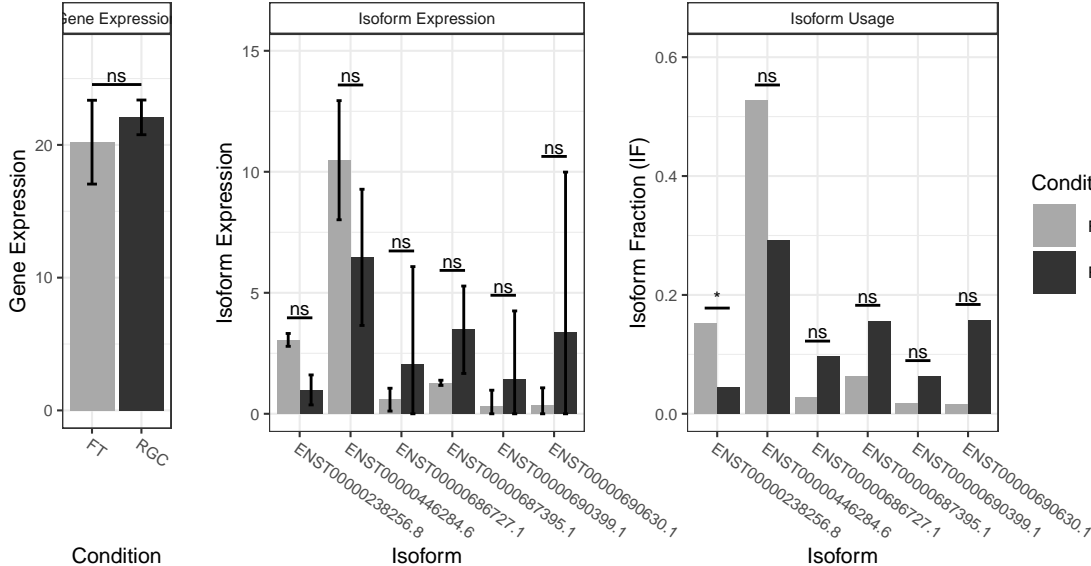
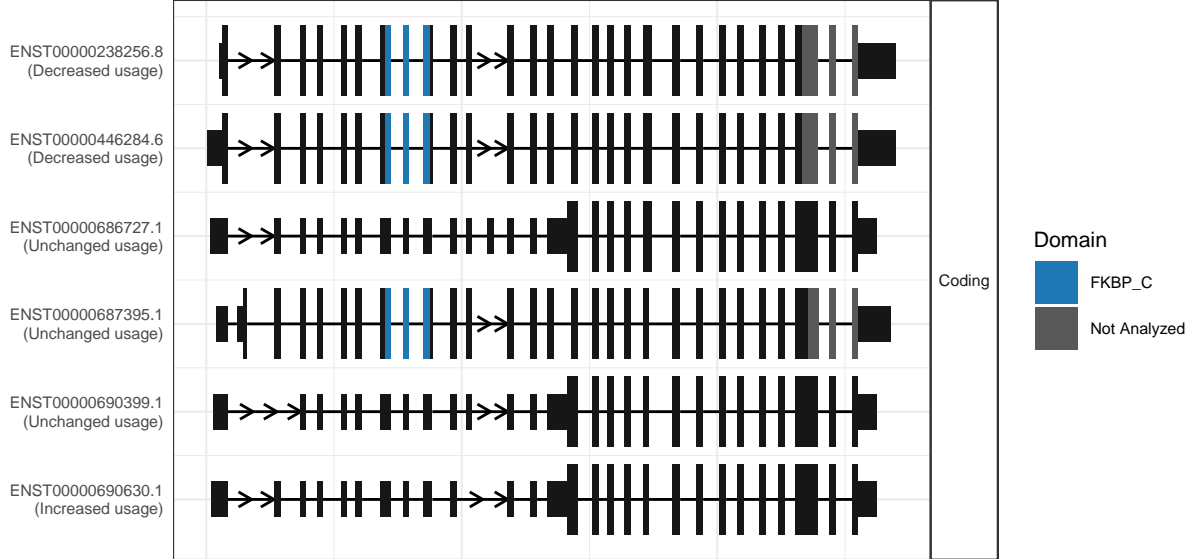


Domain



Pkinase

The isoform switch in FKBP15 (FT vs RGC)



Domain

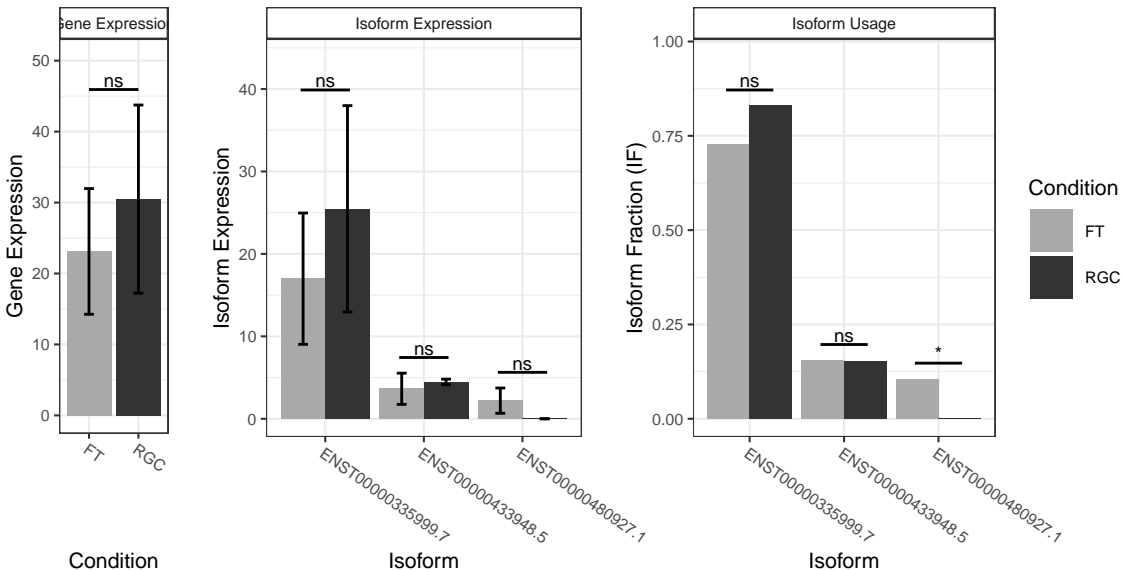
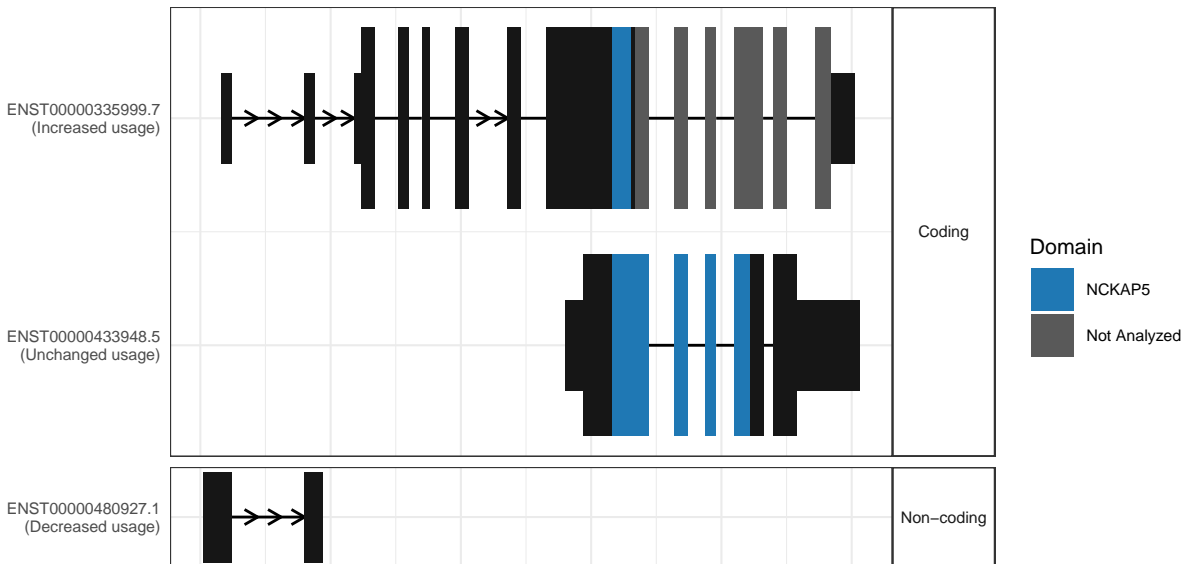


FKBP_C



Not Analyzed

The isoform switch in NCKAP5L (FT vs RGC)



