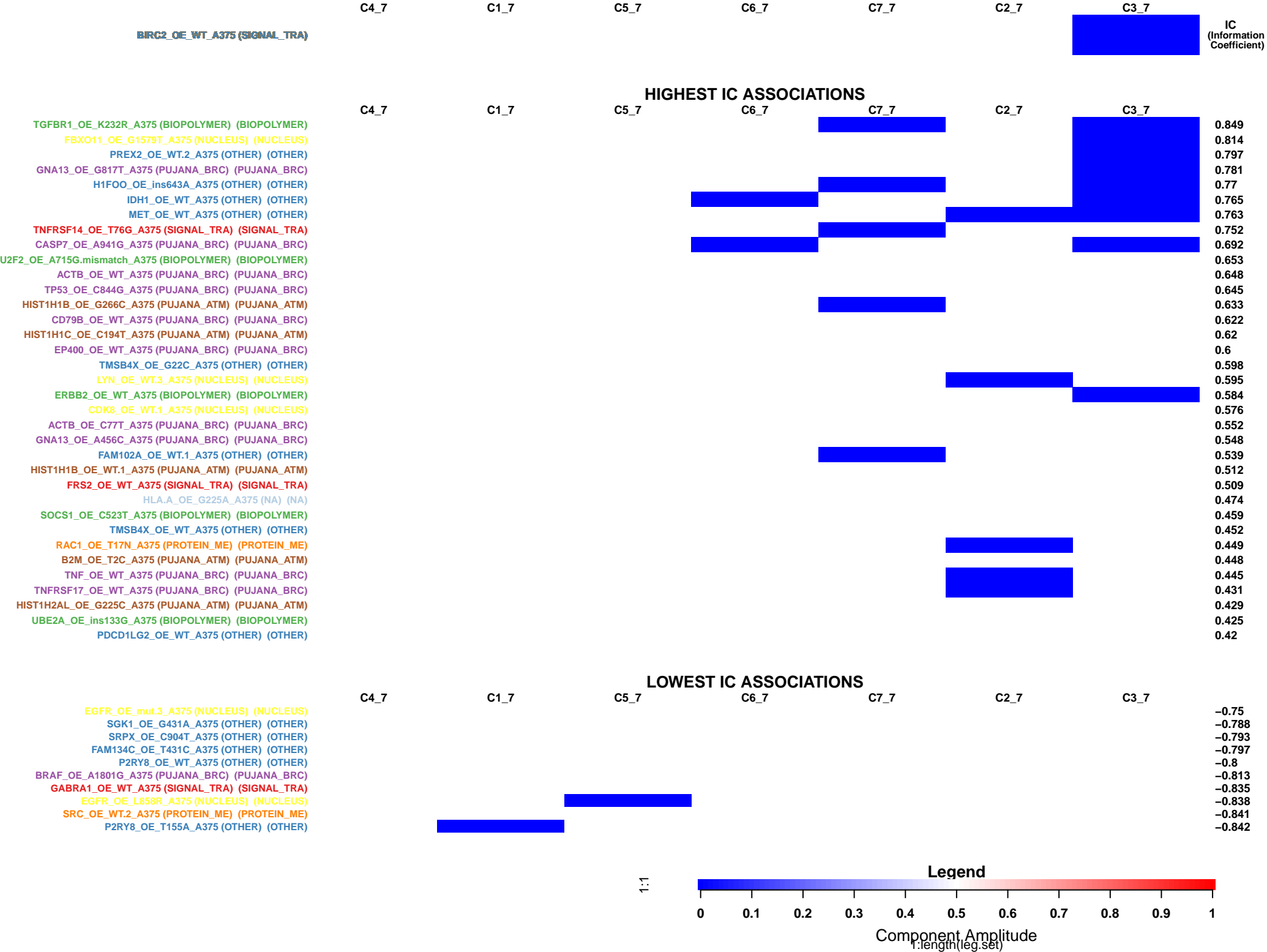
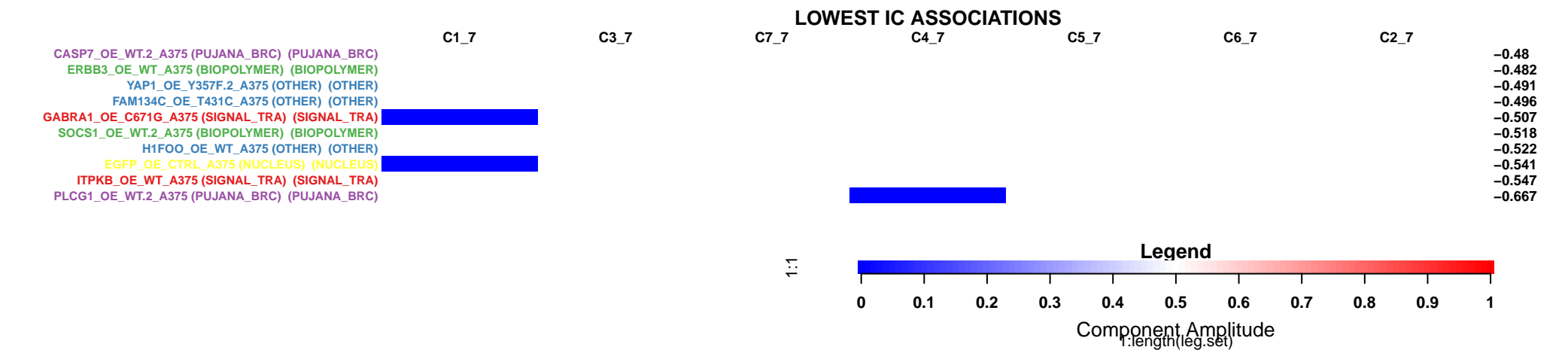
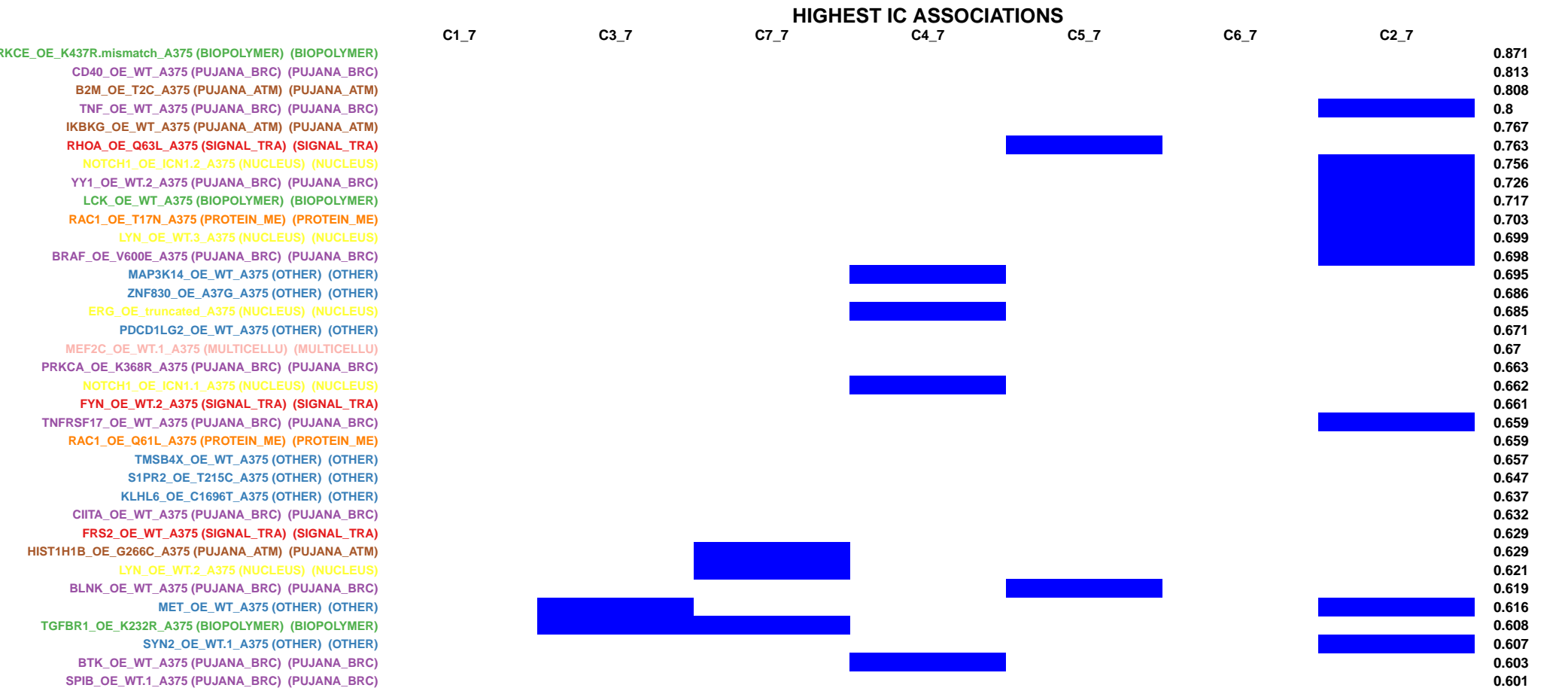


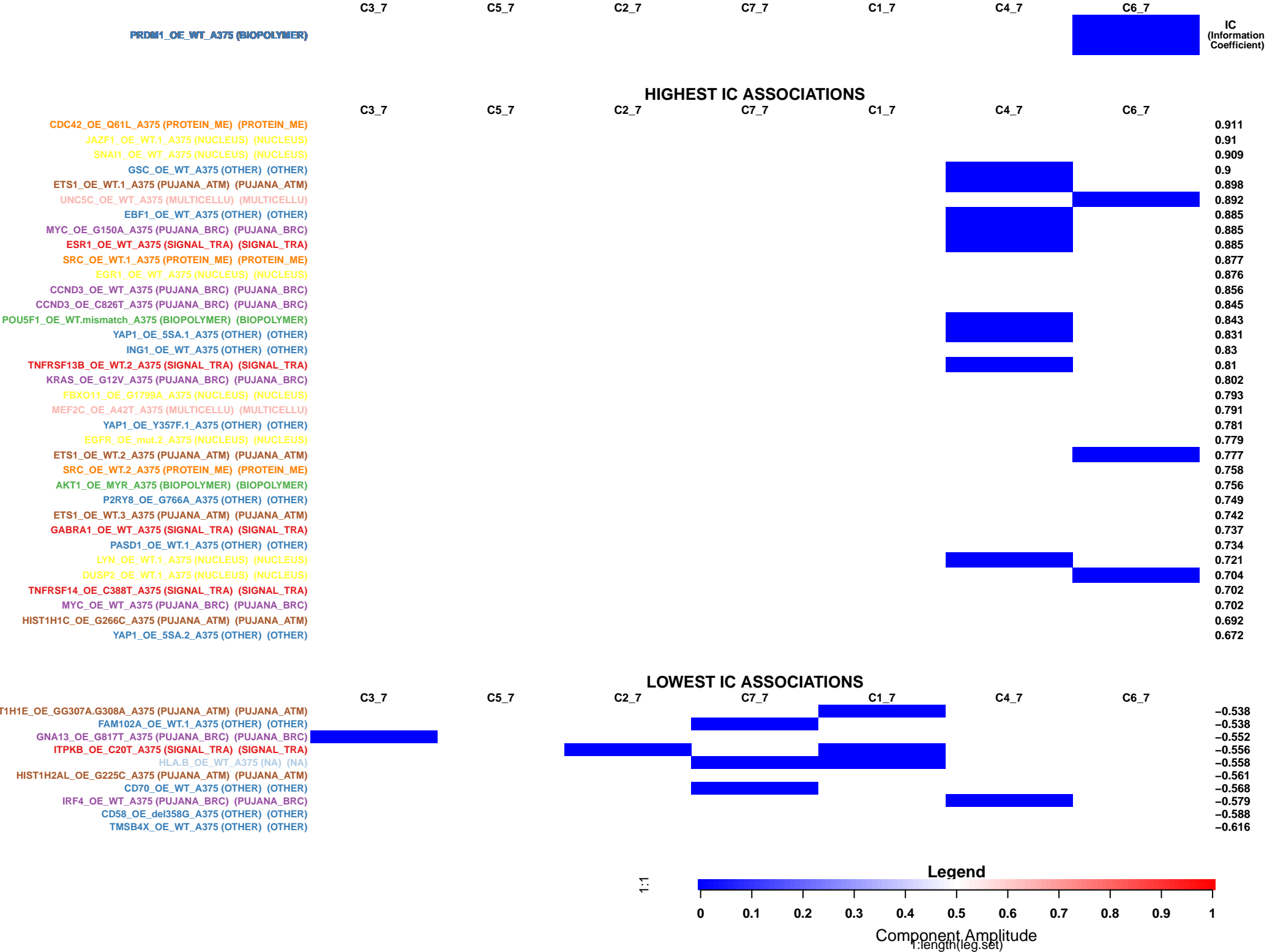
BIRC2\_OE\_WT\_A375 (SIGNAL\_TRA)



C1_7	C3_7	C7_7	C4_7	C5_7	C6_7	C2_7	IC (Information Coefficient)
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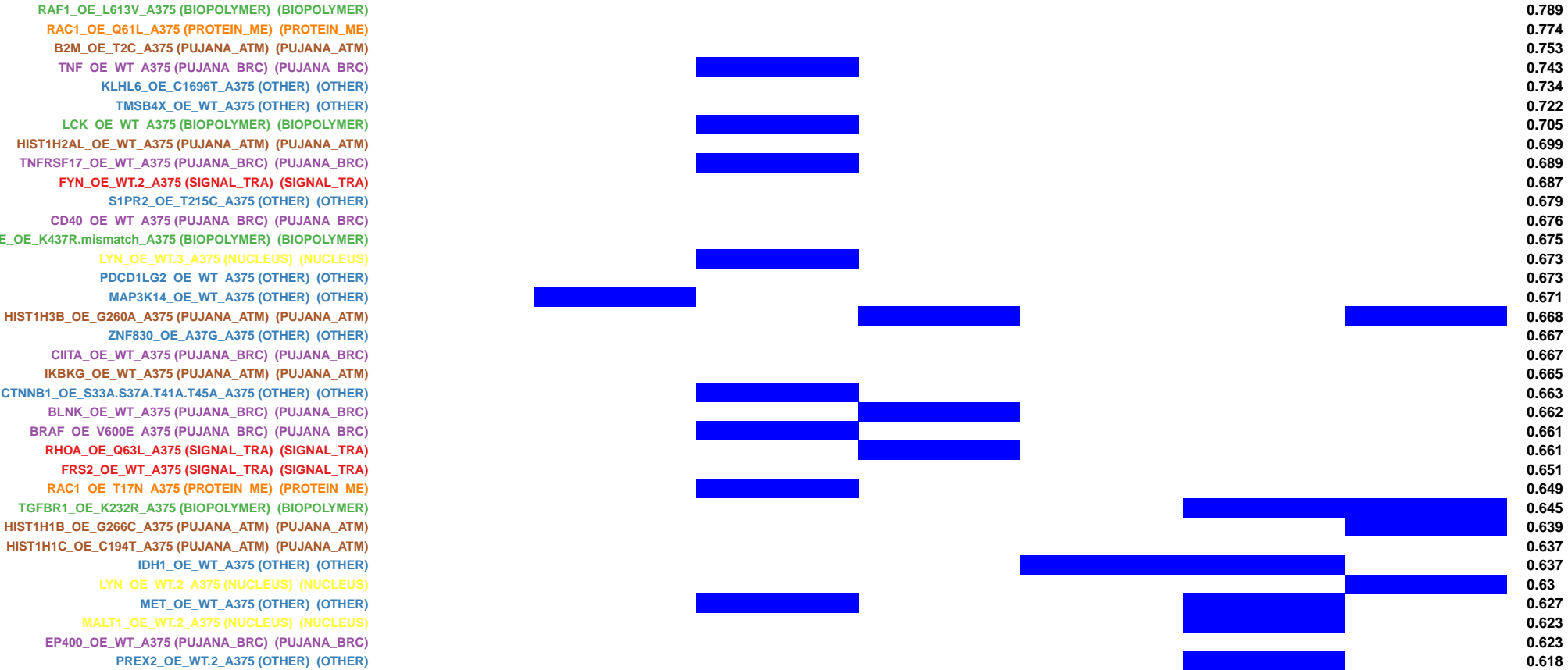
PRDM1\_OE\_WT\_A375 (BIOPOLYMER)



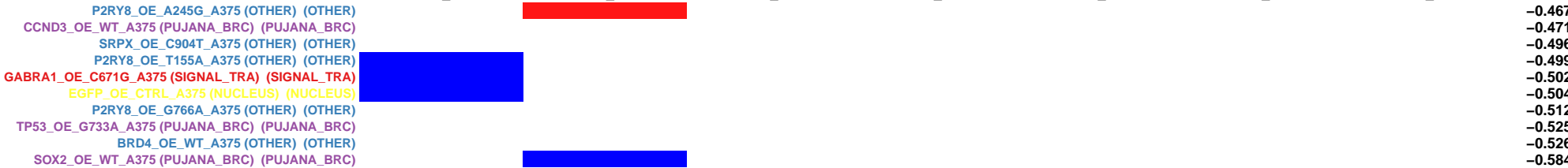
C1_7	C4_7	C2_7	C5_7	C6_7	C3_7	C7_7
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PRKCA\_OE\_K368R\_A375 (PUJANA\_BRC)