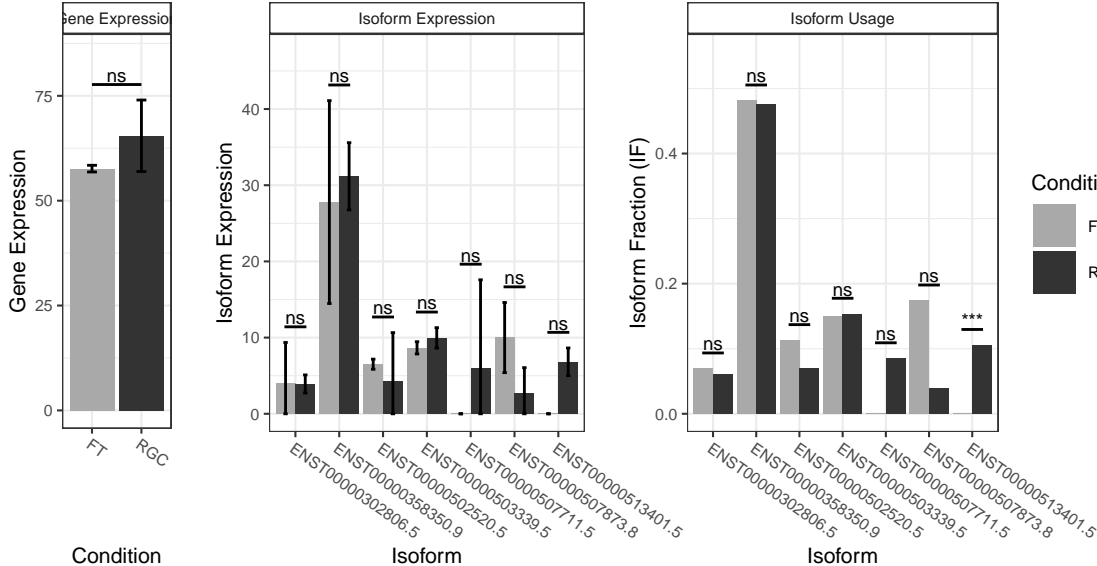
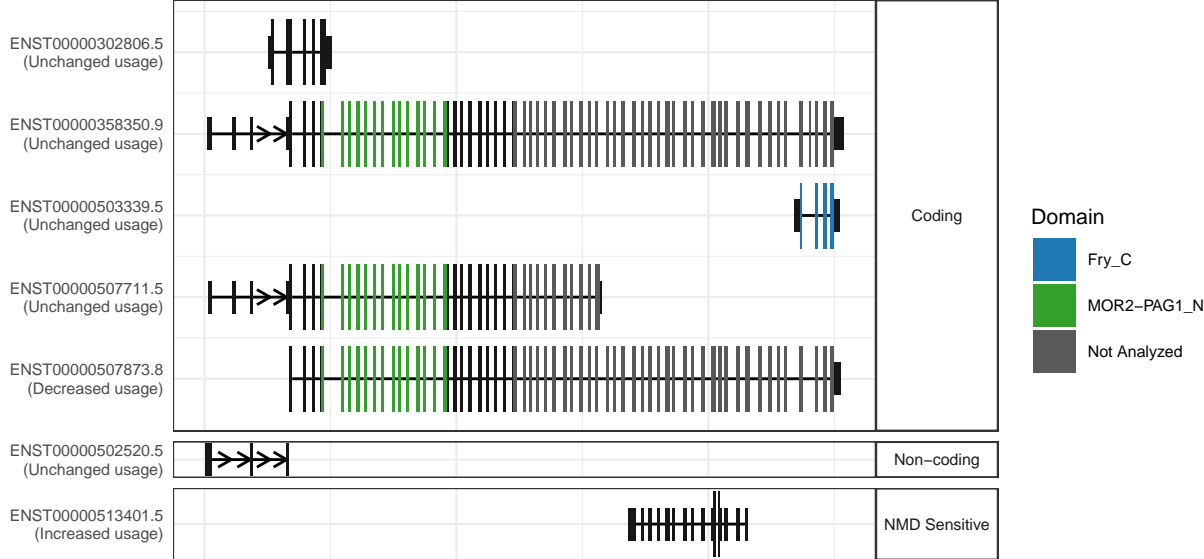
Domain



NCKAP5

Not Analyzed

The isoform switch in FRYL (FT vs RGC)



Domain



Fry_C

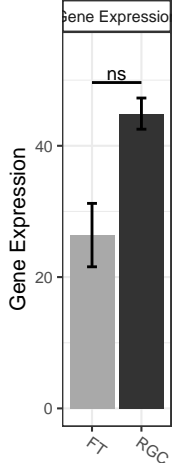
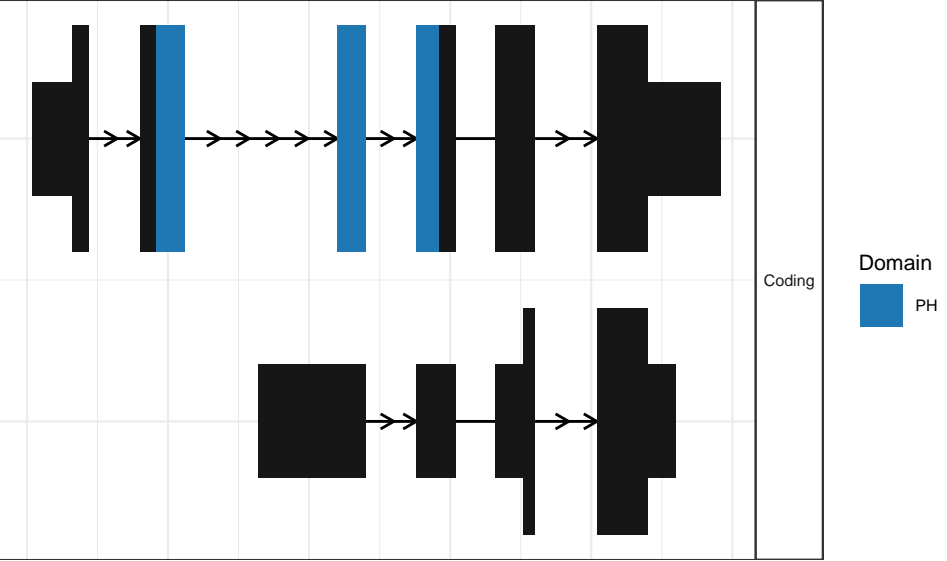


MOR2-PAG1_N

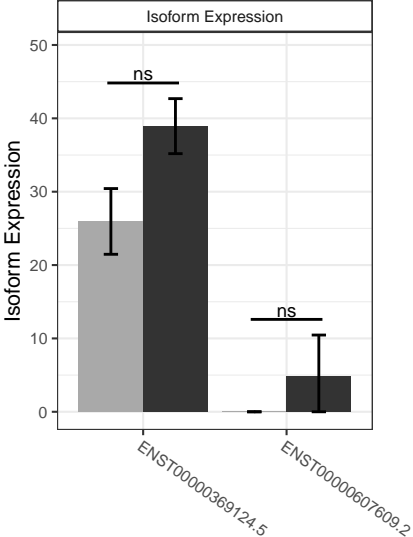


Not Analyzed

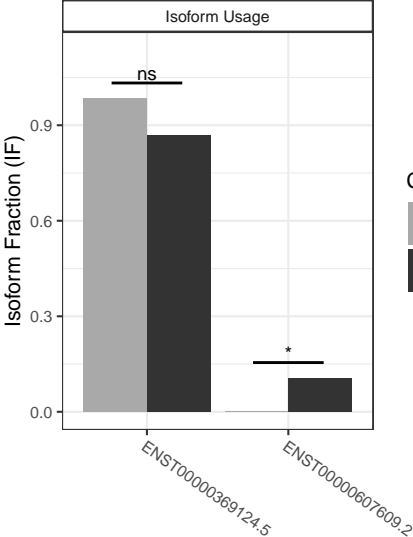
The isoform switch in PLEKHO1 (FT vs RGC)



Condition



Isoform



Isoform

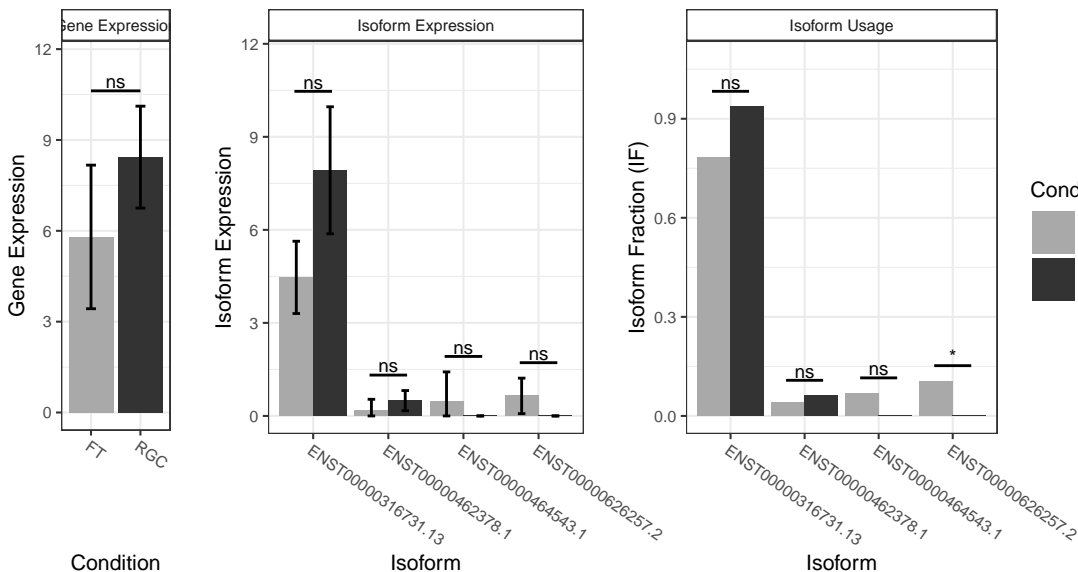
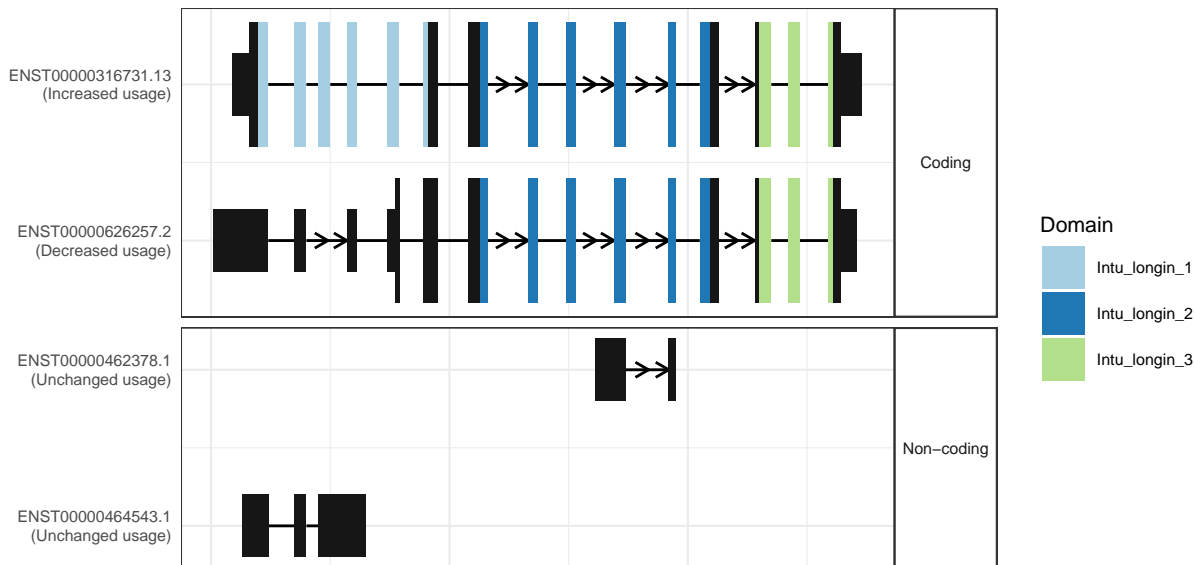