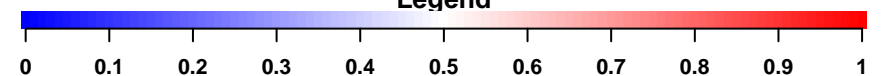
C1_7	C4_7	C2_7	C5_7	C6_7	C3_7	C7_7
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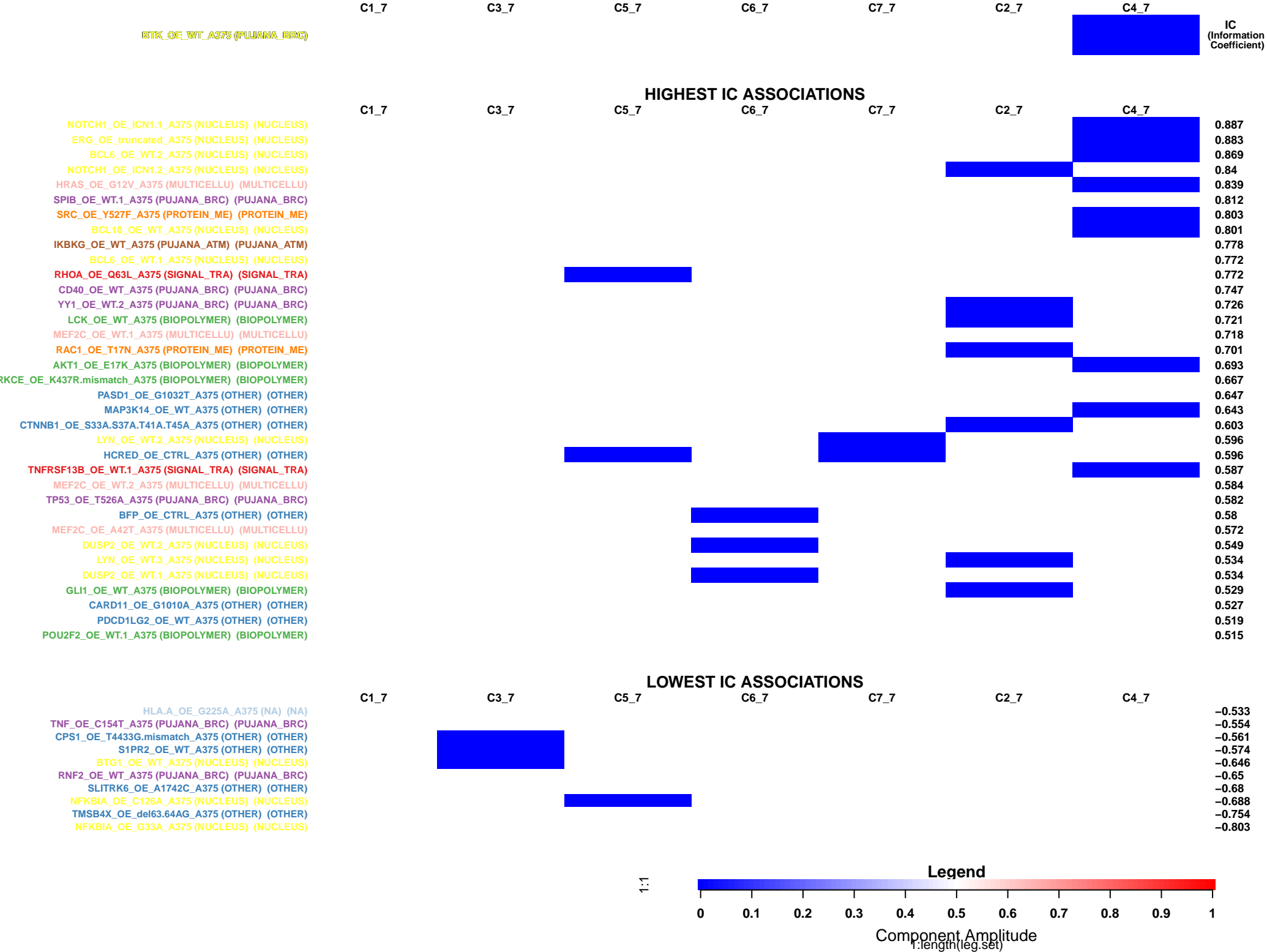
C1_7	C4_7	C2_7	C5_7	C6_7	C3_7	C7_7
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### Legend

Component Amplitude  
1:length(leg.set)

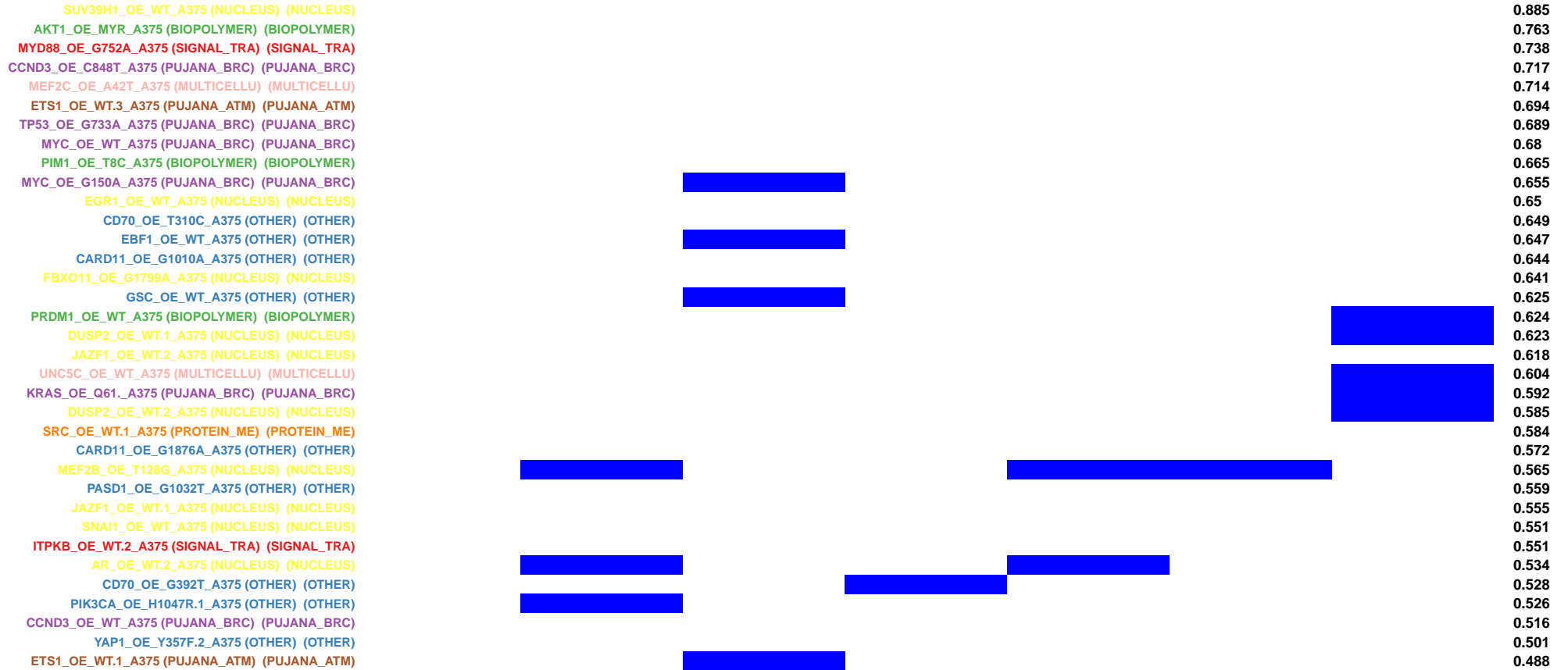
BTK\_OE\_WT\_A375 (PUJANA\_BRC)



C3\_7                      C5\_7                      C4\_7                      C2\_7                      C1\_7                      C7\_7                      C6\_7

**CARD11\_OE\_F123Lmismatch\_A375 (OTHER)**

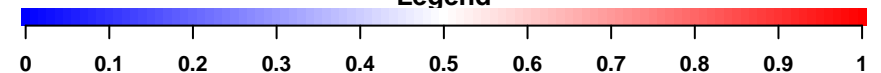
C3\_7                      C5\_7                      C4\_7                      C2\_7                      C1\_7                      C7\_7                      C6\_7



C3\_7                      C5\_7                      C4\_7                      C2\_7                      C1\_7                      C7\_7                      C6\_7

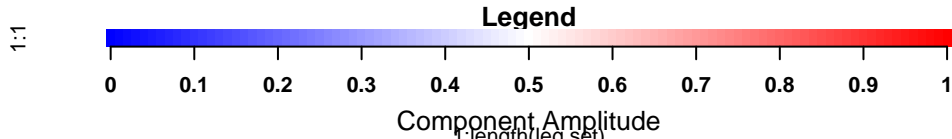
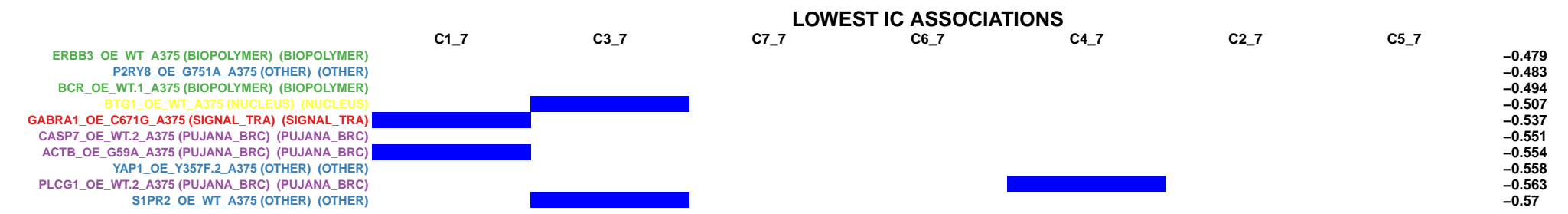
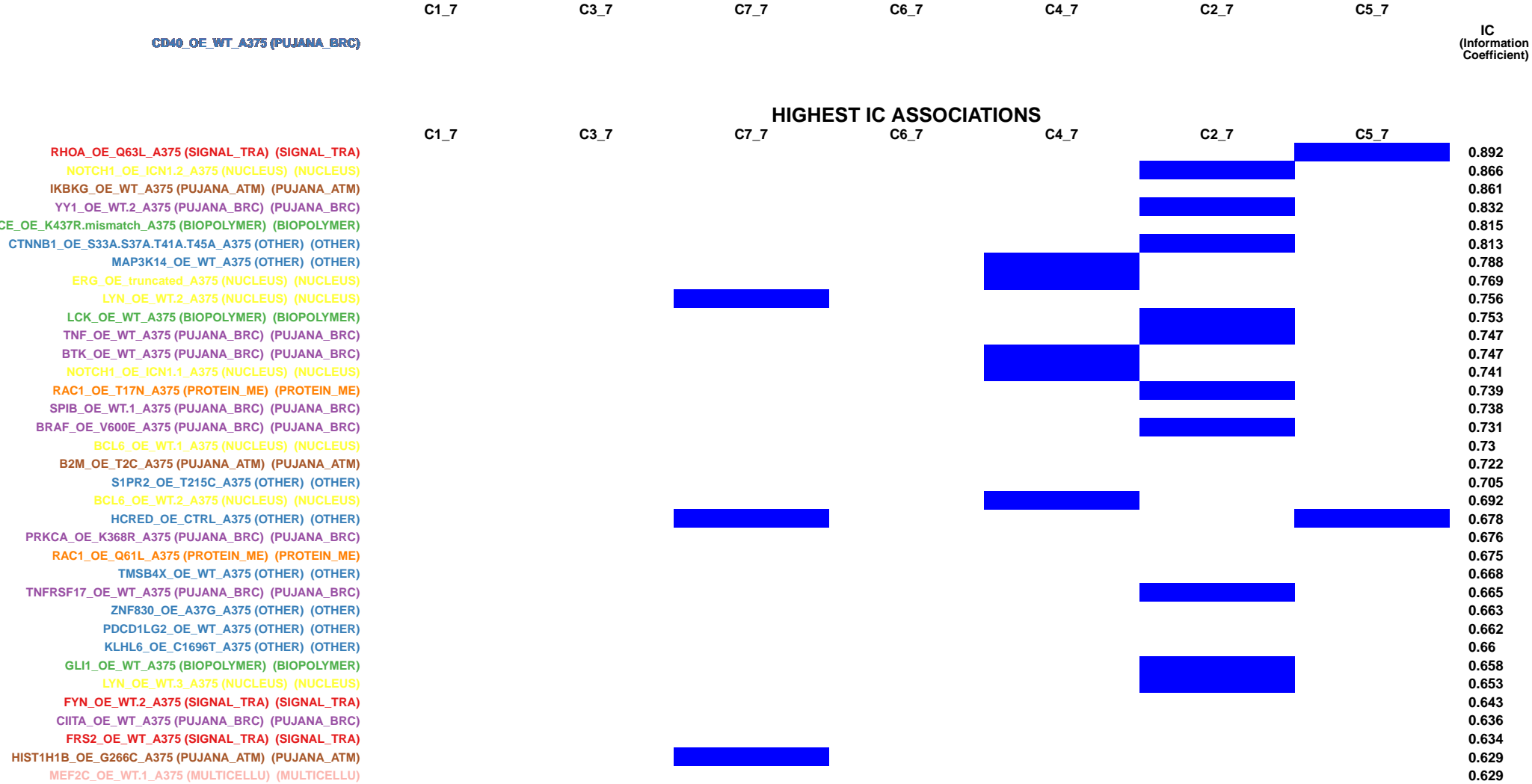


### Legend



```
1:length(leg.set)
```

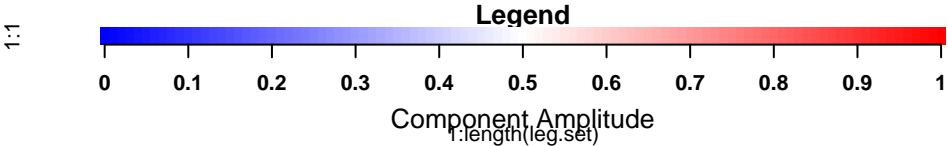
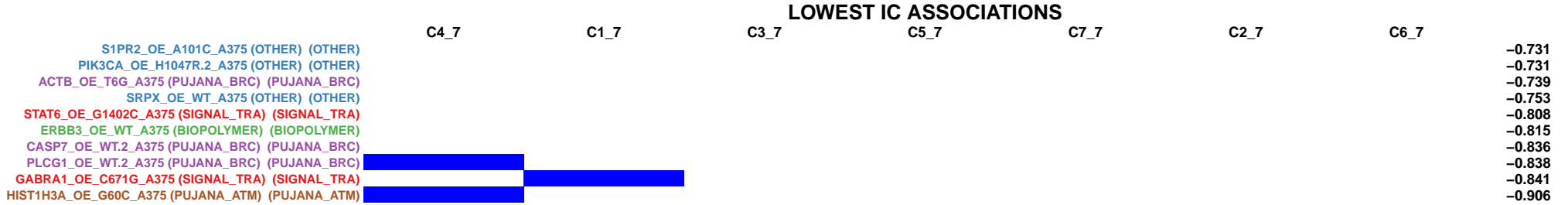
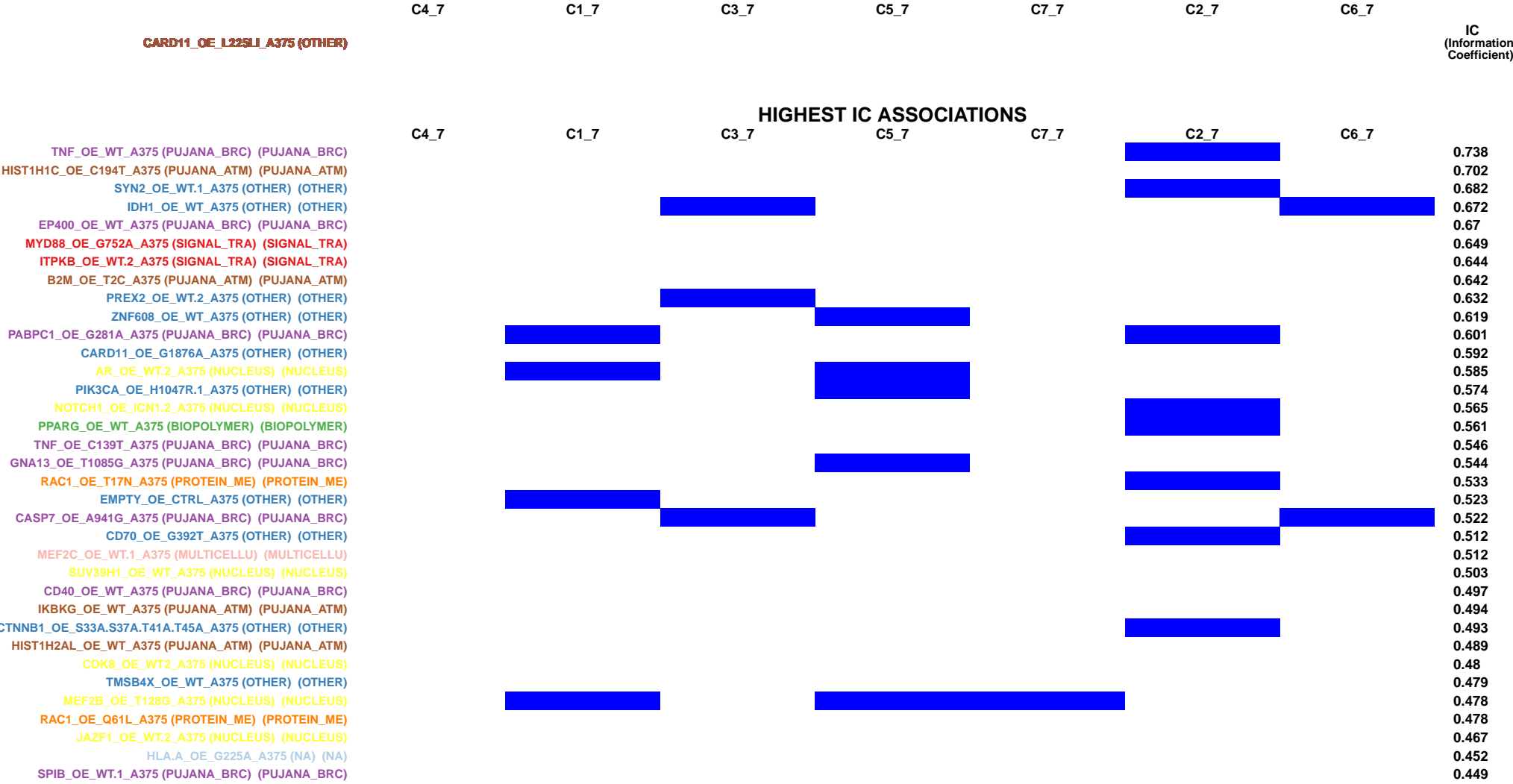
CD40\_OE\_WT\_A375 (PUJANA\_BRC)



CARD11\_OE\_L225LI\_A375 (OTHER)

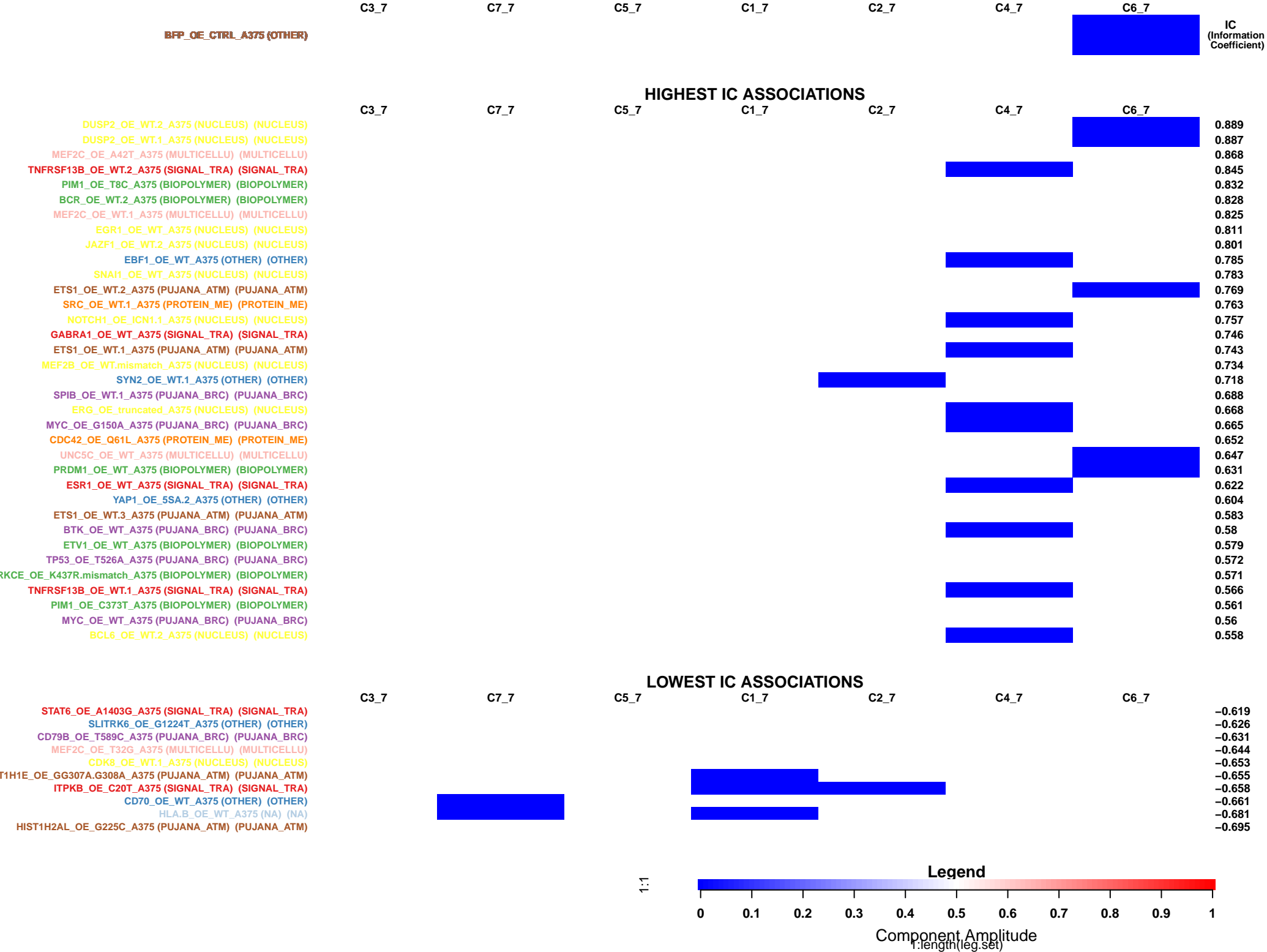
CARD11\_OE\_L225LI\_A375 (OTHER)

IC  
(Information  
Coefficient)

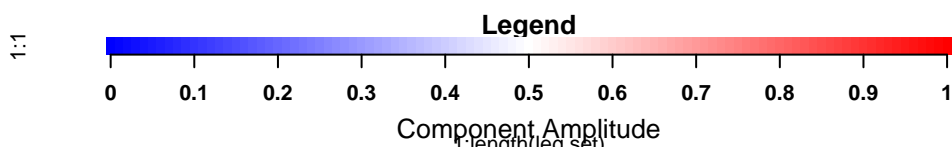
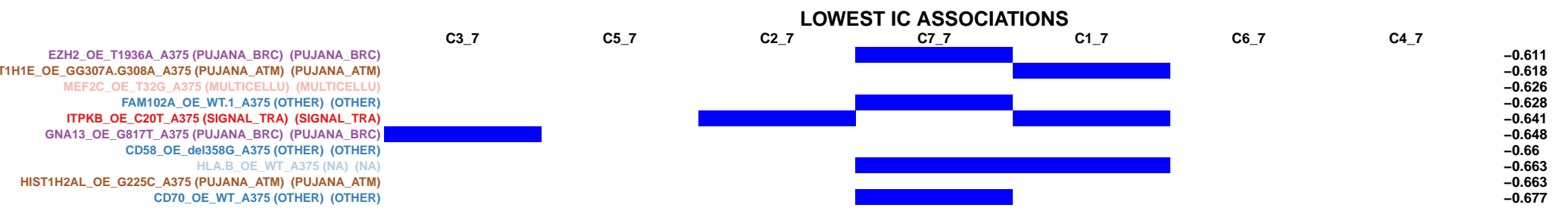
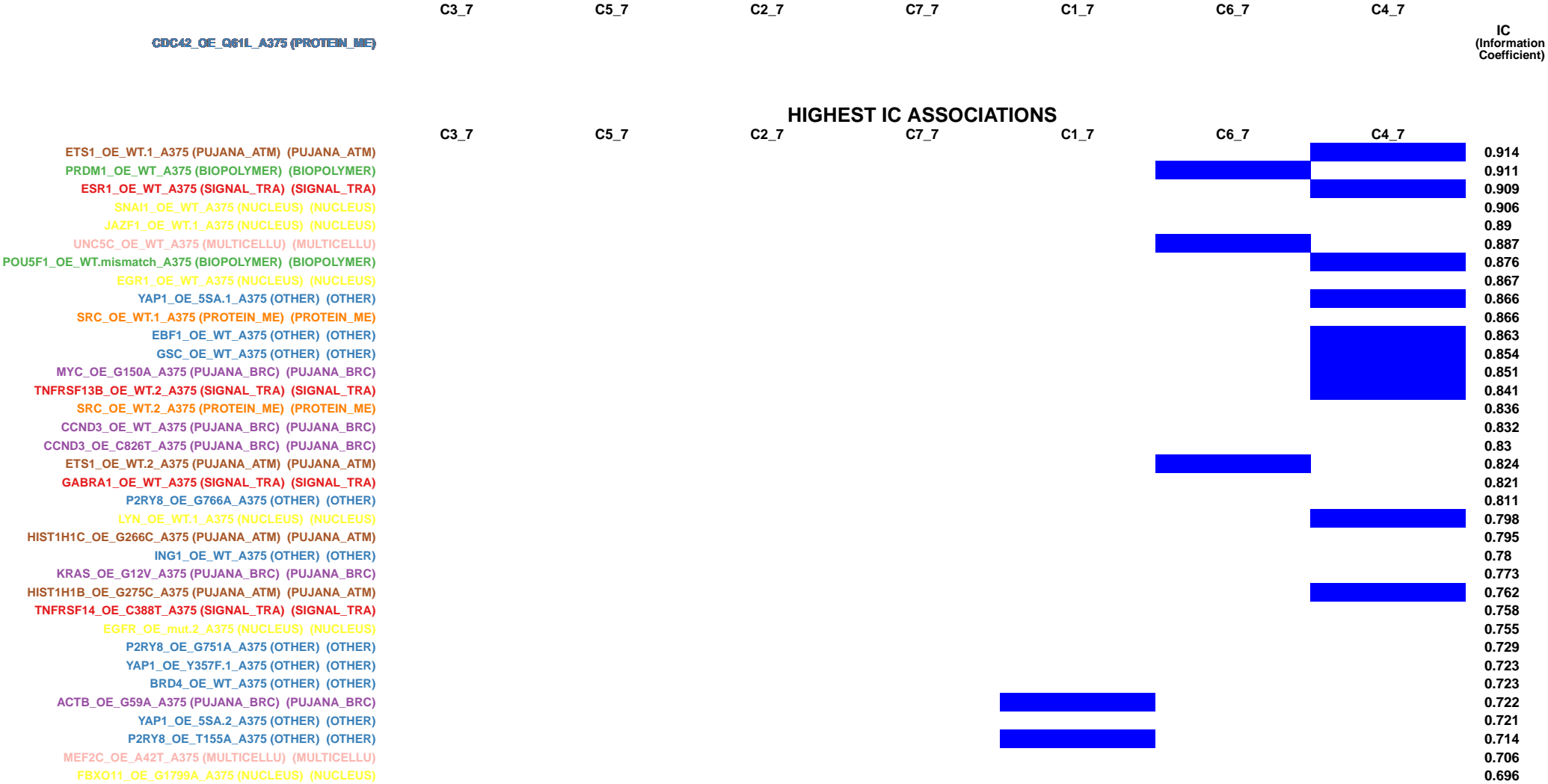




BFP\_OE\_CTRL\_A375 (OTHER)

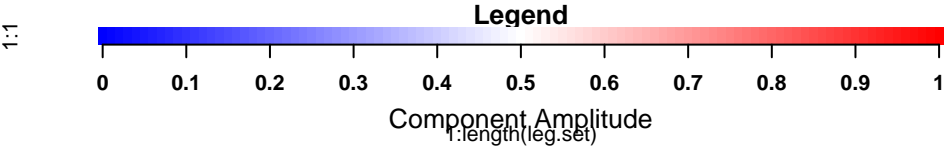
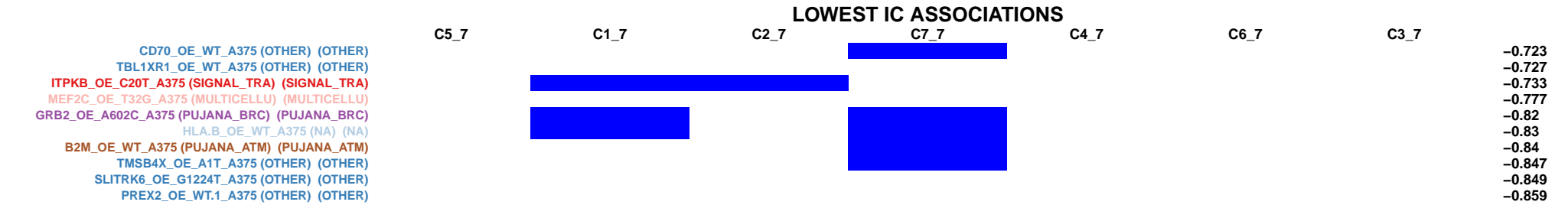
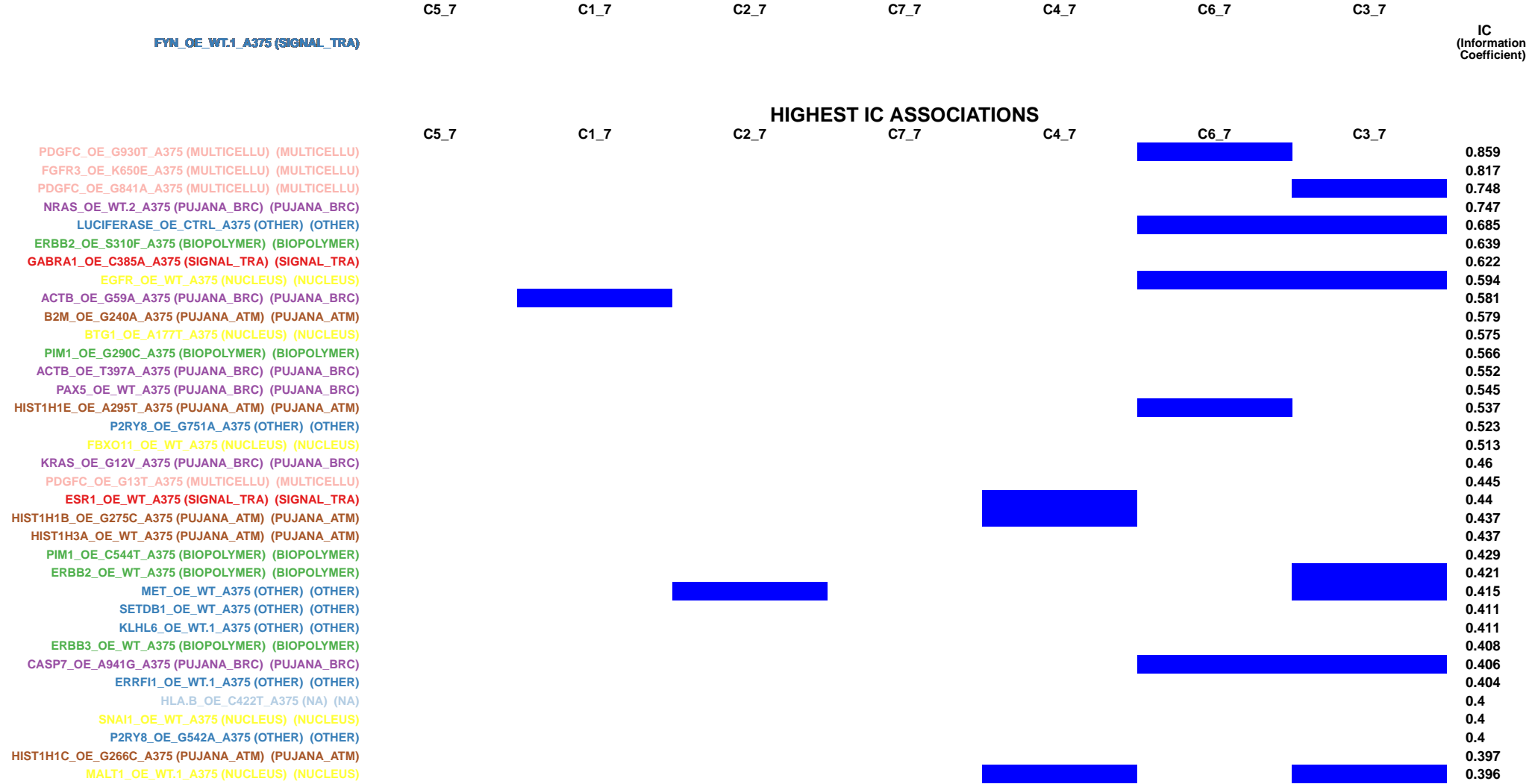


CDC42\_OE\_Q61L\_A375 (PROTEIN\_ME)