Domain



PH

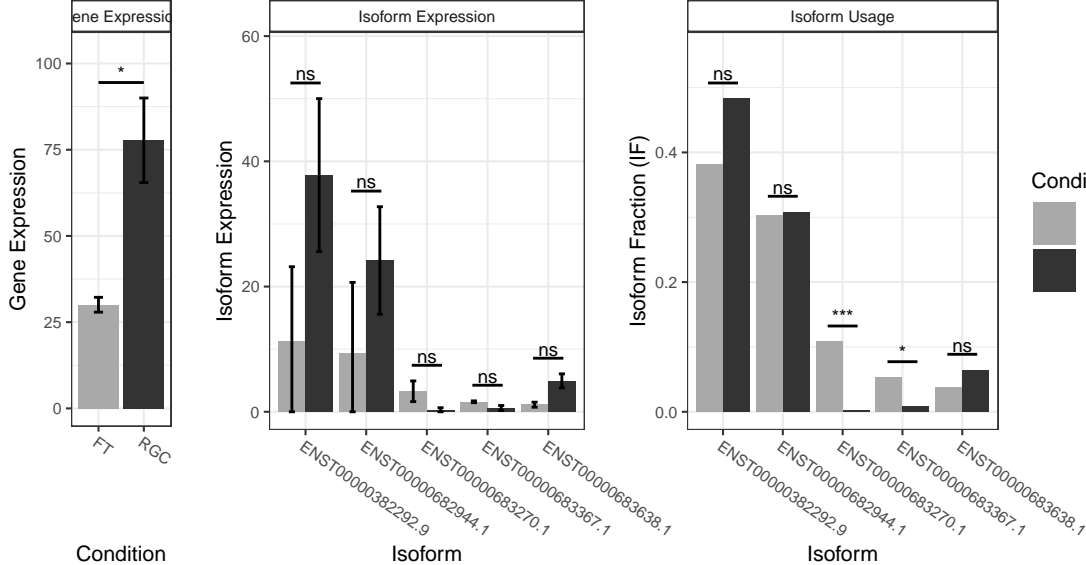
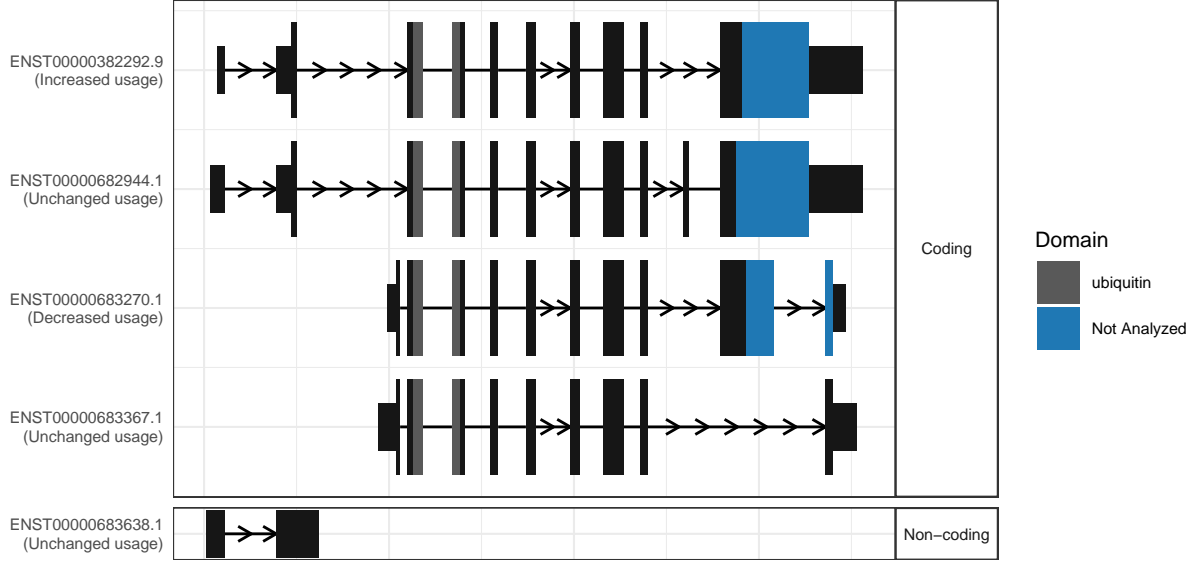
The isoform switch in CCZ1B (FT vs RGC)



Domain

	Intu_longin_1
	Intu_longin_2
	Intu_longin_3

The isoform switch in SACS (FT vs RGC)



Domain

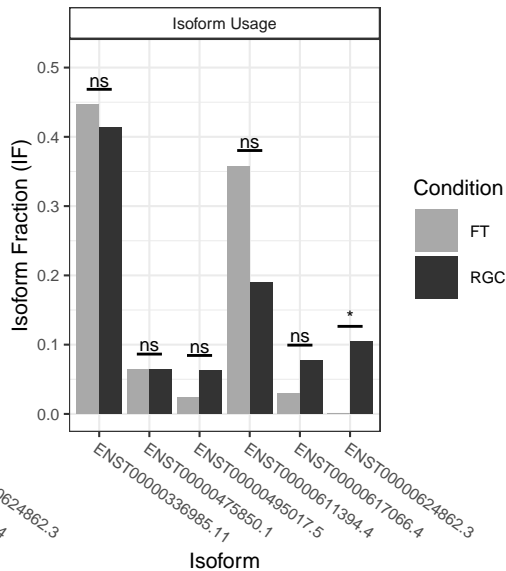
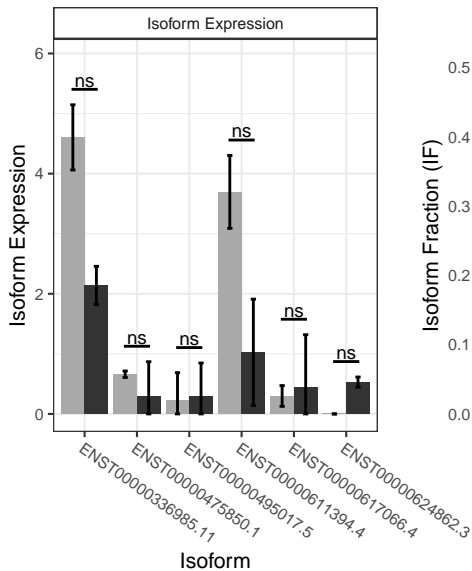
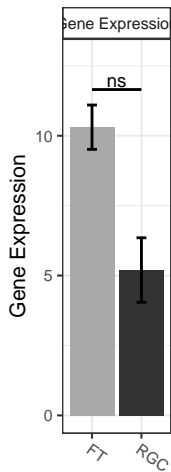
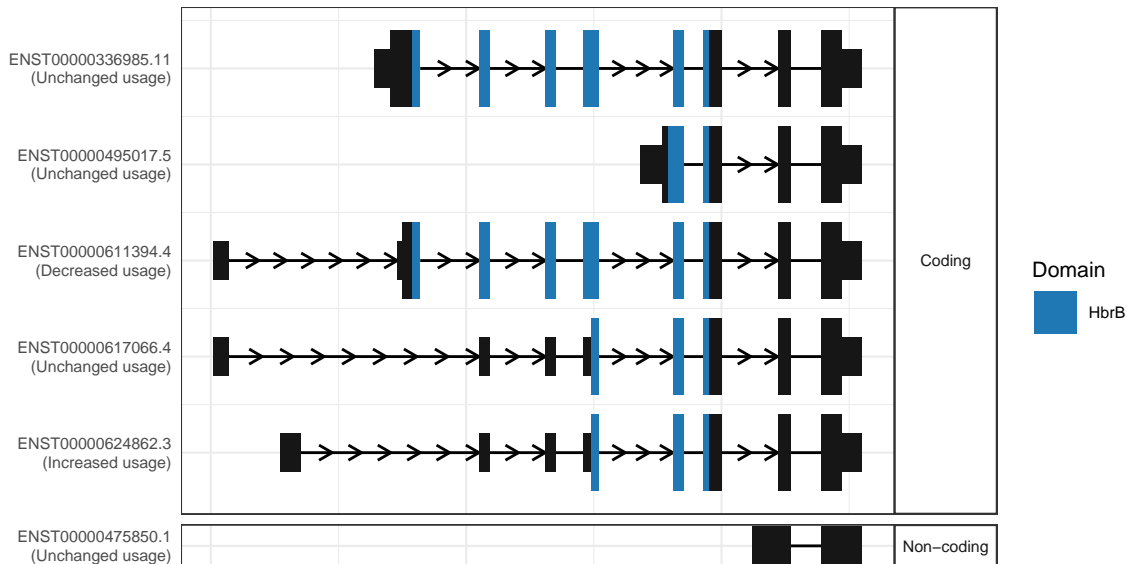


ubiquitin



Not Analyzed

The isoform switch in PRR5 (FT vs RGC)