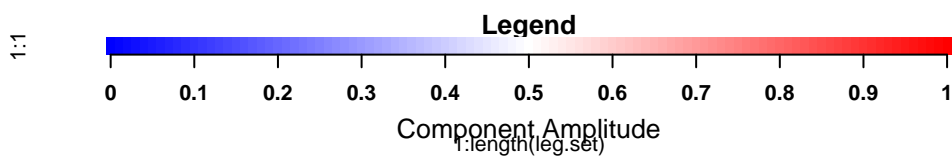
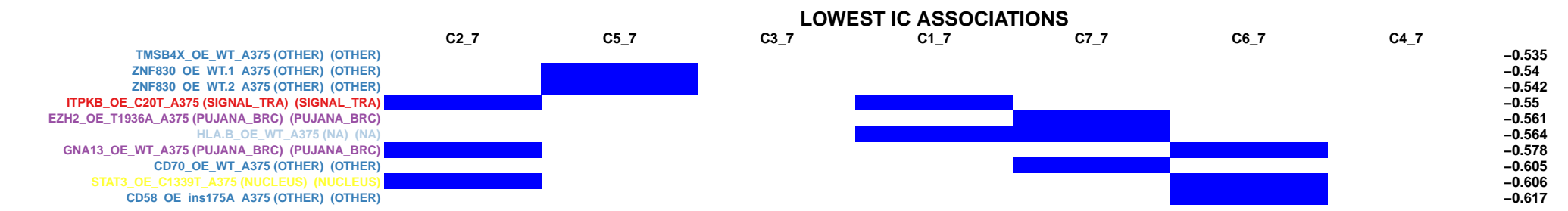
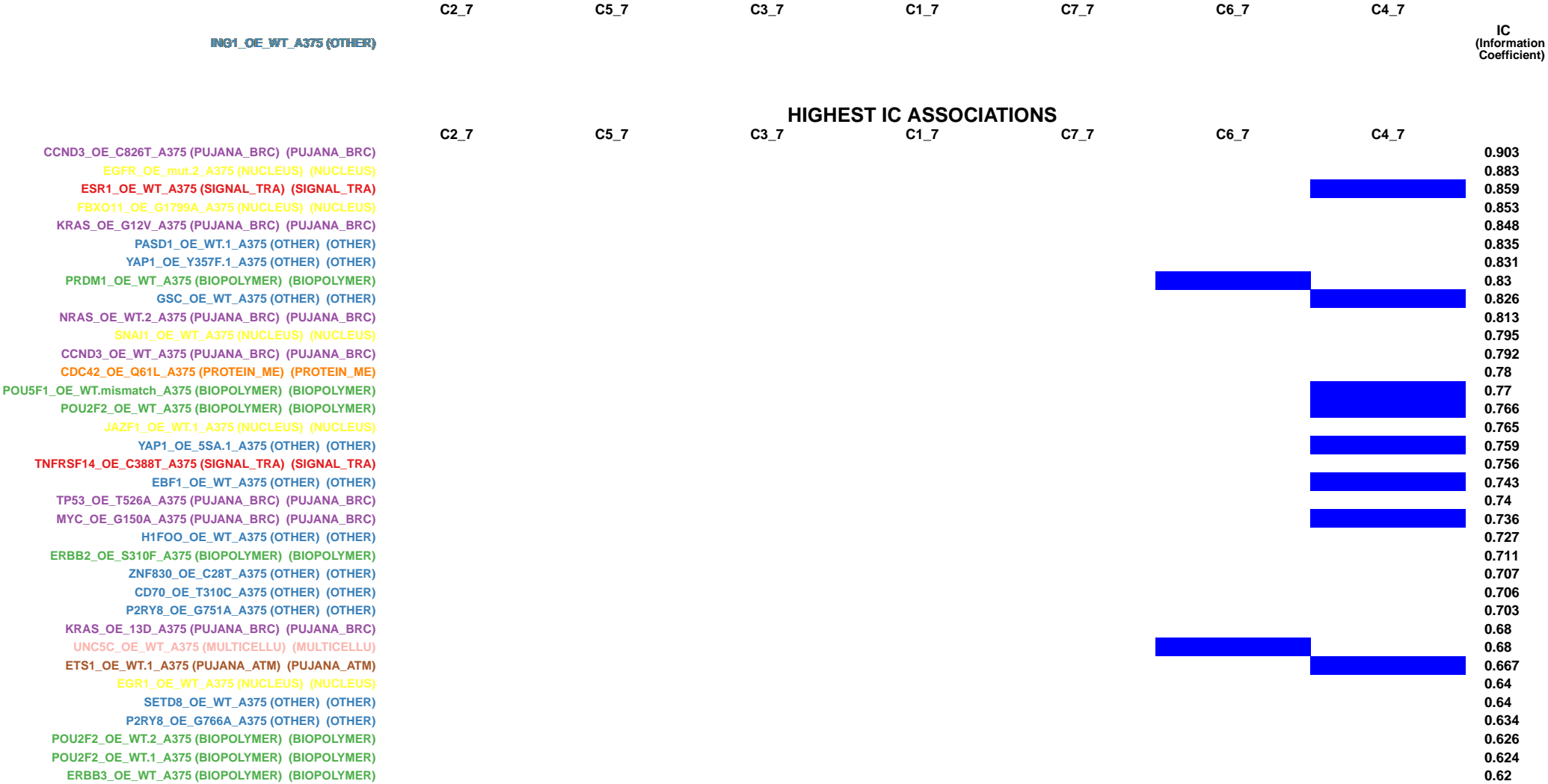


FYN\_OE\_WT.1\_A375 (SIGNAL\_TRA)

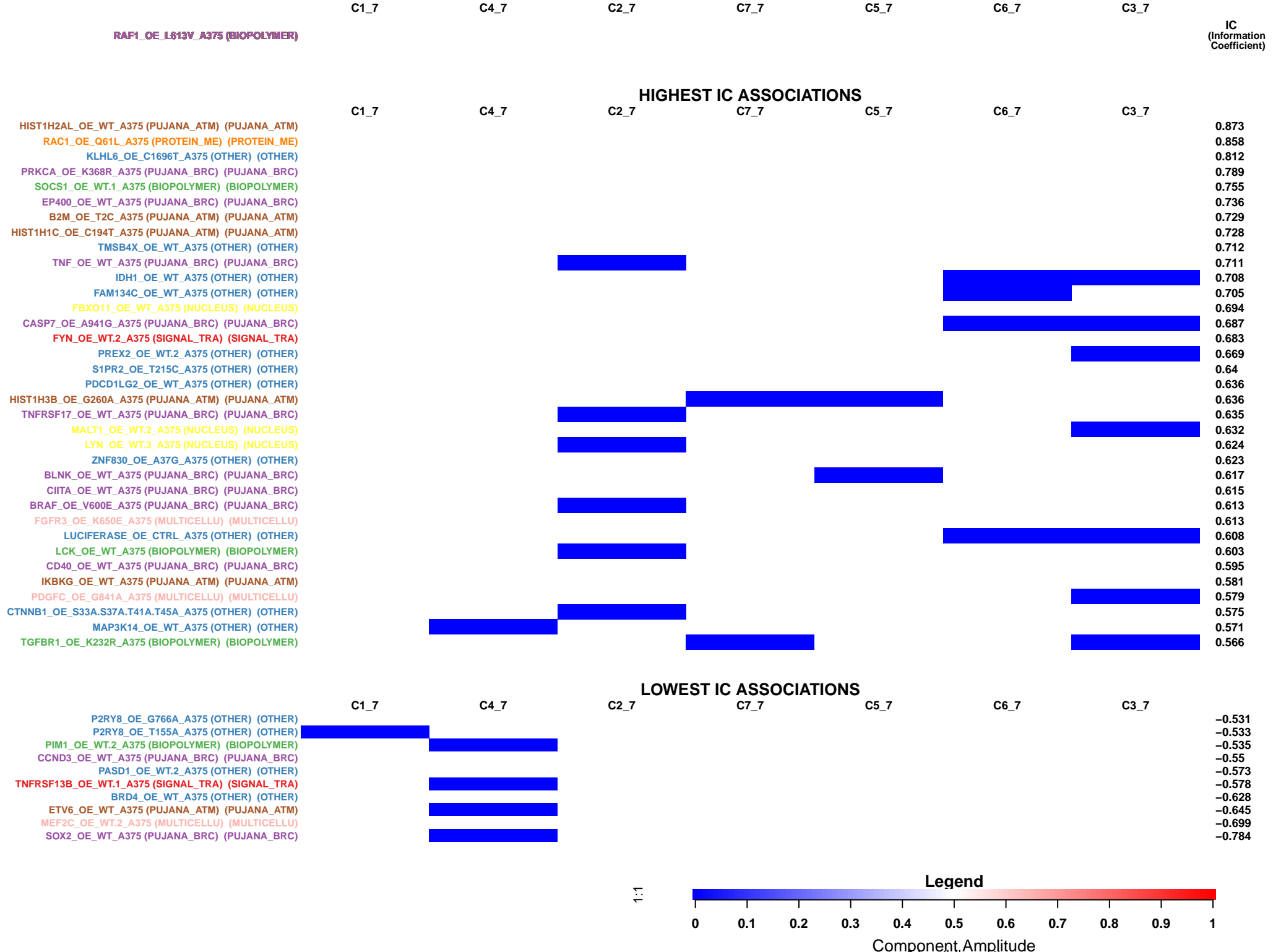
FYN\_OE\_WT.1\_A375 (SIGNAL\_TRA)



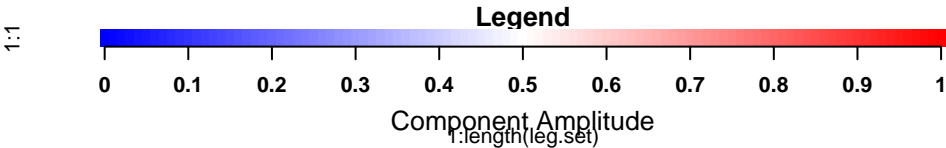
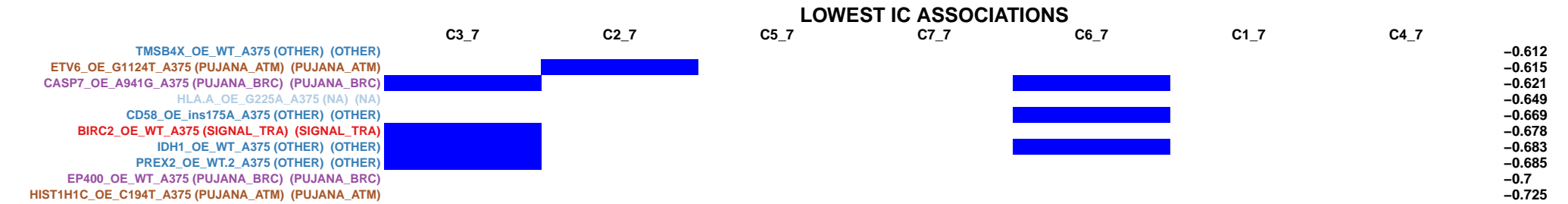
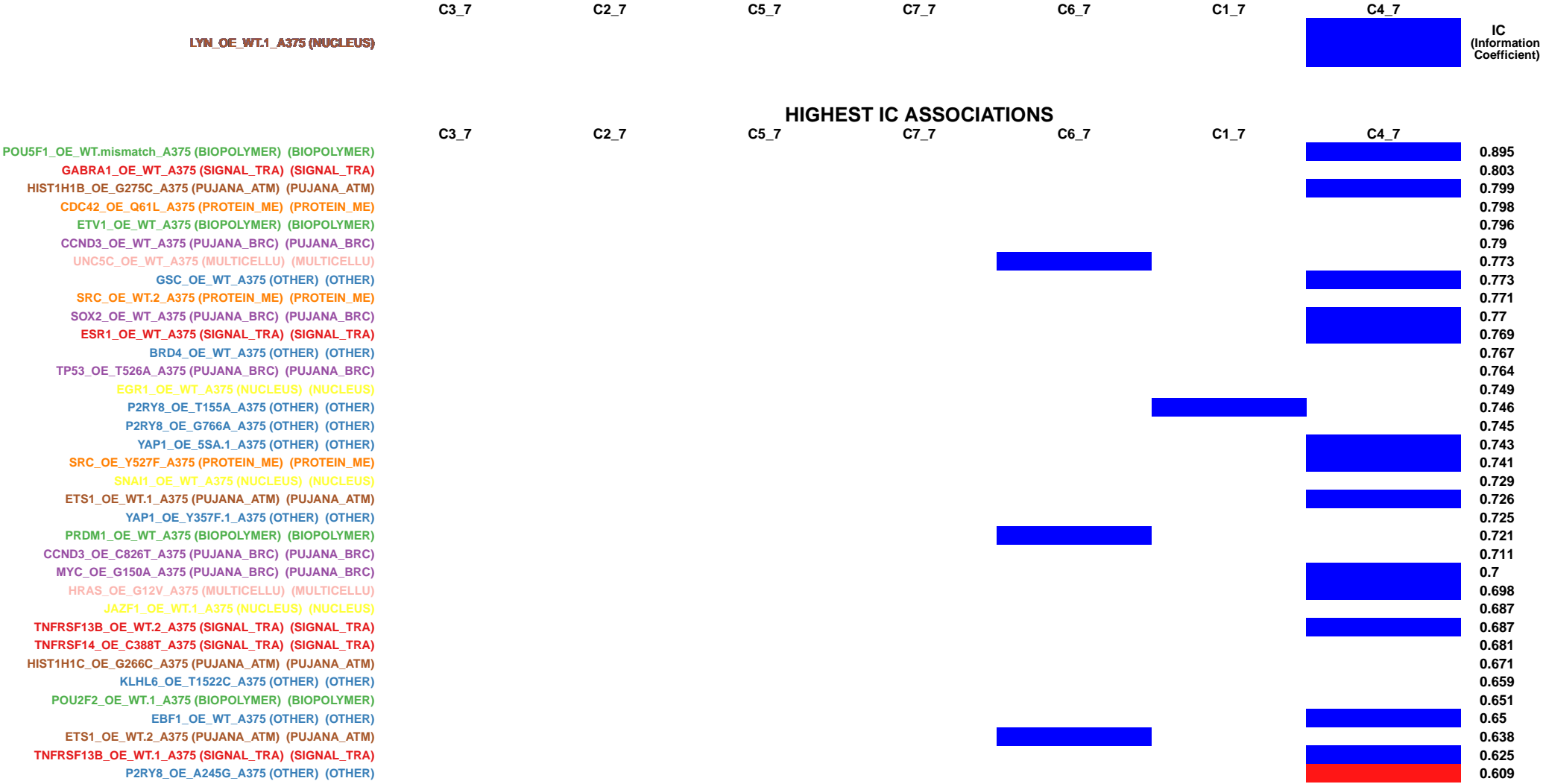
ING1\_OE\_WT\_A375 (OTHER)



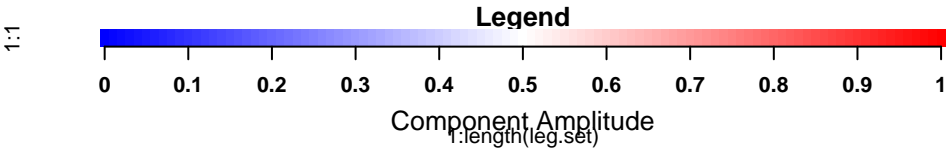
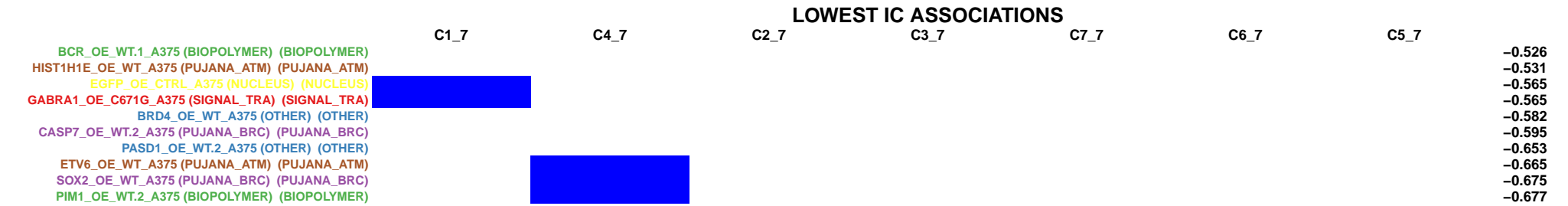
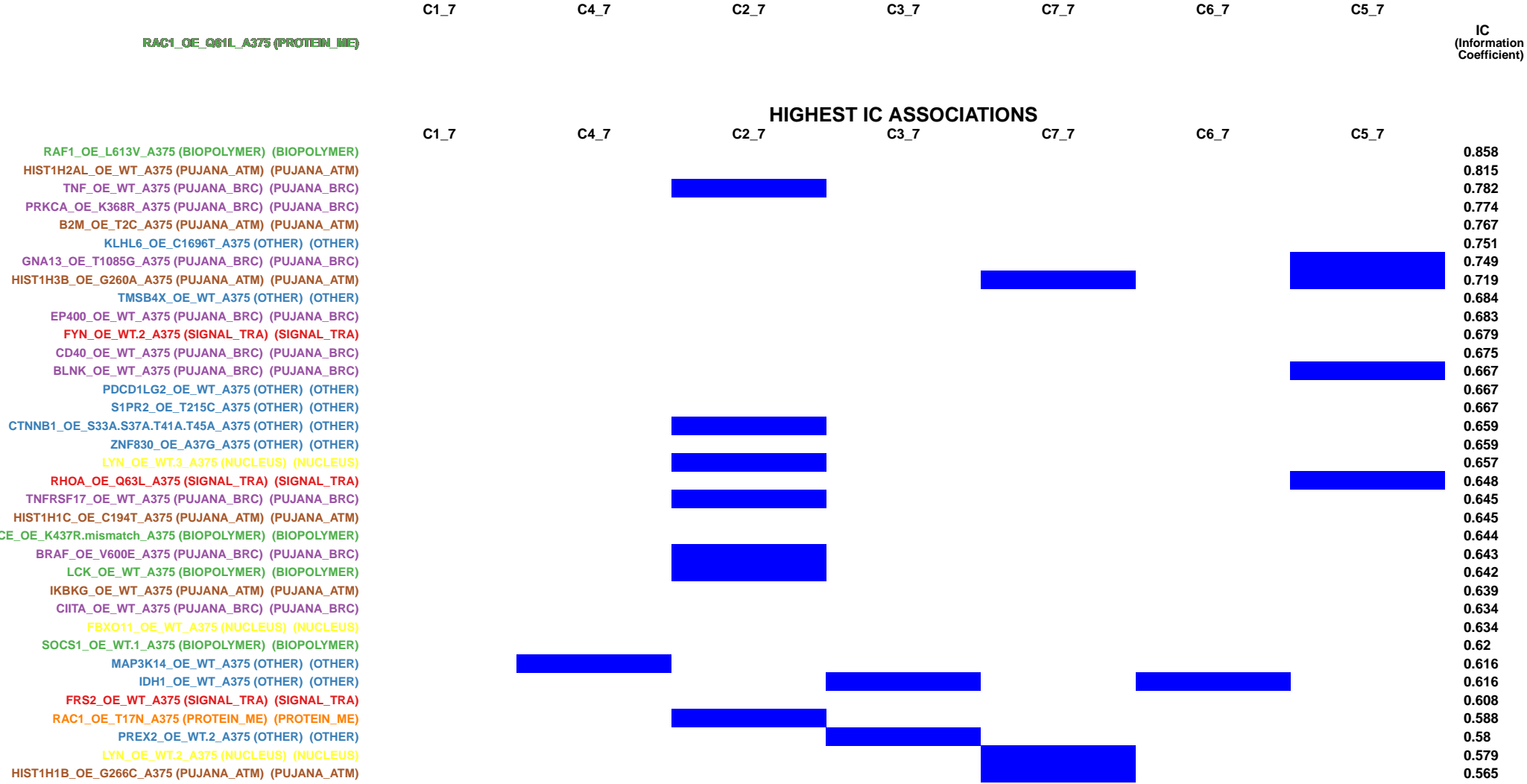
RAF1\_OE\_L613V\_A375 (BIOPOLYMER)