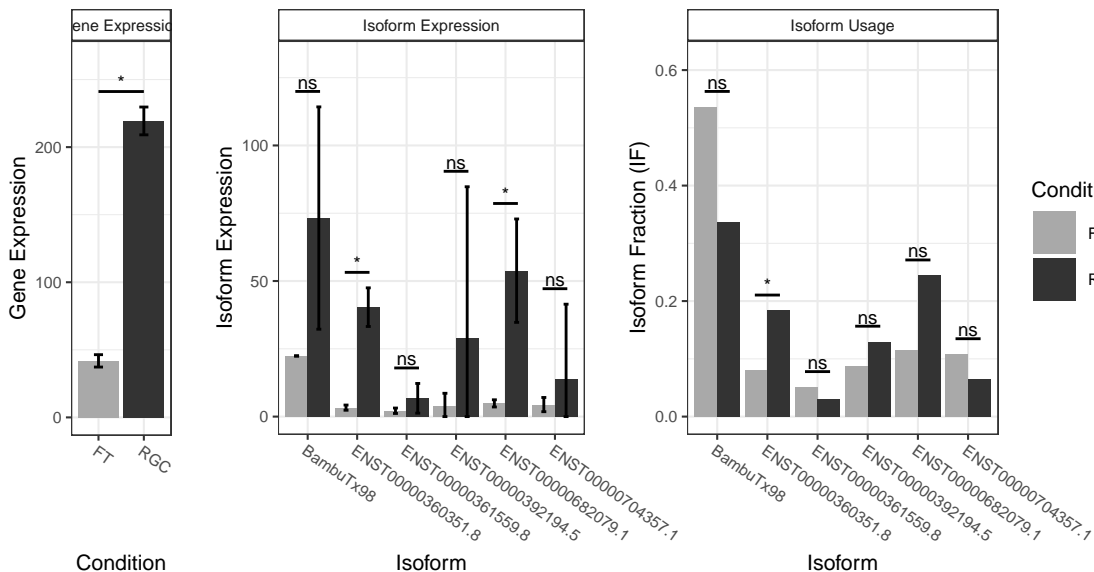
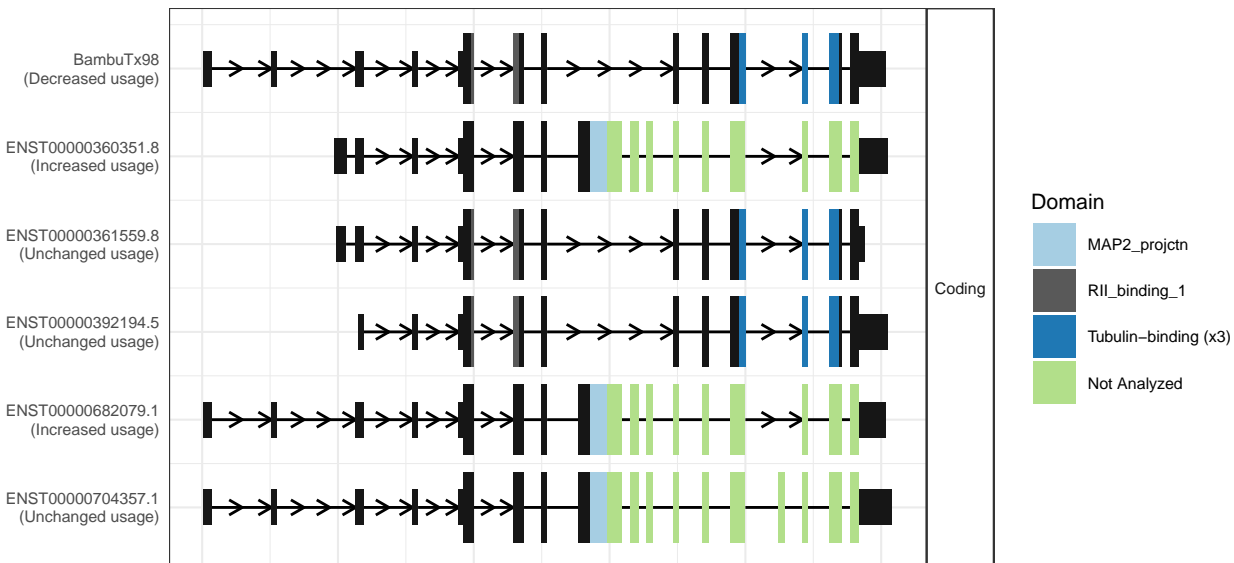


Domain

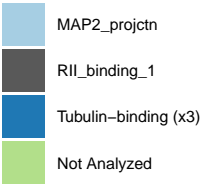


HbrB

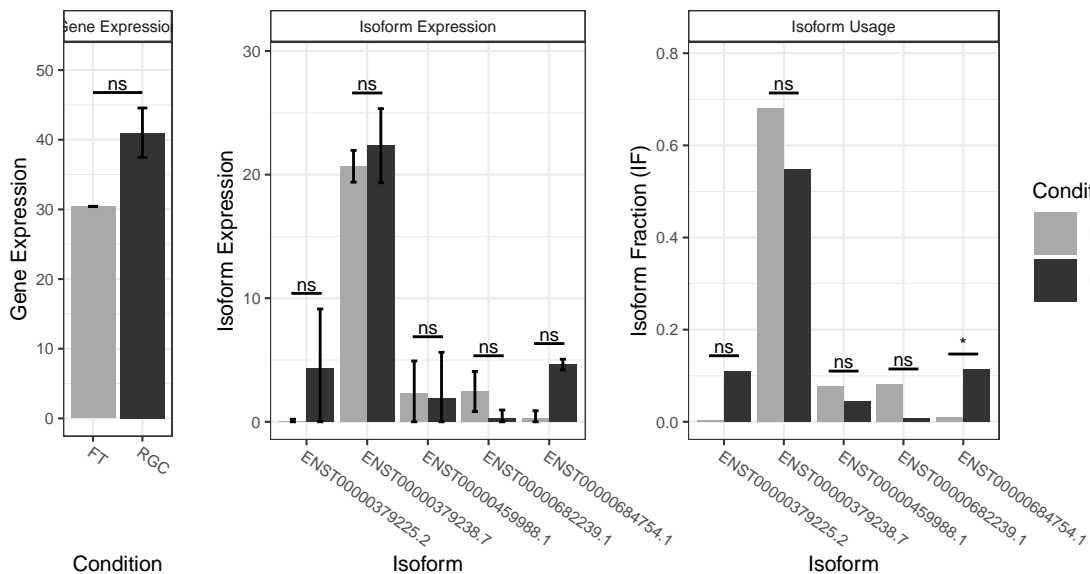
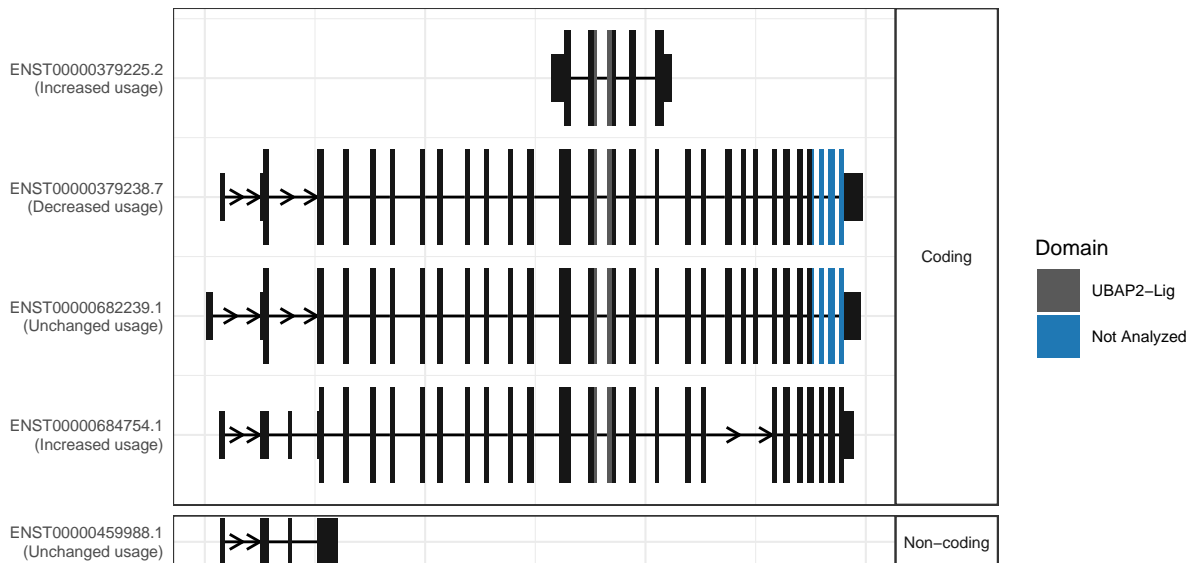
The isoform switch in MAP2 (FT vs RGC)



Domain



The isoform switch in UBAP2 (FT vs RGC)



Domain

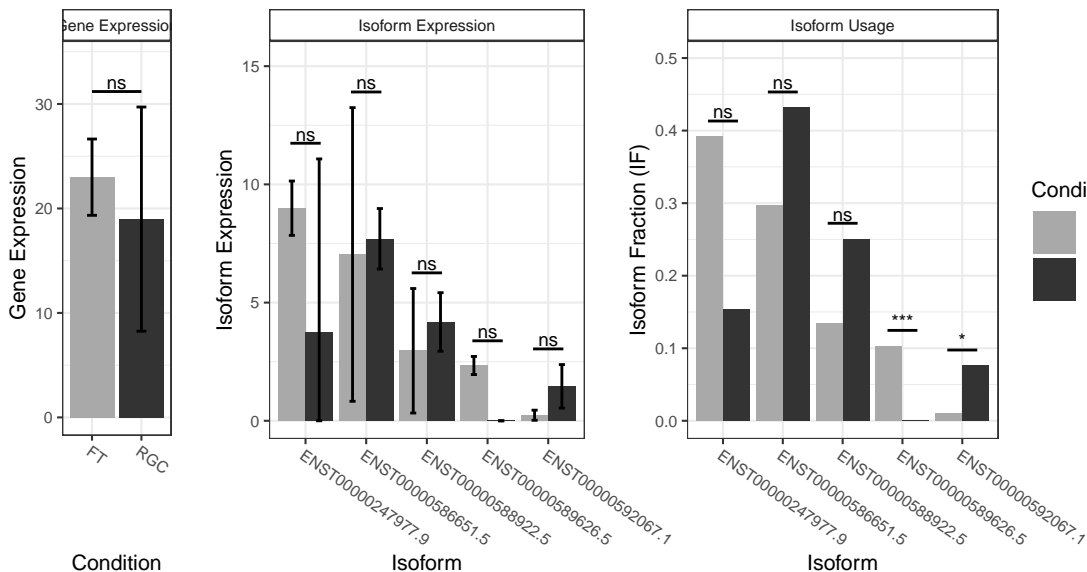
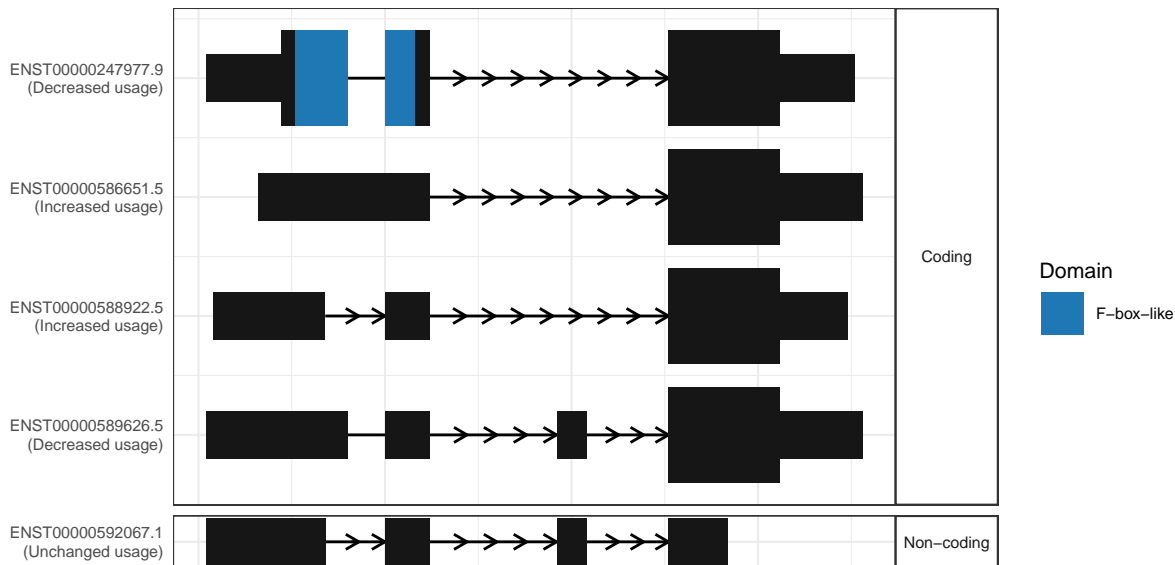


UBAP2-Lig



Not Analyzed

The isoform switch in FBXL12 (FT vs RGC)

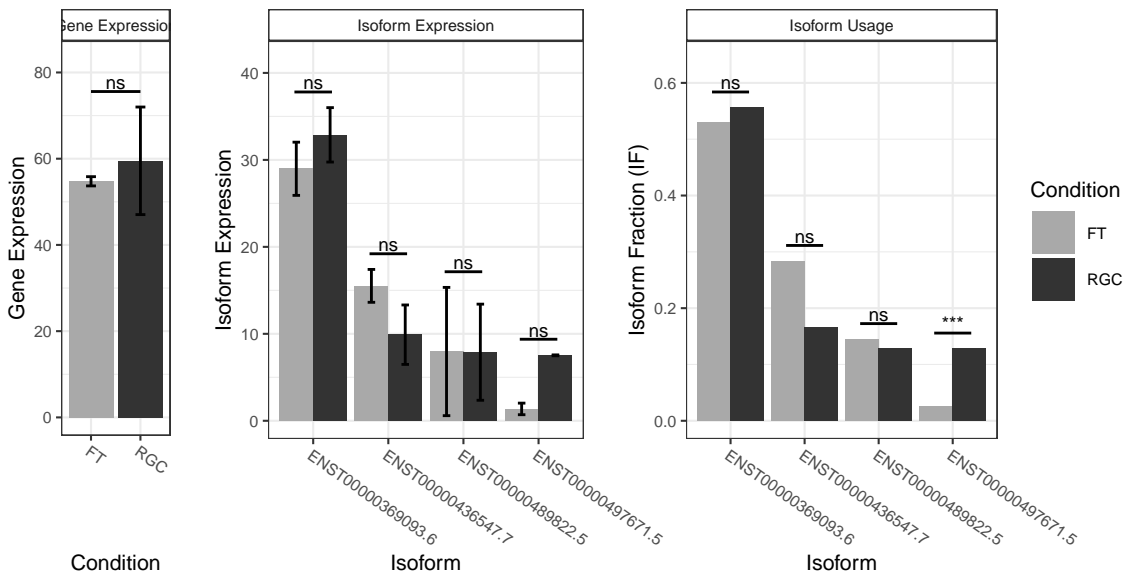
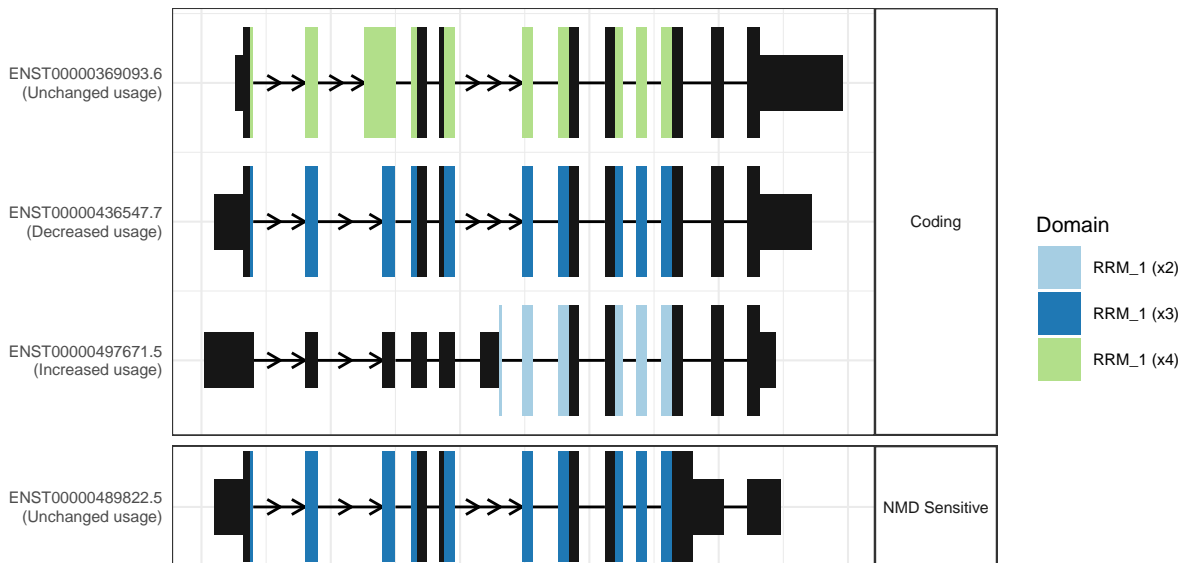