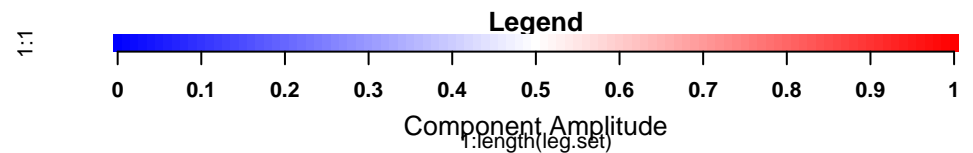
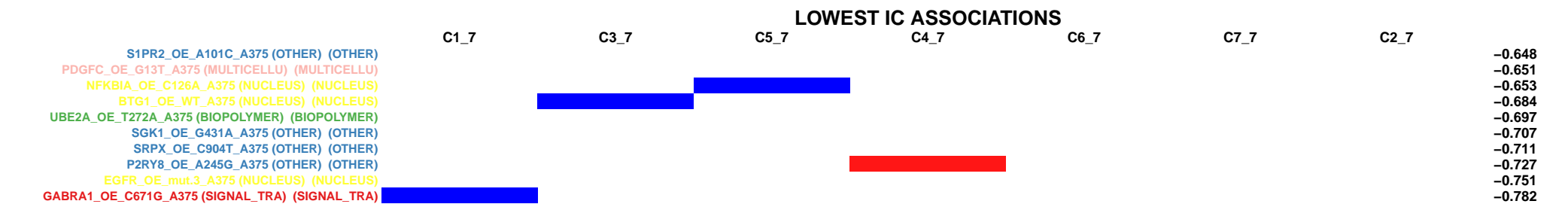
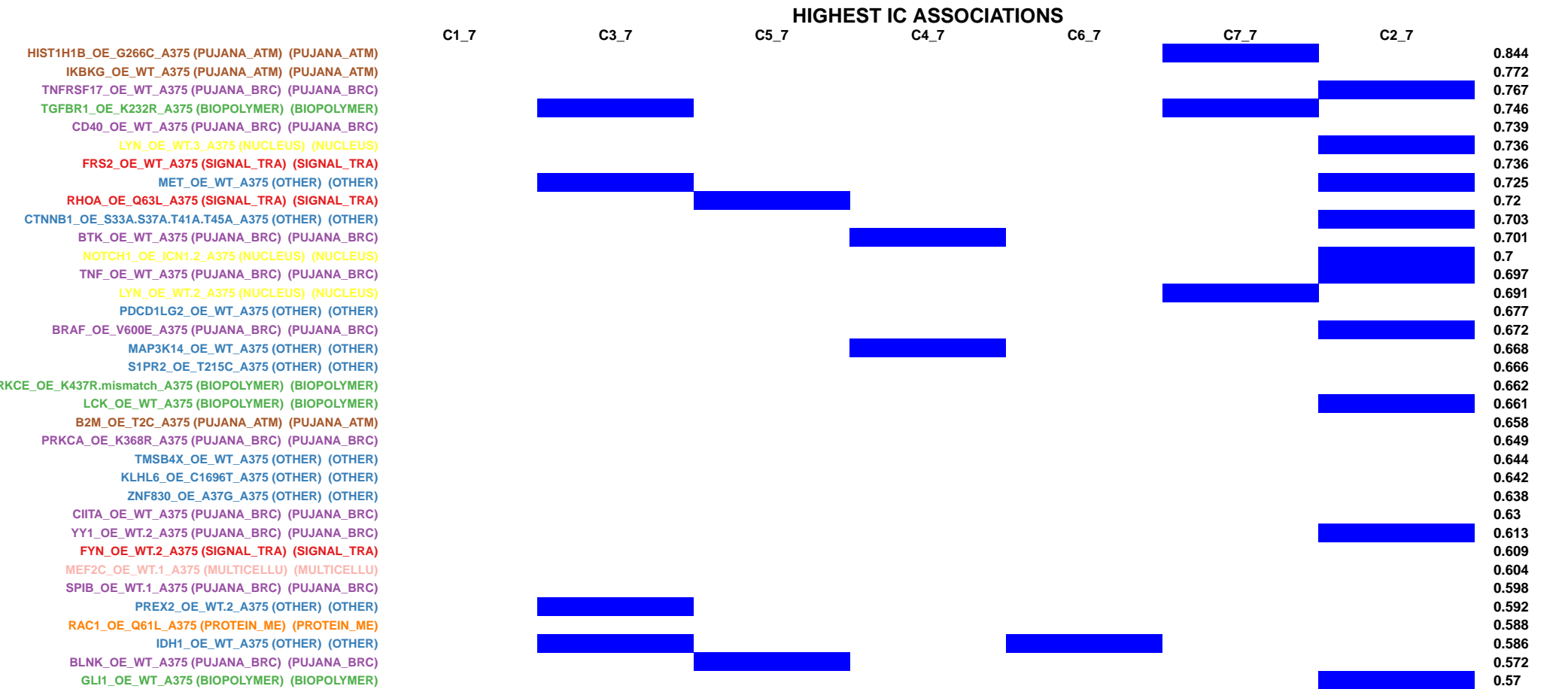
LYN\_OE\_WT.1\_A375 (NUCLEUS)



RAC1\_OE\_Q61L\_A375 (PROTEIN\_ME)



C1_7	C3_7	C5_7	C4_7	C6_7	C7_7	C2_7	IC (Information Coefficient)

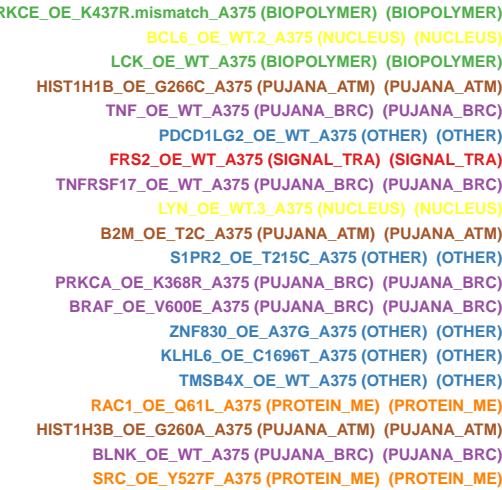




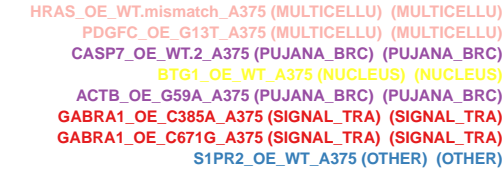
C1_7	C3_7	C6_7	C4_7	C7_7	C5_7	C2_7
------	------	------	------	------	------	------

0.868  
0.861  
0.844  
0.829  
0.828  
0.8  
0.79  
0.778  
0.772  
0.767  
0.761  
0.748  
0.745  
0.742  
0.736  
0.733  
0.727  
0.707  
0.705  
0.701  
0.699  
0.691  
0.69  
0.686  
0.671  
0.671  
0.665  
0.665  
0.66  
0.658  
0.649  
0.639  
0.637  
0.635  
0.631

C1\_7                      C3\_7                      C6\_7                      C4\_7                      C7\_7                      C5\_7                      C2\_7



C1_7	C3_7	C6_7	C4_7	C7_7	C5_7	C2_7
------	------	------	------	------	------	------

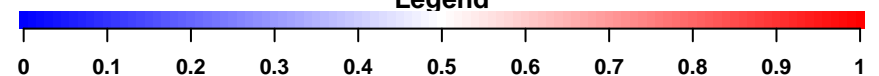


Legend

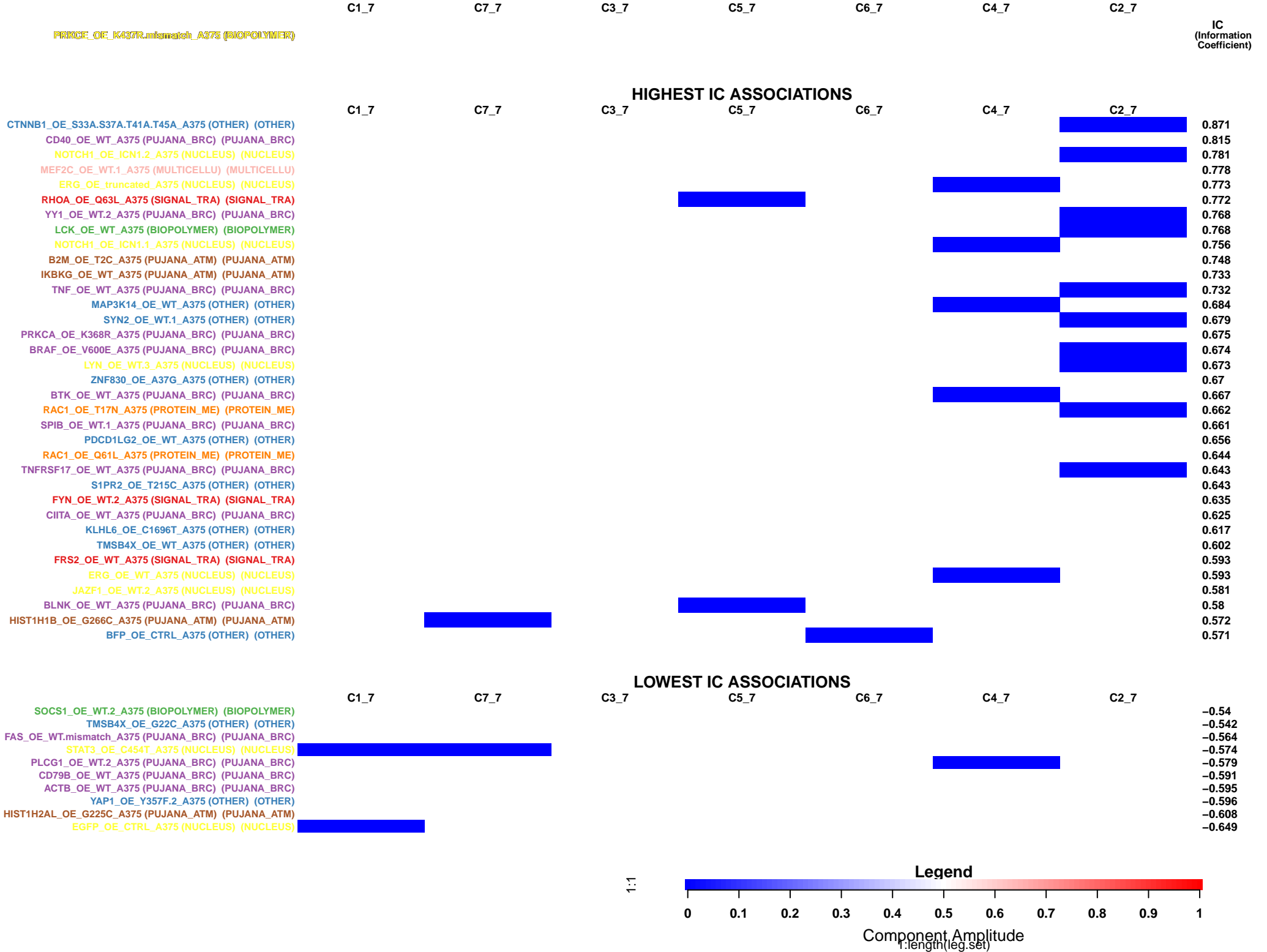
0.4 0.5 0.6

Component Amplitude

1:length(leg.set)



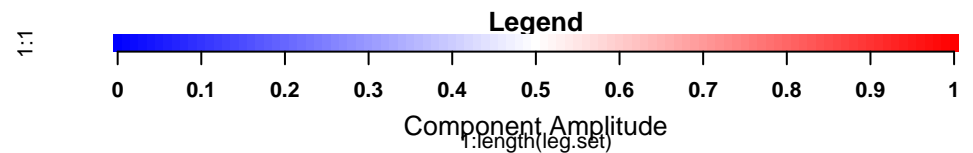
PRKCE\_OE\_K437R.mismatch\_A375 (BIOPOLYMER)



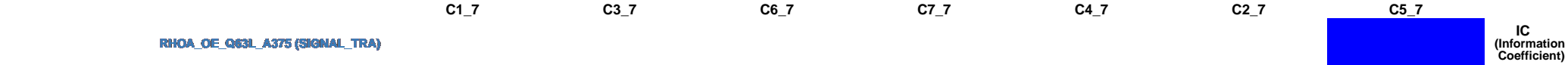
C3_7	C6_7	C1_7	C7_7	C5_7	C2_7	C4_7	IC (Information Coefficient)
------	------	------	------	------	------	------	---------------------------------

C3_7	C6_7	C1_7	C7_7	C5_7	C2_7	C4_7
						0.922
						0.868
						0.833
						0.832
						0.821
						0.816
						0.801
						0.8
						0.746
						0.739
						0.714
						0.708
						0.69
						0.688
						0.675
						0.64
						0.632
						0.631
						0.626
						0.623
						0.618
						0.617
						0.613
						0.611
						0.608
						0.547
						0.547
						0.544
						0.532
						0.529
						0.527
						0.514
						0.476
						0.475
						0.467

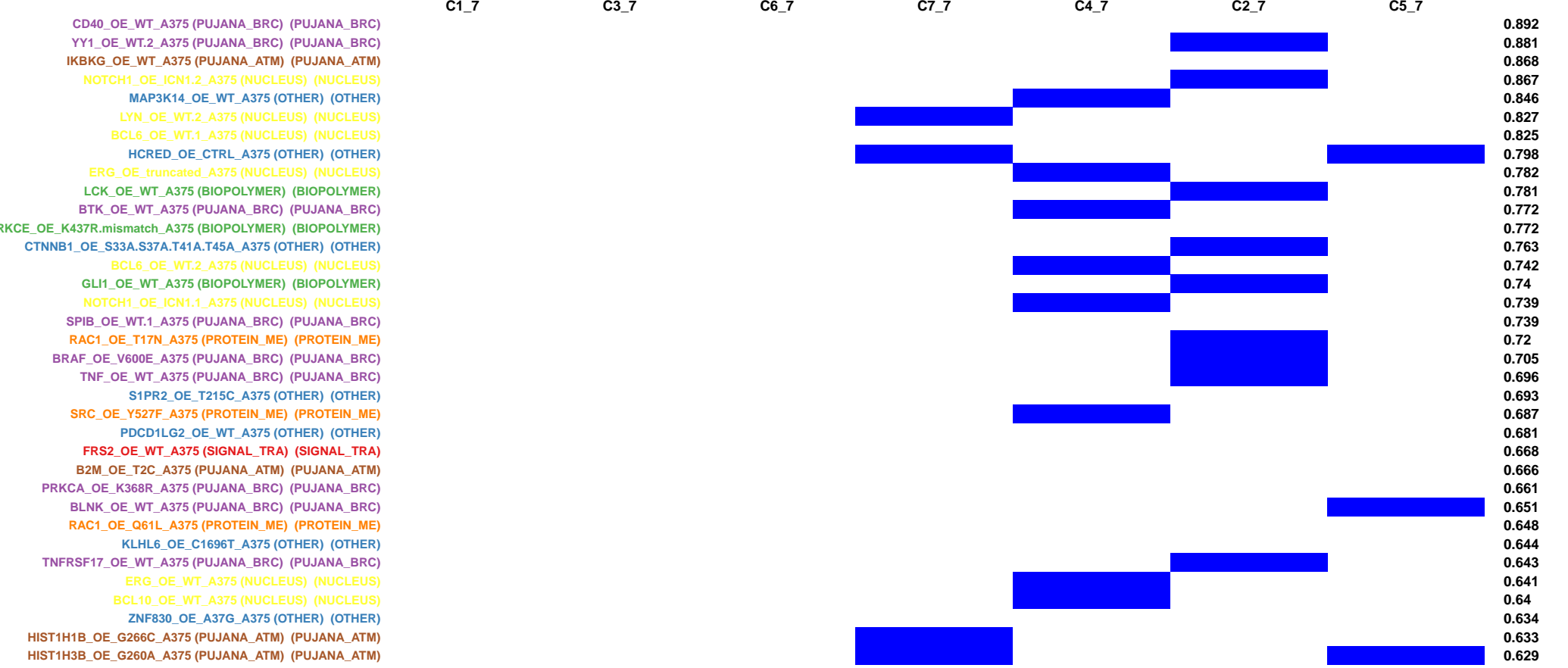
	C2_7	C3_7	C4_7	C5_7	C6_7	C7_7	C1_7
PDGFC_OE_G13T_A375 (MULTICELLU) (MULTICELLU)	-0.551						
SOC51_OE_WT.1_A375 (BIOPOLYMER) (BIOPOLYMER)	-0.558						
HLA.A_OE_G225A_A375 (NA) (NA)	-0.564						
LUCIFERASE_OE_CTRL_A375 (OTHER) (OTHER)	-0.566						
RNF2_OE_WT_A375 (PUJANA_BRC) (PUJANA_BRC)	-0.658						
CPS1_OE_T4433G.mismatch_A375 (OTHER) (OTHER)	-0.663						
S1PR2_OE_WT_A375 (OTHER) (OTHER)	-0.677						
SLITRK6_OE_A1742C_A375 (OTHER) (OTHER)	-0.713						
NFKBIA_OE_G33A_A375 (NUCLEUS) (NUCLEUS)	-0.735						
TMSB4X_OE_del63.64AG_A375 (OTHER) (OTHER)	-0.809						



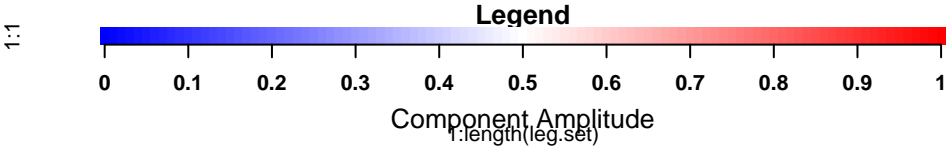
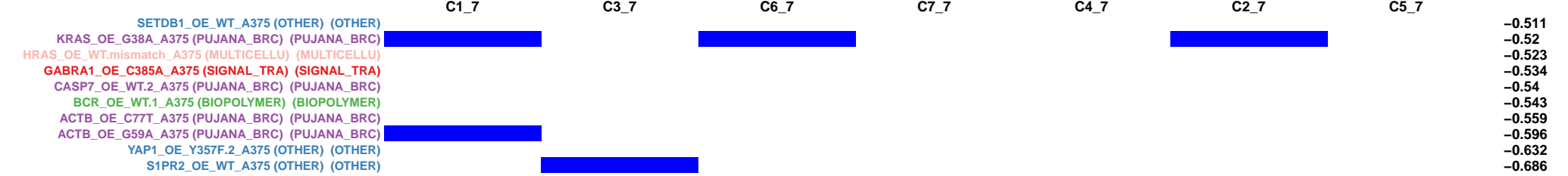
RHOA\_OE\_Q63L\_A375 (SIGNAL\_TRA)



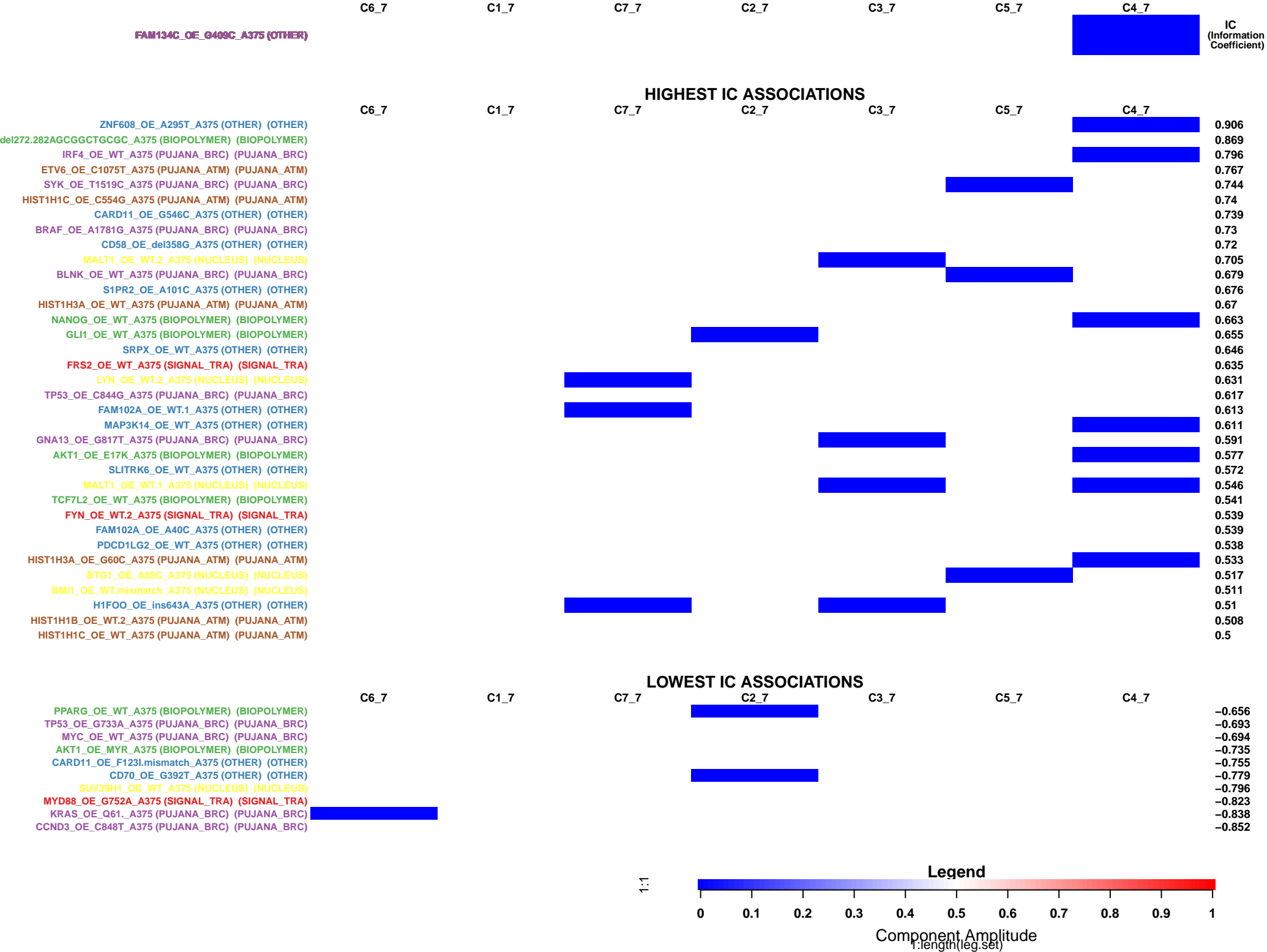
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS



FAM134C\_OE\_G409C\_A375 (OTHER)



C2\_7                      C4\_7                      C1\_7                      C5\_7                      C6\_7                      C3\_7                      C7\_7

**SOCS1\_OE\_WT.1\_A375 (BIOPOLYMER)**

C2\_7                      C4\_7                      C1\_7                      C5\_7                      C6\_7                      C3\_7                      C7\_7



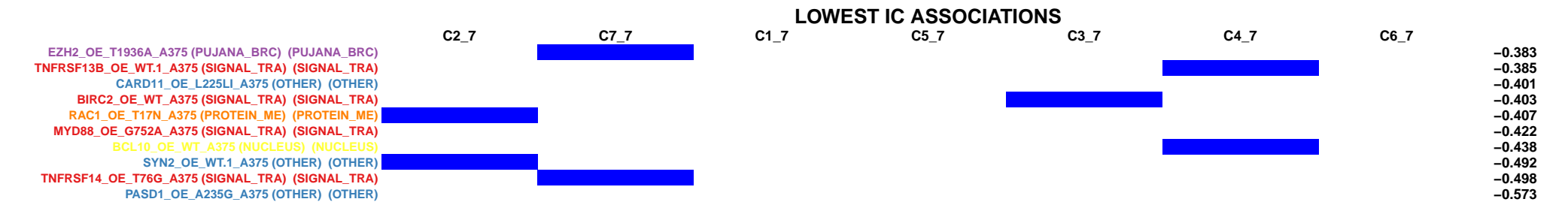
**C2\_7                      C4\_7                      C1\_7                      C5\_7                      C6\_7                      C3\_7                      C7\_7**



0      0.1      0.2      0.3      0.4      0.5      0.6      0.7      0.8      0.9      1

Component Amplitude  
1:length(leg.set)

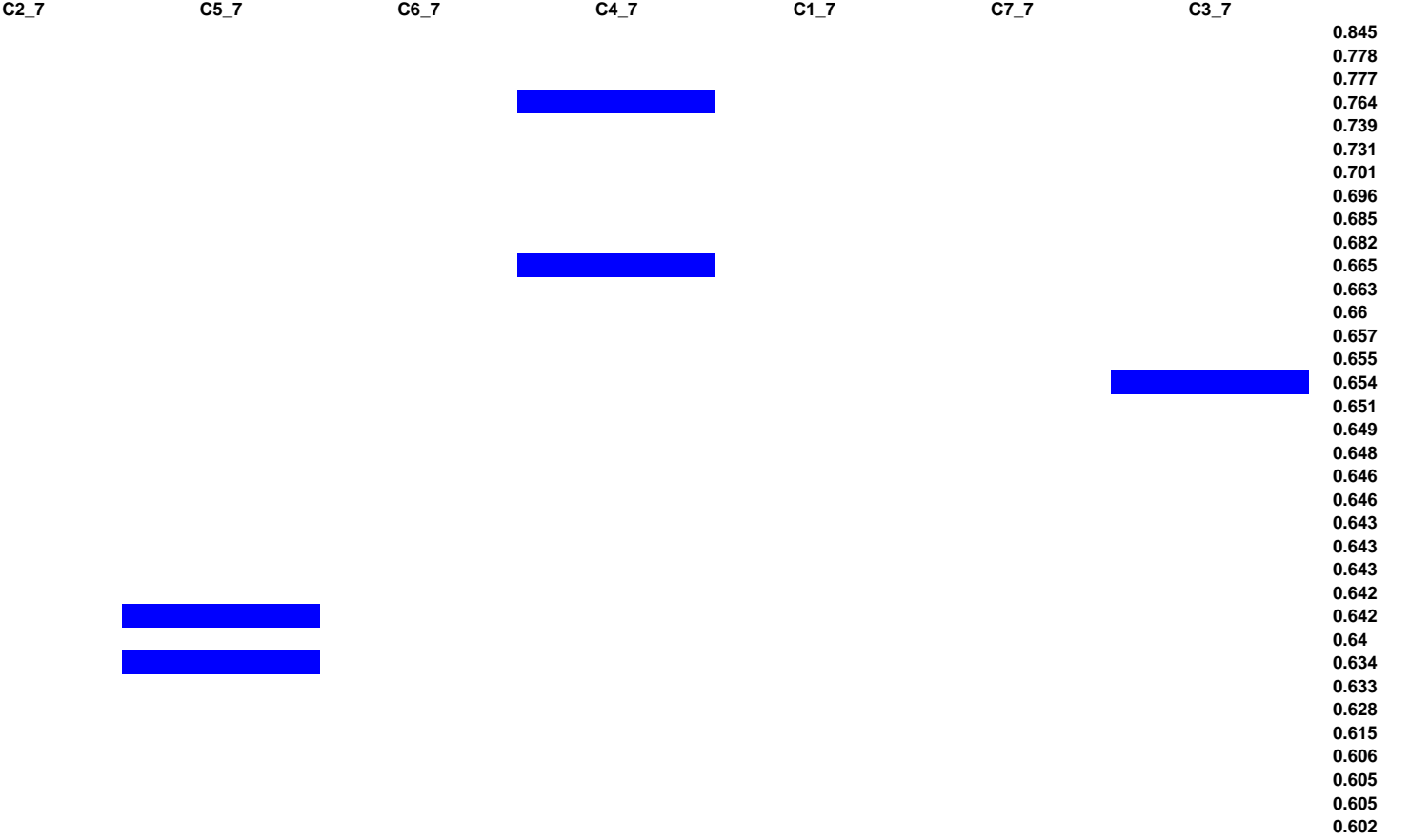
C2_7	C7_7	C1_7	C5_7	C3_7	C4_7	C6_7	IC (Information Coefficient)
------	------	------	------	------	------	------	---------------------------------



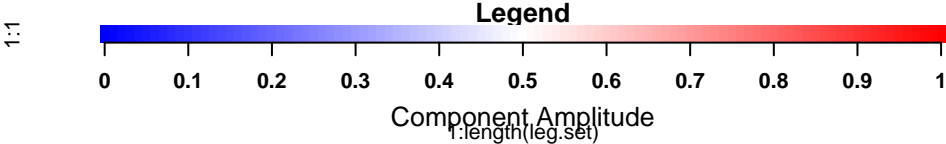
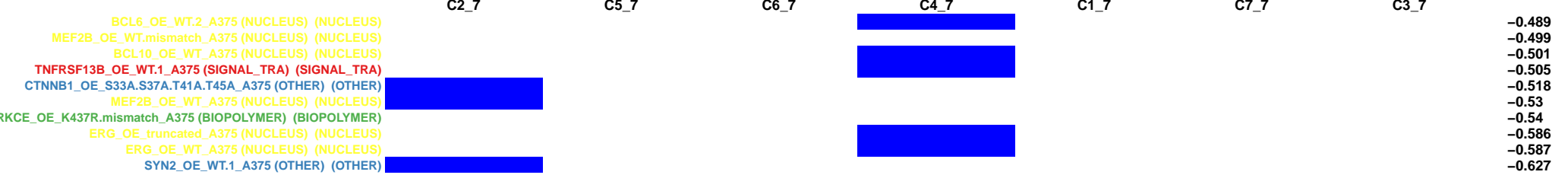
SOCS1\_OE\_WT.2\_A375 (BIOPOLYMER)

C2\_7C5\_7C6\_7C4\_7C1\_7C7\_7C3\_7IC  
(Information  
Coefficient)

HIGHEST IC ASSOCIATIONS

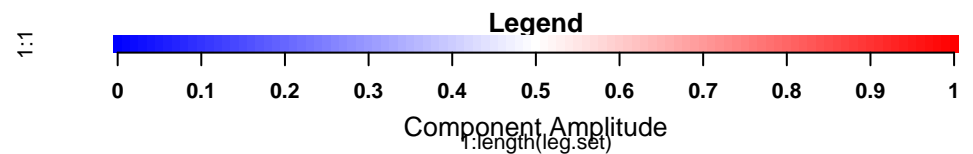
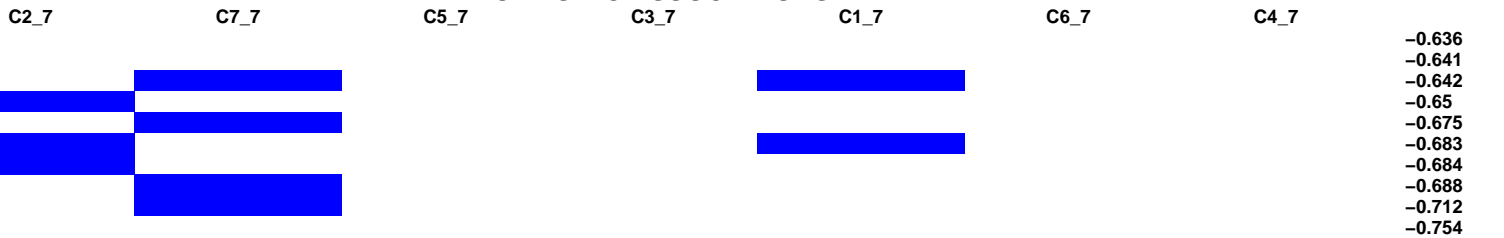
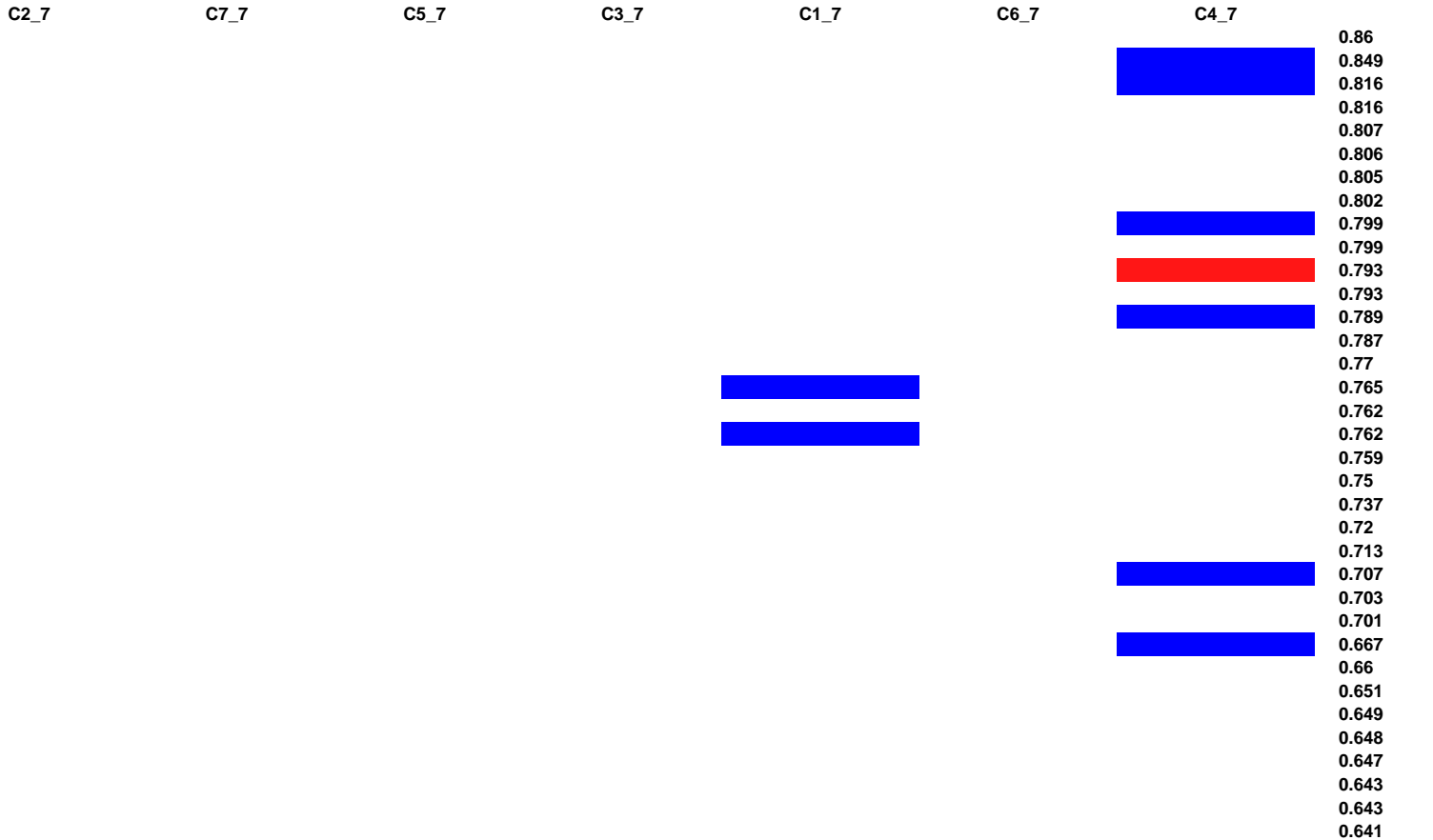