Domain

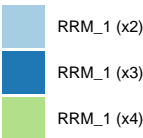


F-box-like

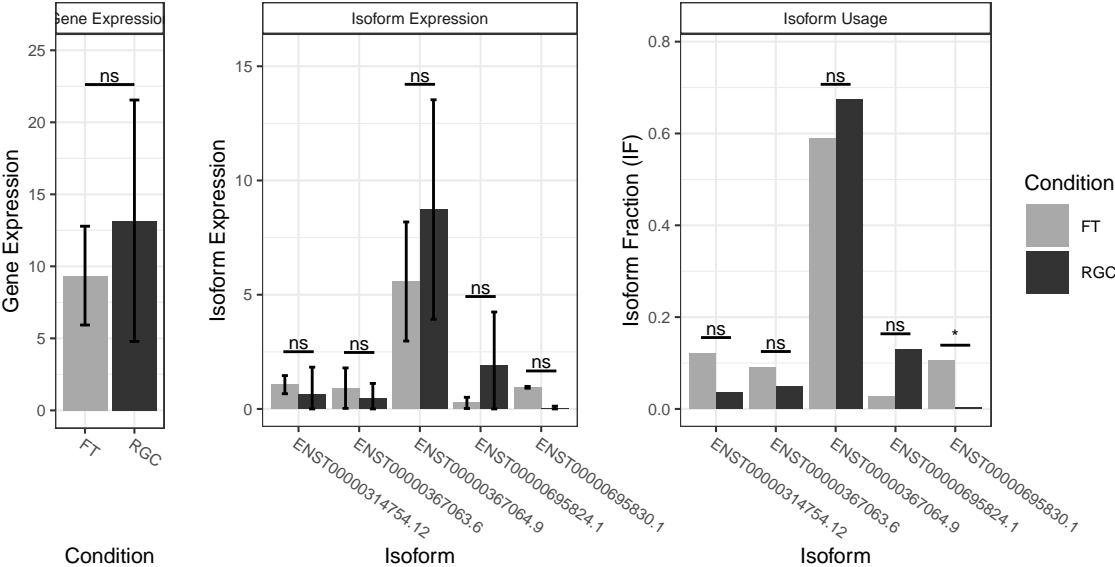
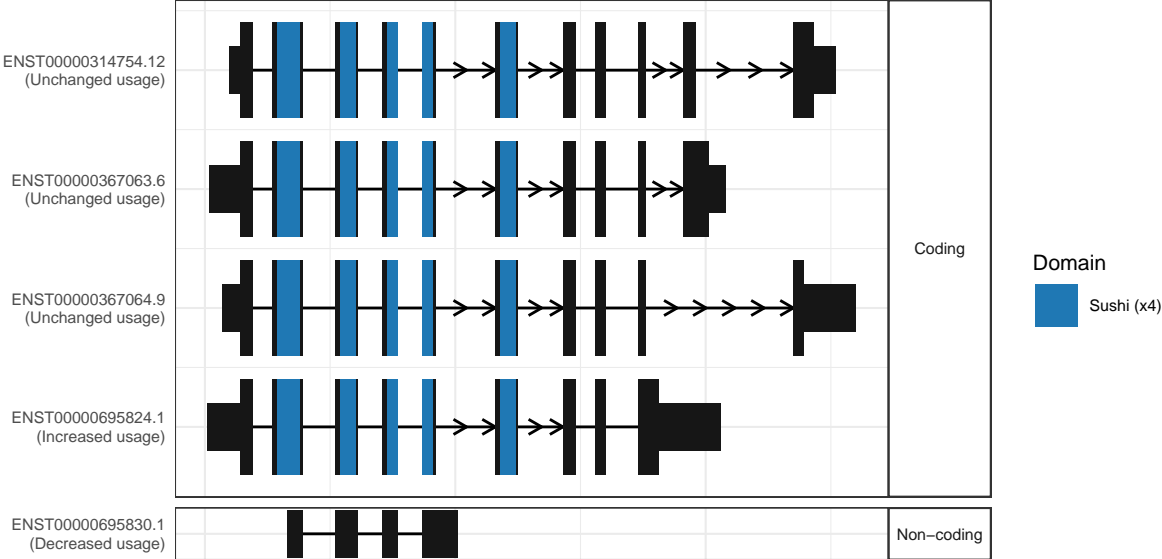
The isoform switch in TIAL1 (FT vs RGC)



Domain



The isoform switch in CD55 (FT vs RGC)

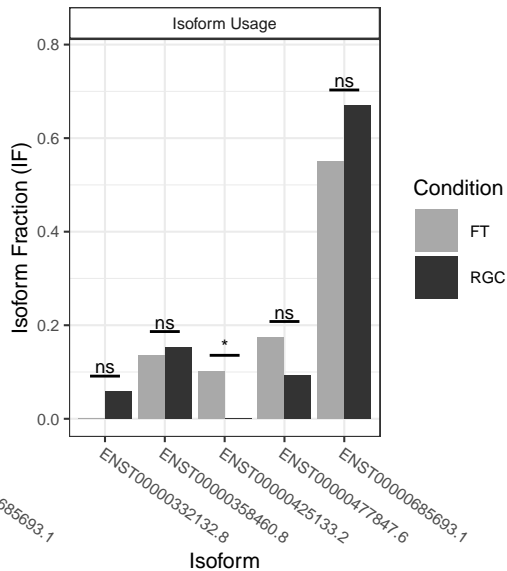
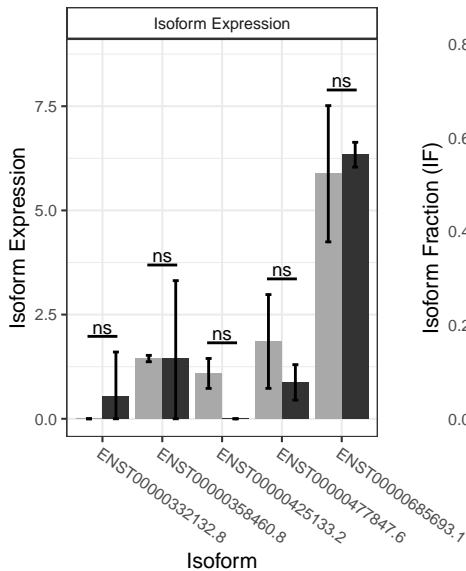
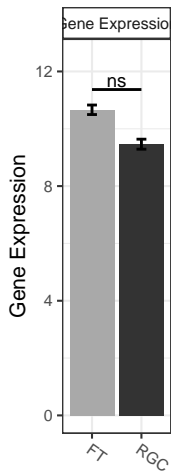
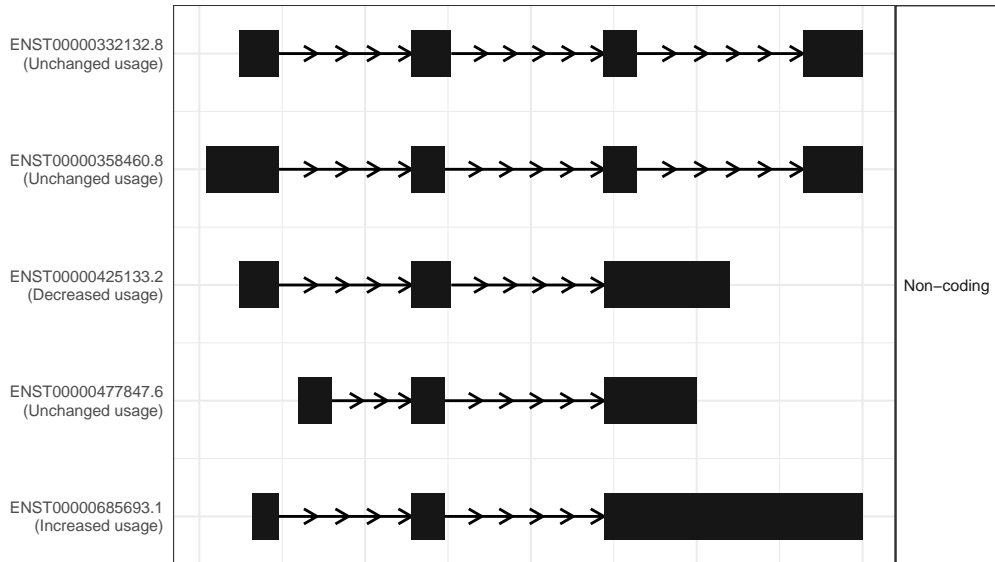


Domain

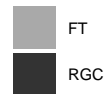


Sushi (x4)

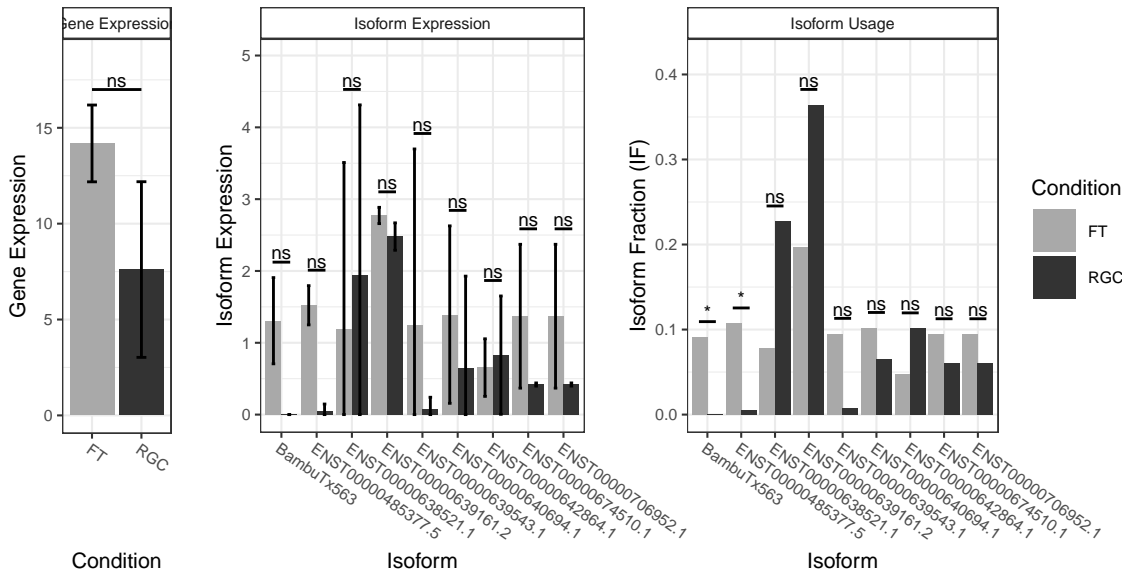
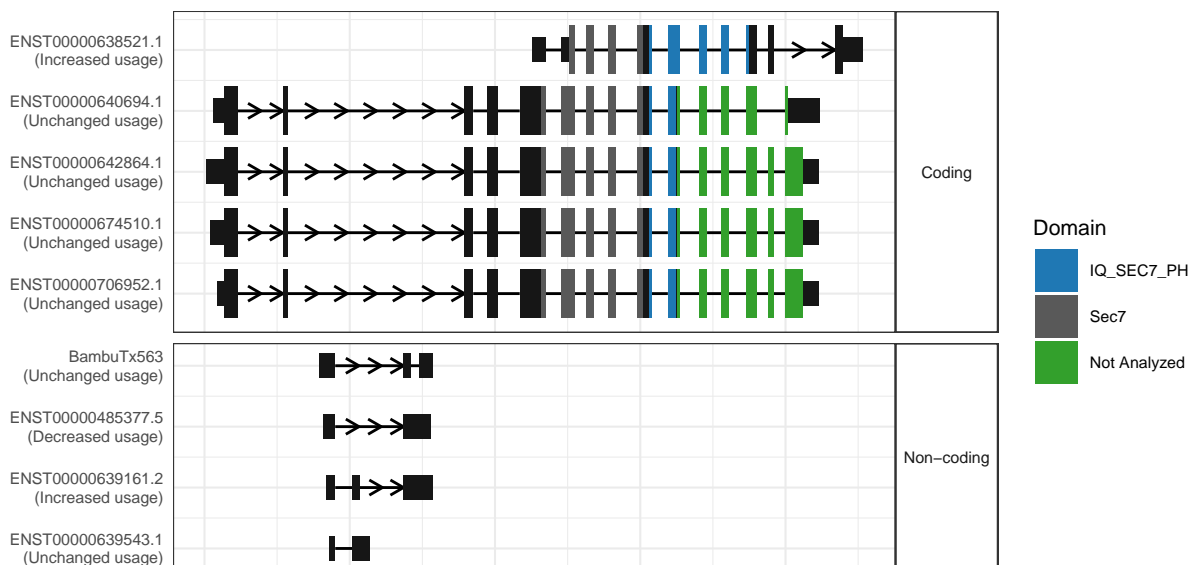
The isoform switch in VCF2 (FT vs RGC)