LOWEST IC ASSOCIATIONS





C2_7	C7_7	C5_7	C3_7	C1_7	C6_7	C4_7	IC (Information Coefficient)
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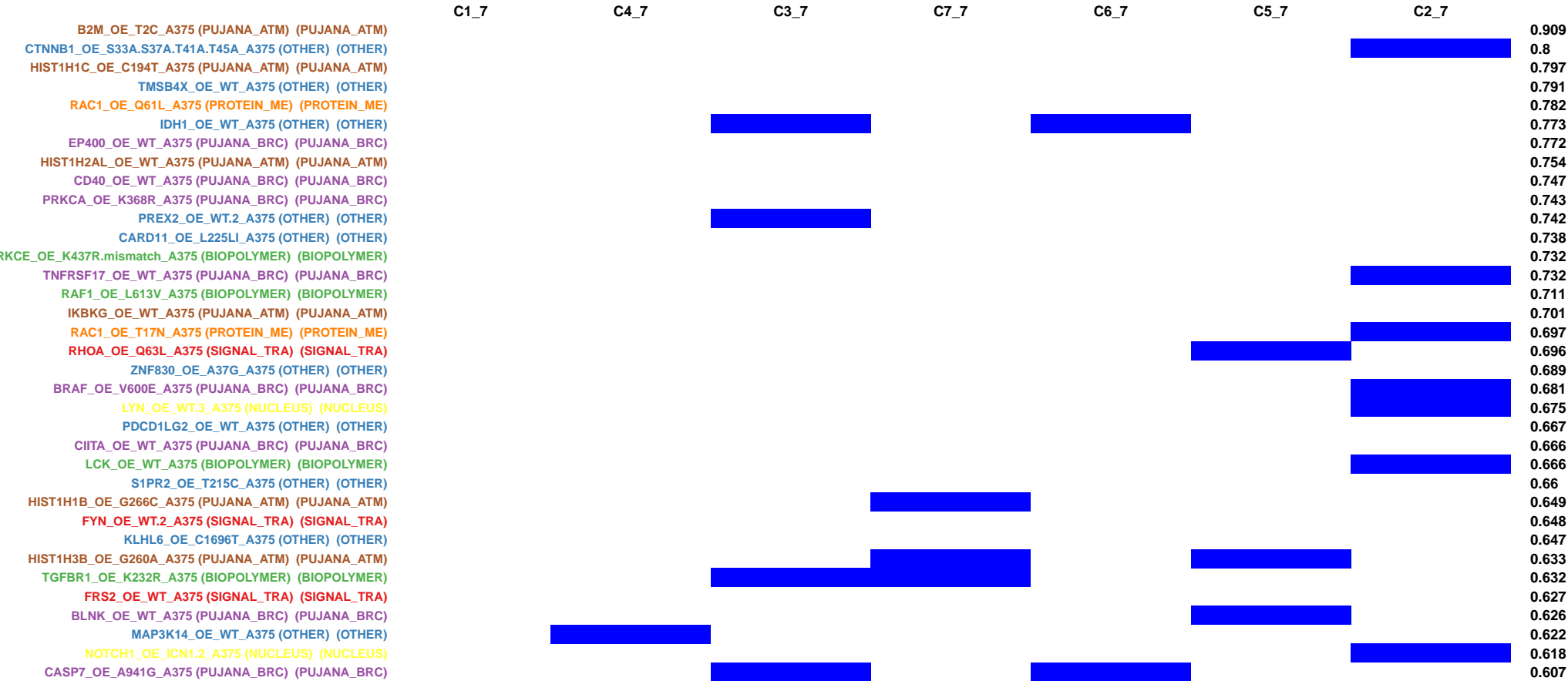


TNF\_OE\_WT\_A375 (PUJANA\_BRC)

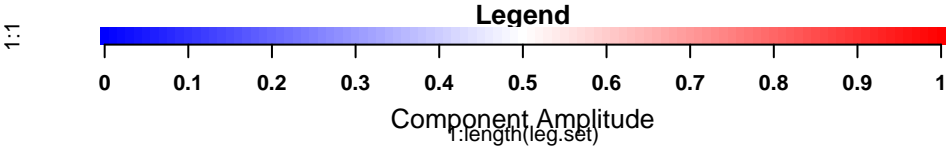
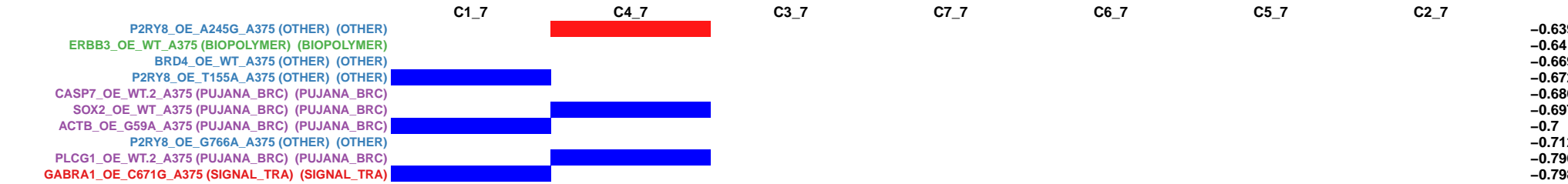
TNF\_OE\_WT\_A375 (PUJANA\_BRC)

IC  
(Information  
Coefficient)

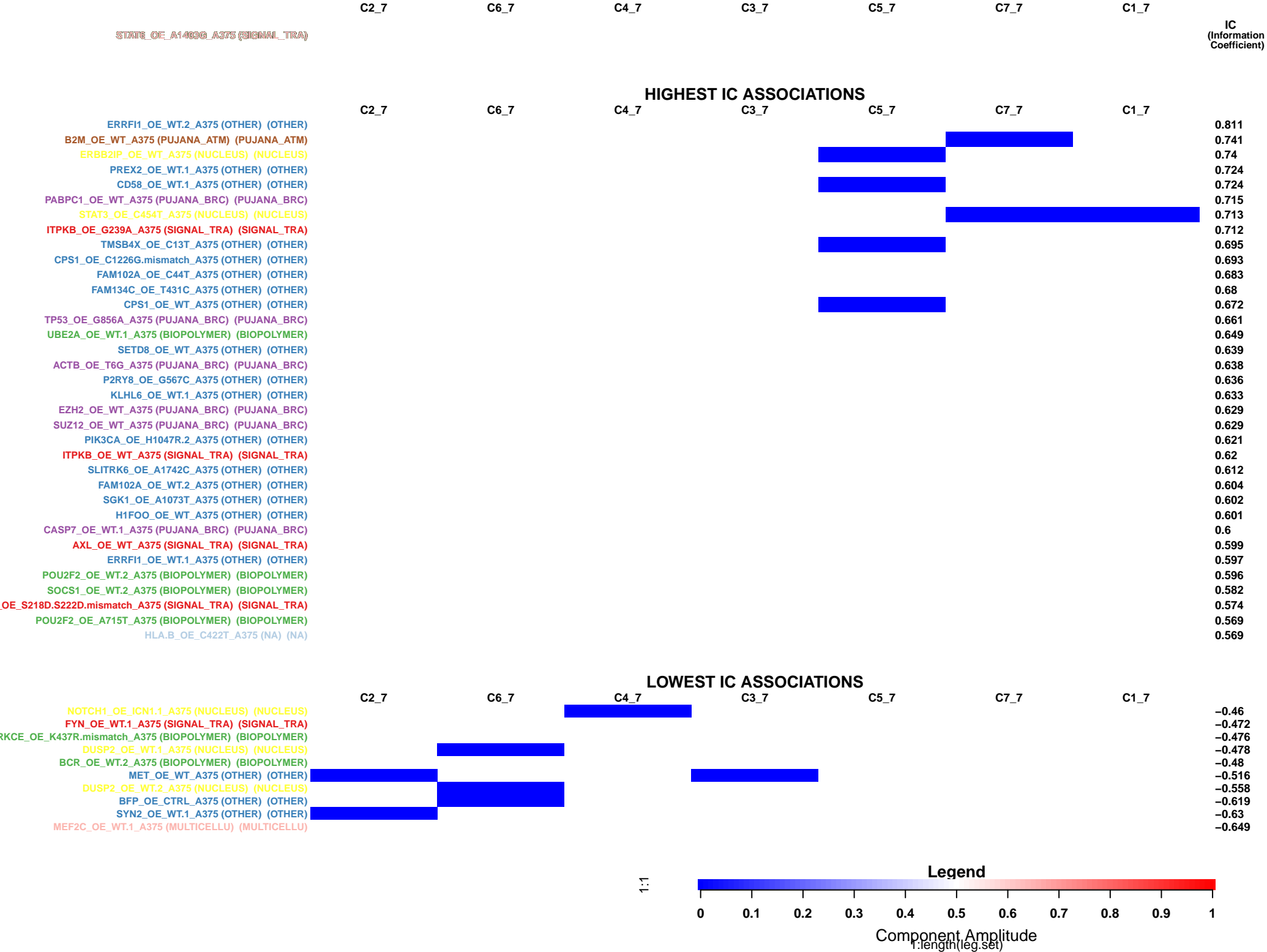
HIGHEST IC ASSOCIATIONS



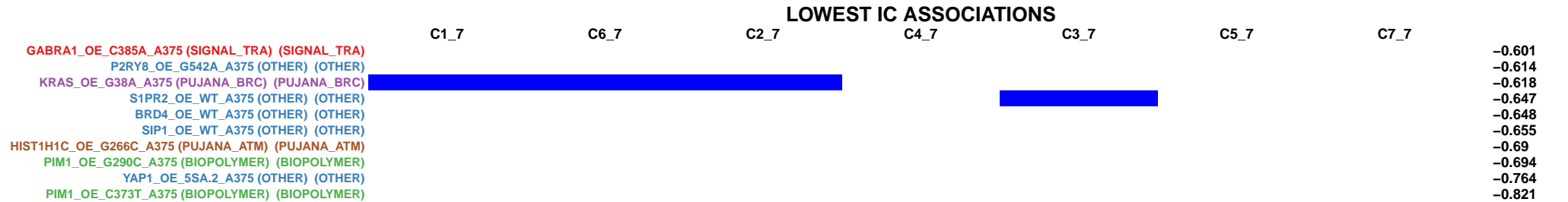
LOWEST IC ASSOCIATIONS



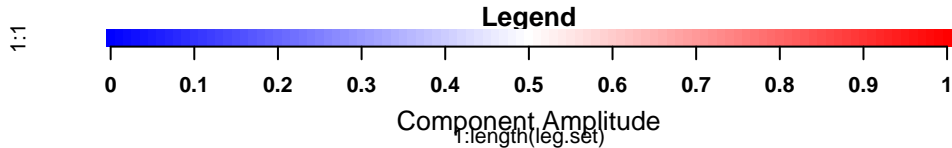
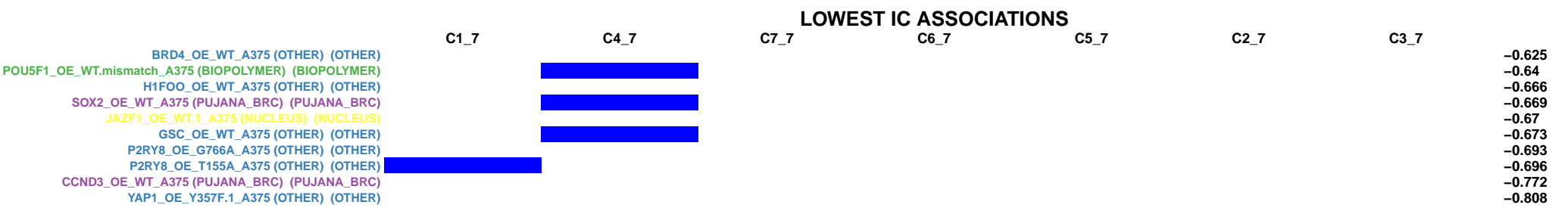
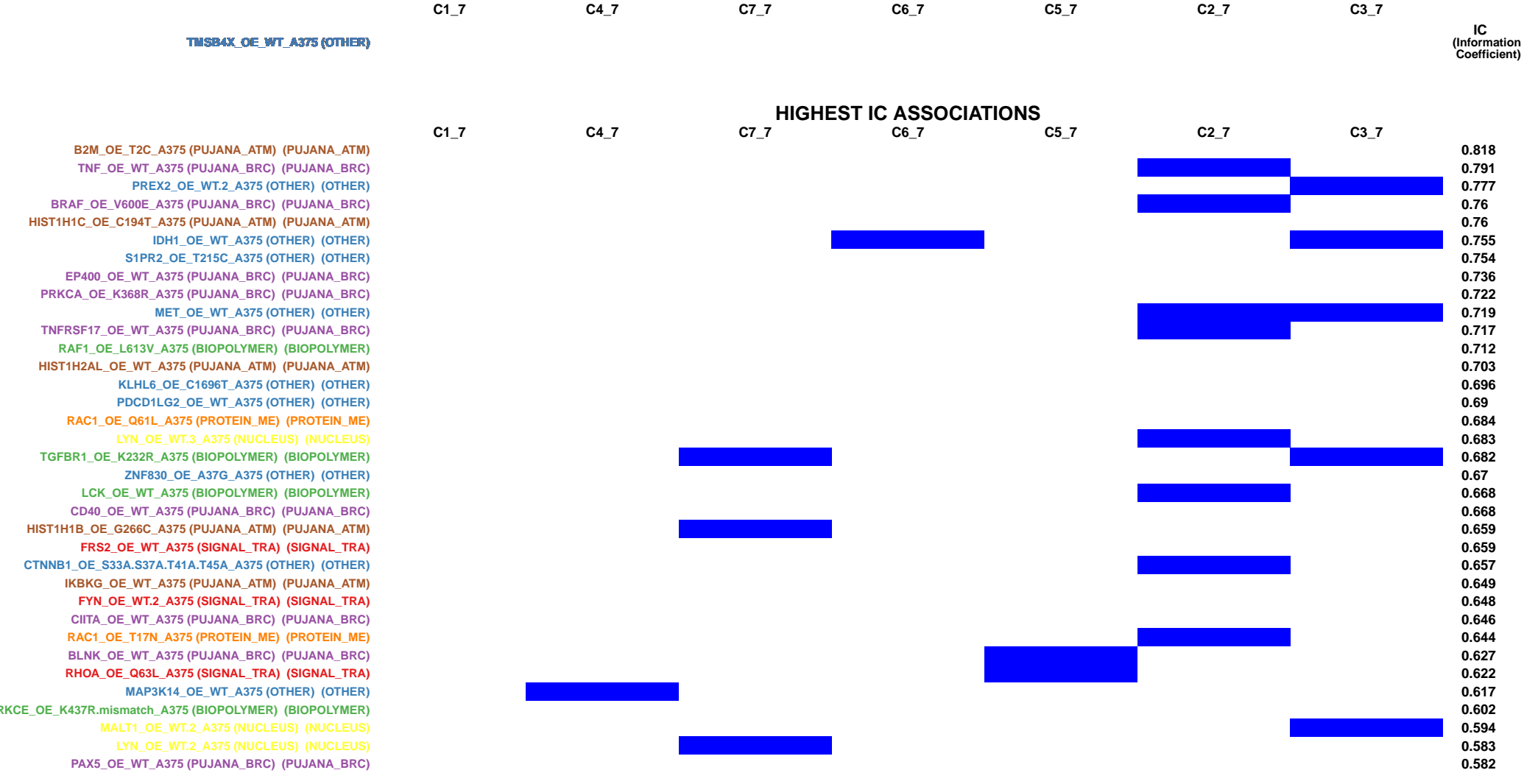
STAT6\_OE\_A1403G\_A375 (SIGNAL\_TRA)



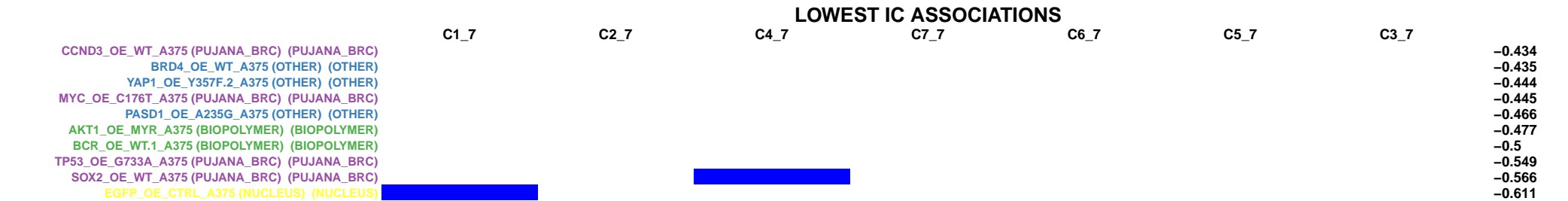
C1_7	C6_7	C2_7	C4_7	C3_7	C5_7	C7_7	IC (Information Coefficient)