Condition



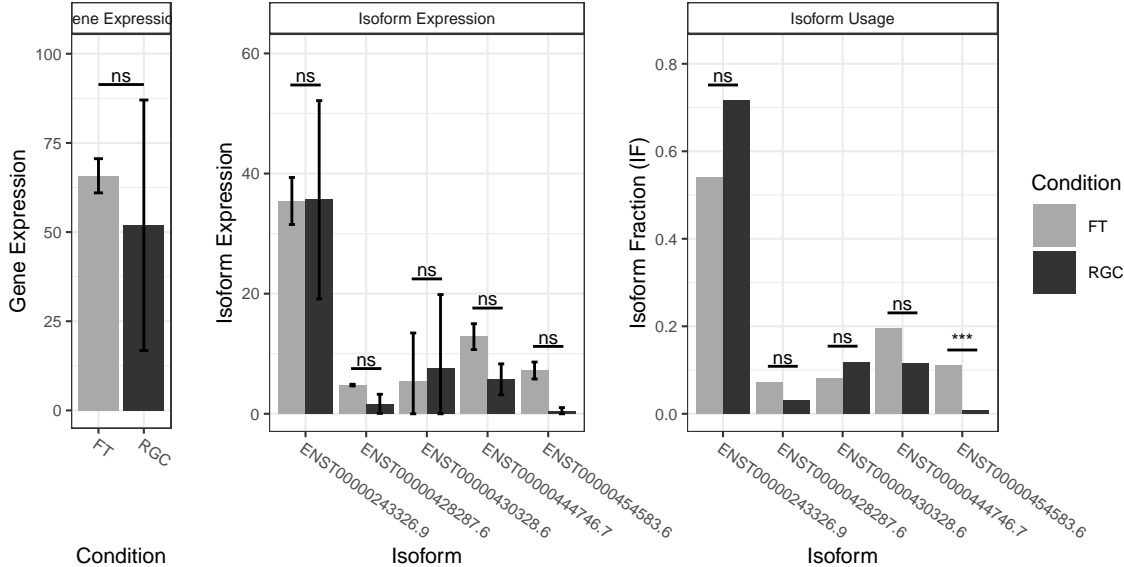
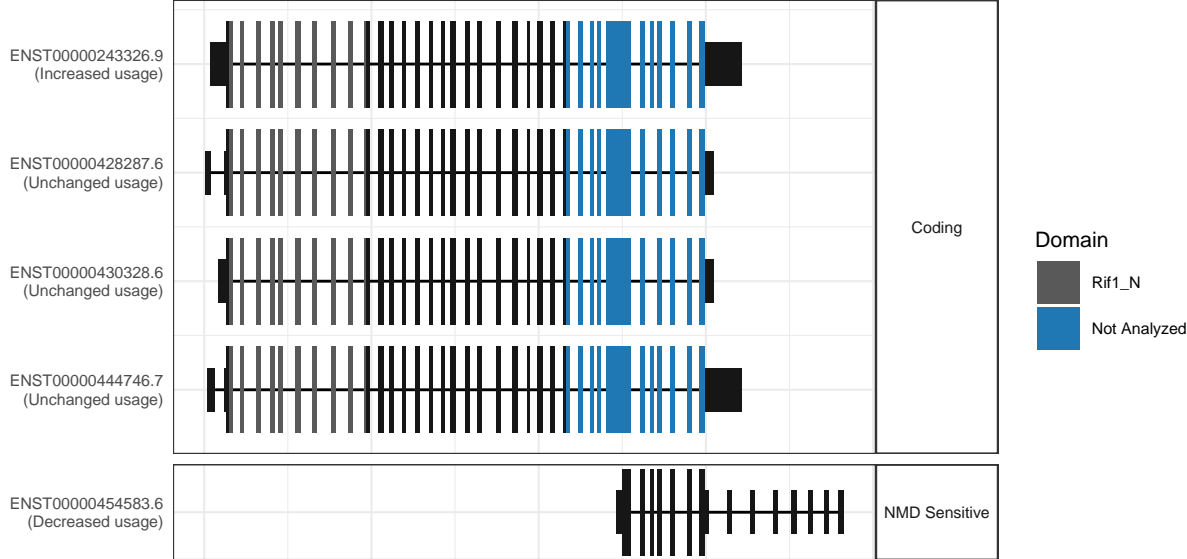
The isoform switch in IQSEC2 (FT vs RGC)



Domain

	IQ_SEC7_PH
	Sec7
	Not Analyzed

The isoform switch in RIF1 (FT vs RGC)



Domain

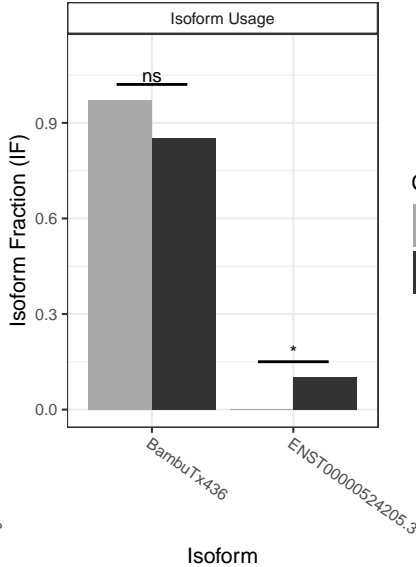
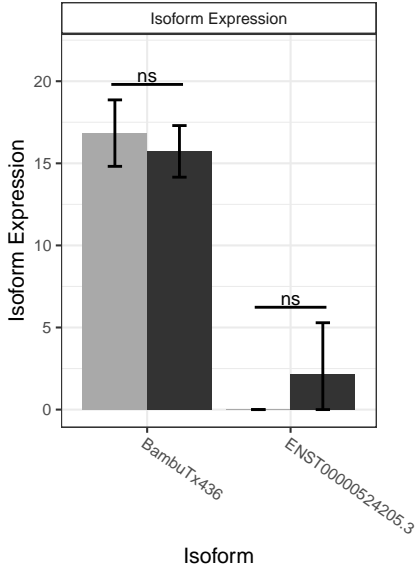
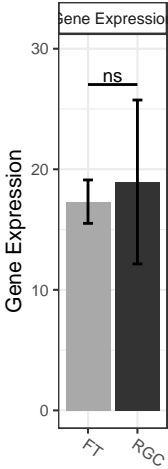
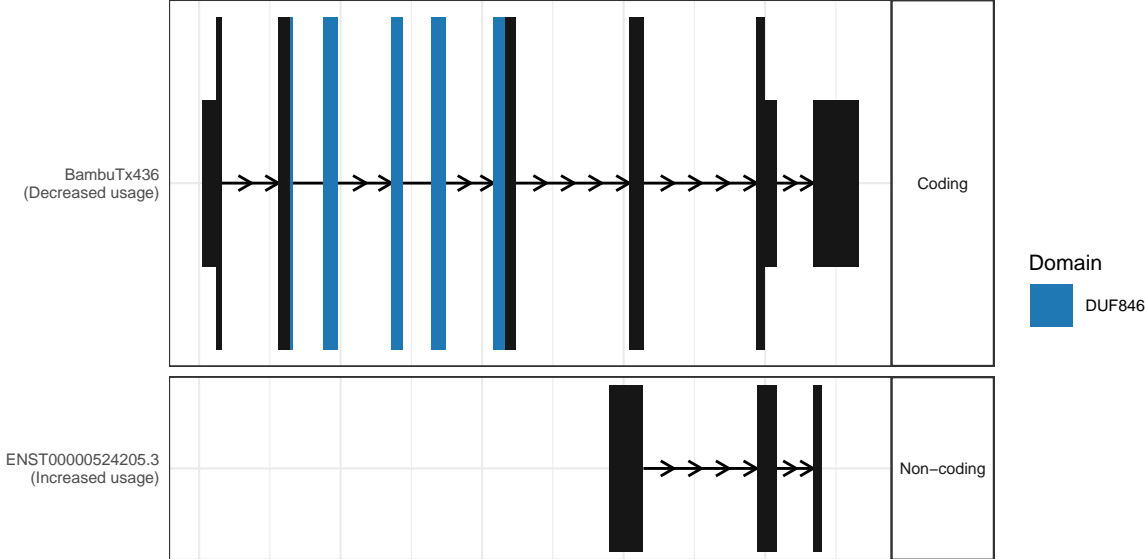


Rif1_N



Not Analyzed

The isoform switch in TVP23C-CDRT4 (FT vs RGC)

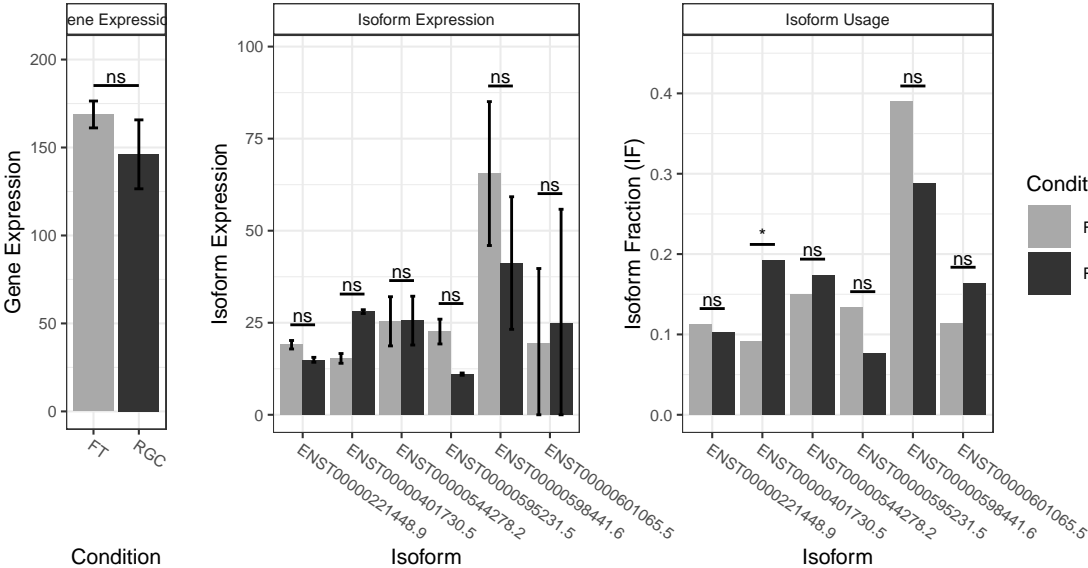
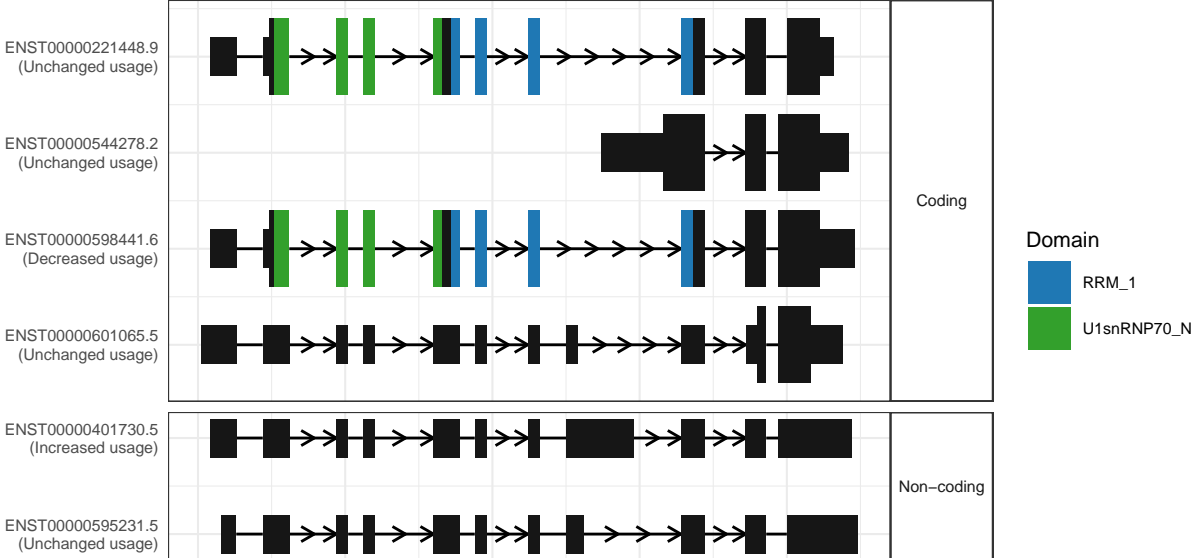


Domain



DUF646

The isoform switch in SNRNP70 (FT vs RGC)



Domain

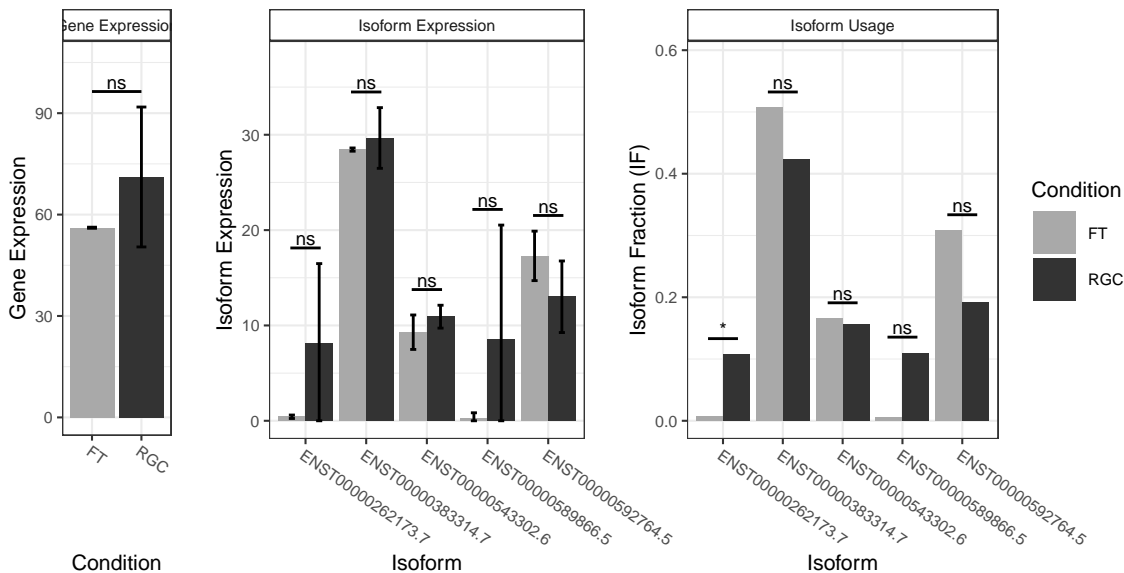
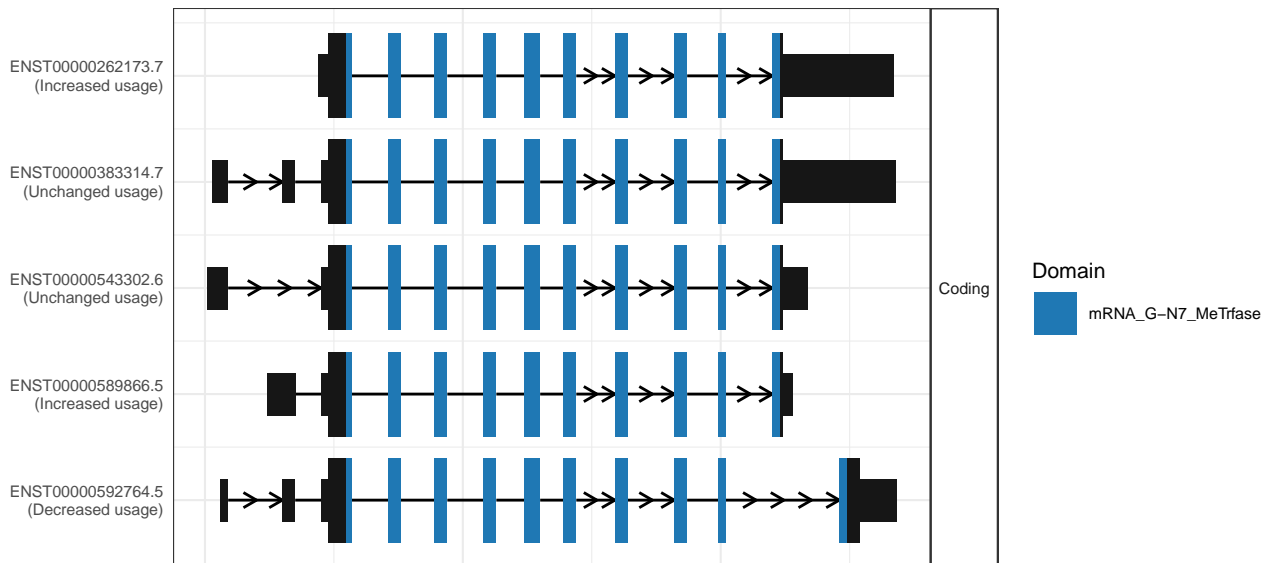


RRM_1



U1snRNP70_N

The isoform switch in RNMT (FT vs RGC)



Domain



mRNA_G-N7_MeTrfase