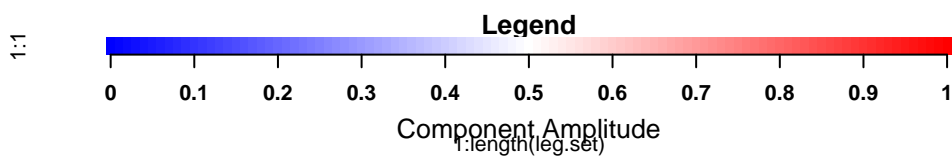
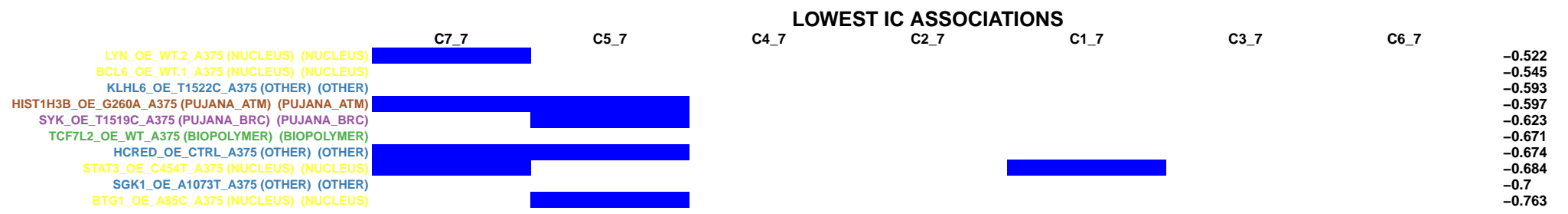
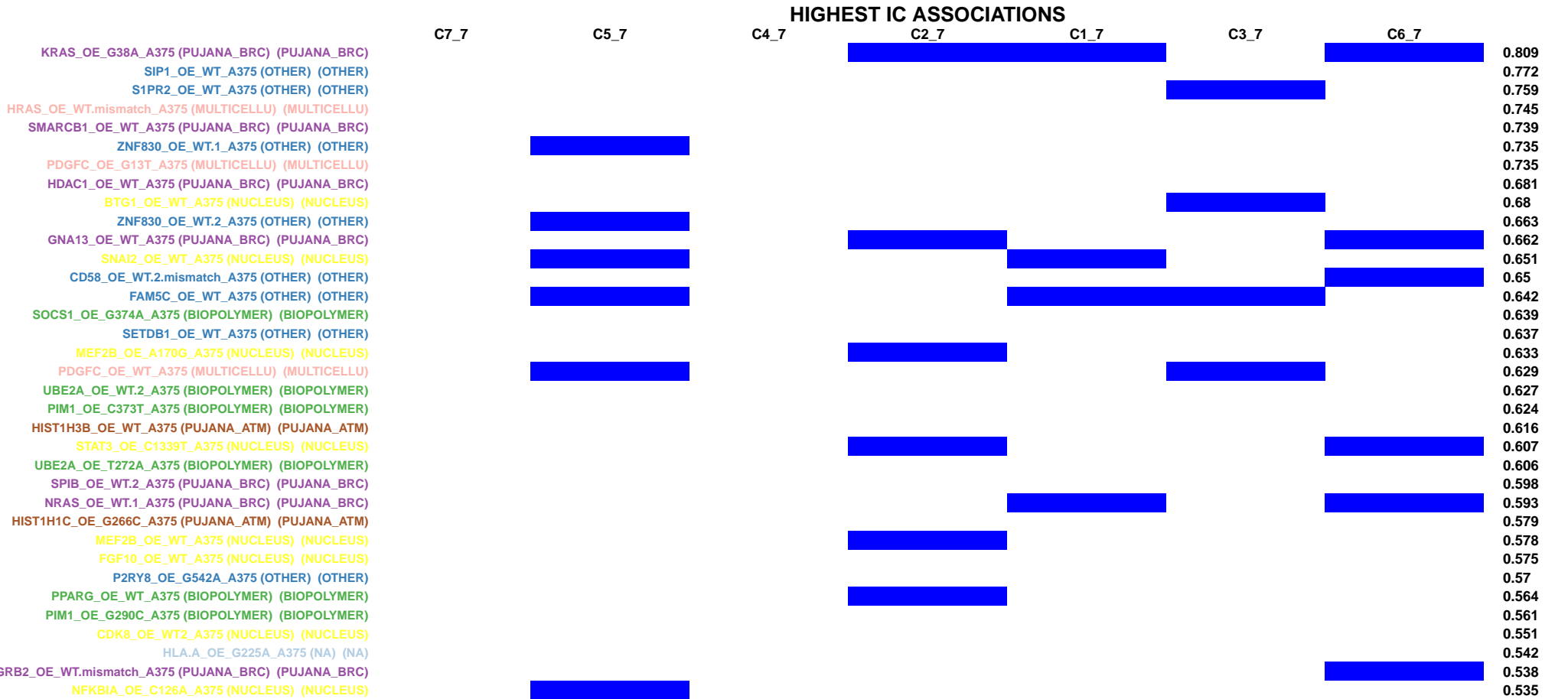
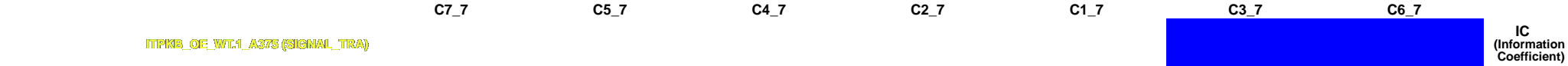
TMSB4X\_OE\_WT\_A375 (OTHER)



C1_7	C2_7	C4_7	C7_7	C6_7	C5_7	C3_7	IC (Information Coefficient)
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ITPKB\_OE\_WT.1\_A375 (SIGNAL\_TRA)

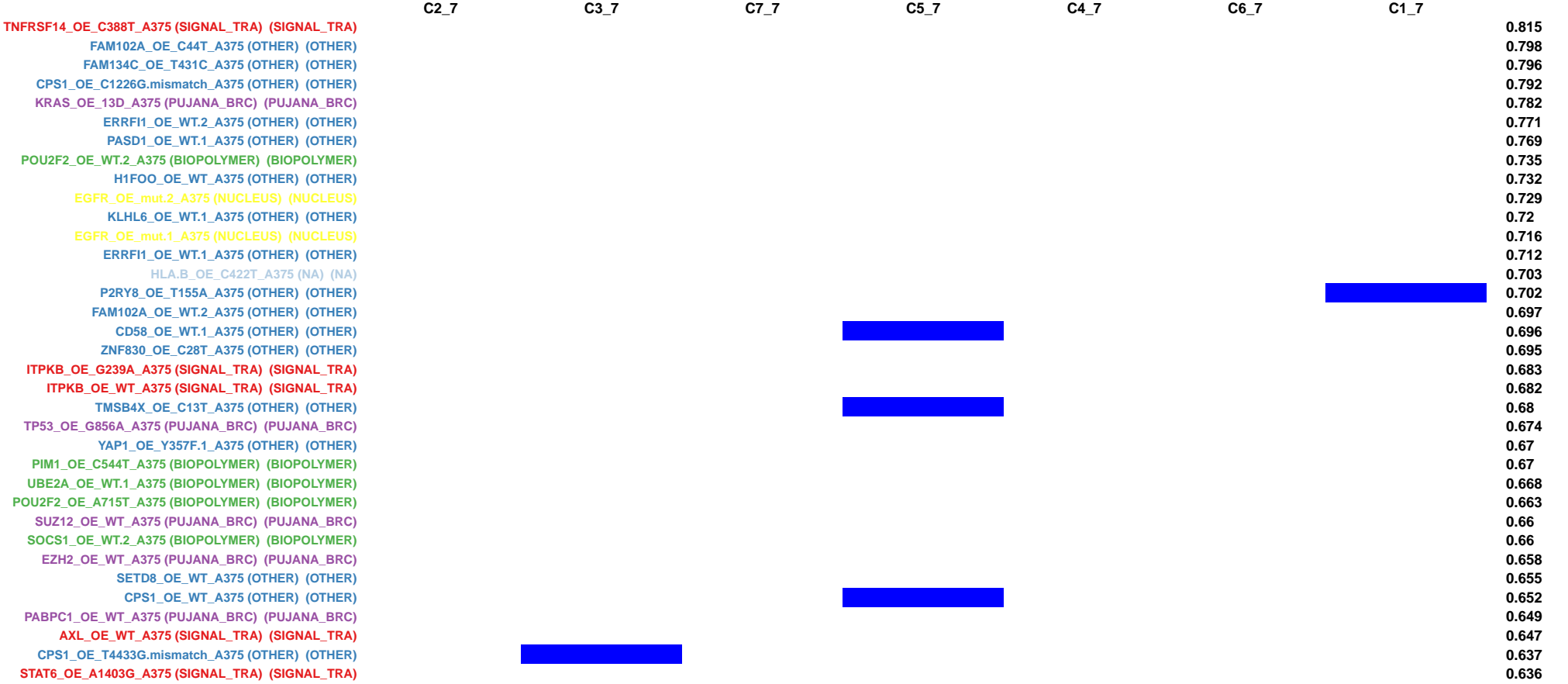


P2RY8\_OE\_G567C\_A375 (OTHER)

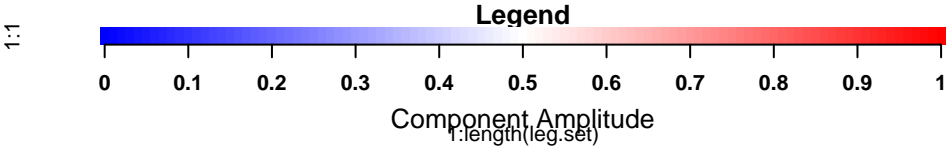
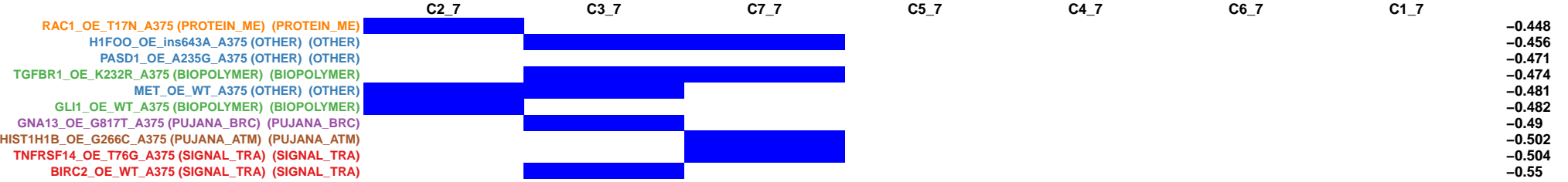
P2RY8\_OE\_G567C\_A375 (OTHER)



HIGHEST IC ASSOCIATIONS

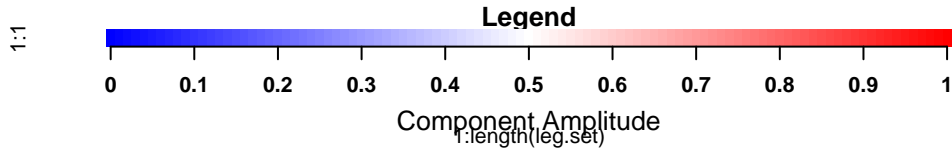
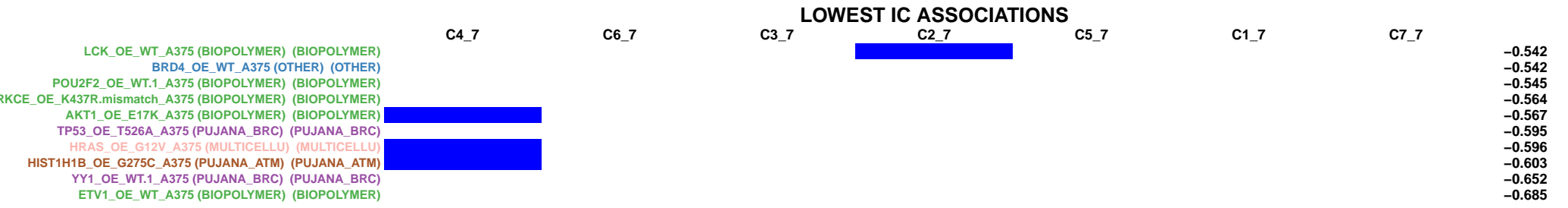
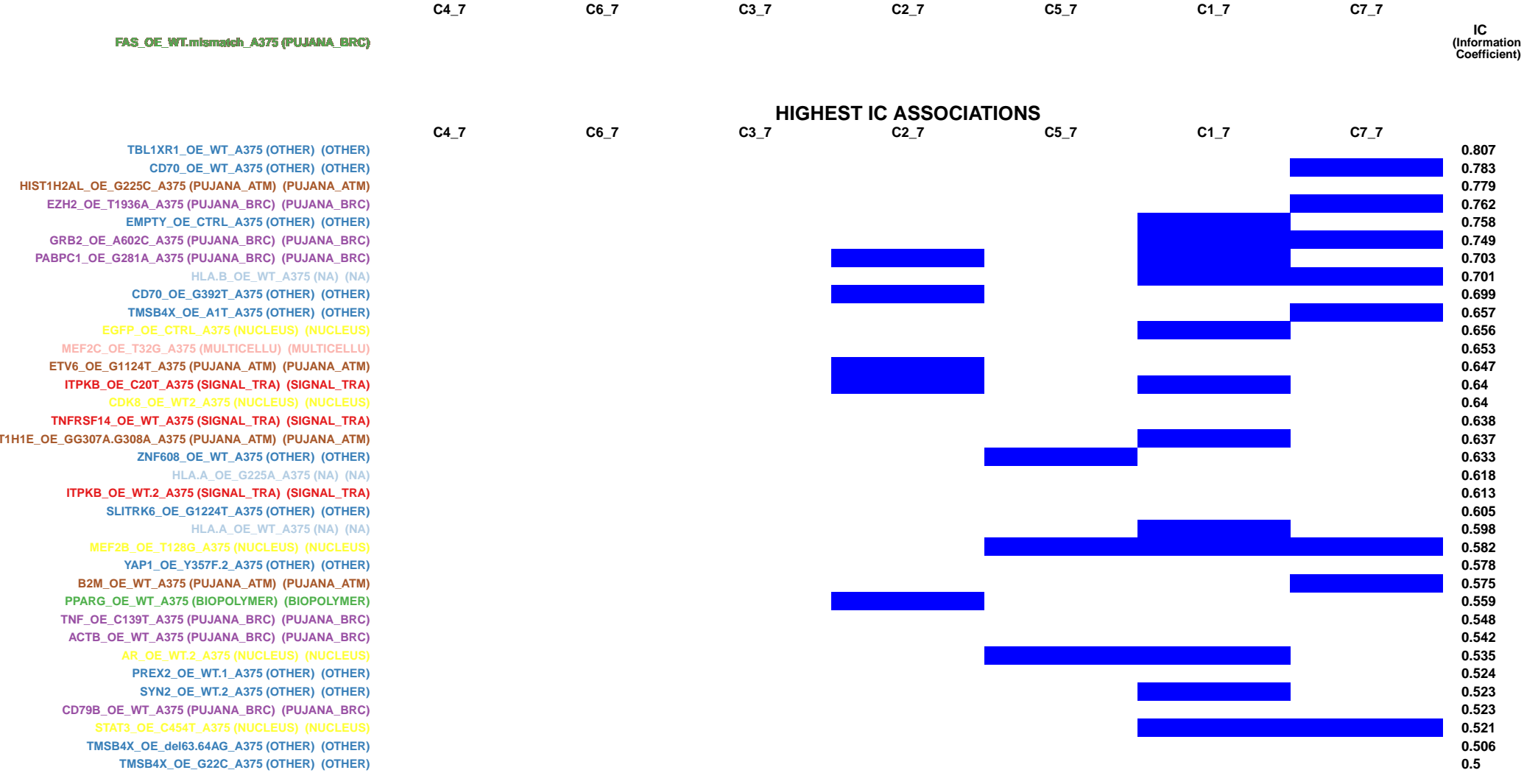


LOWEST IC ASSOCIATIONS

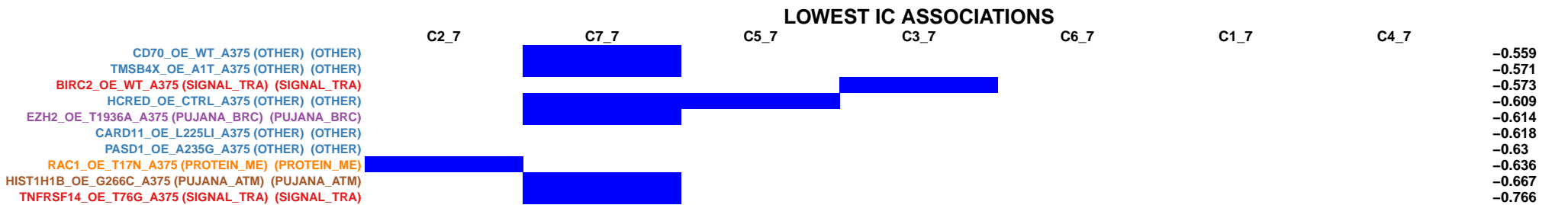




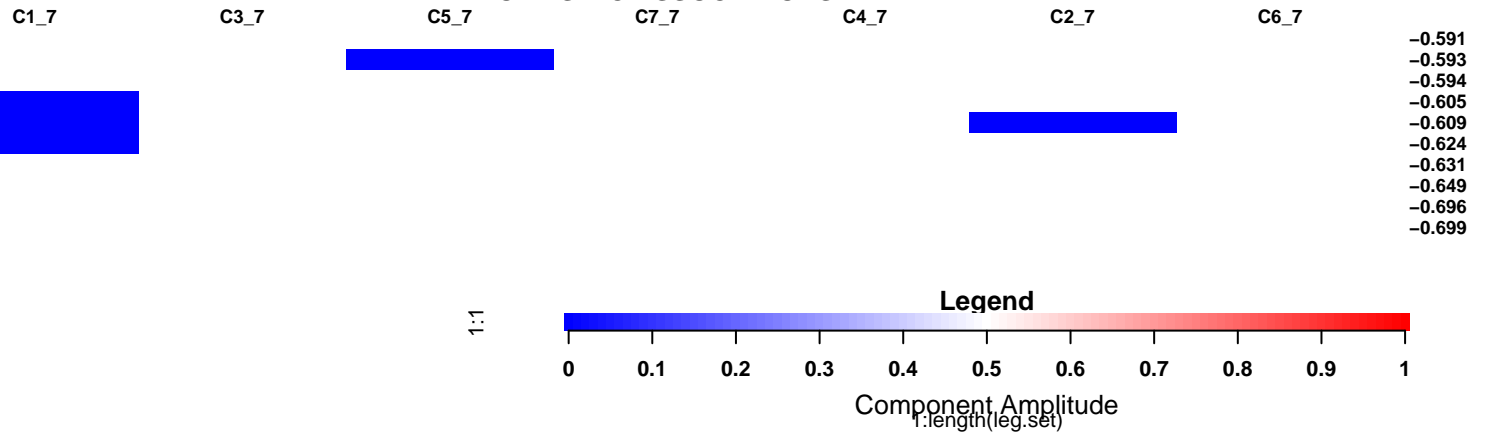
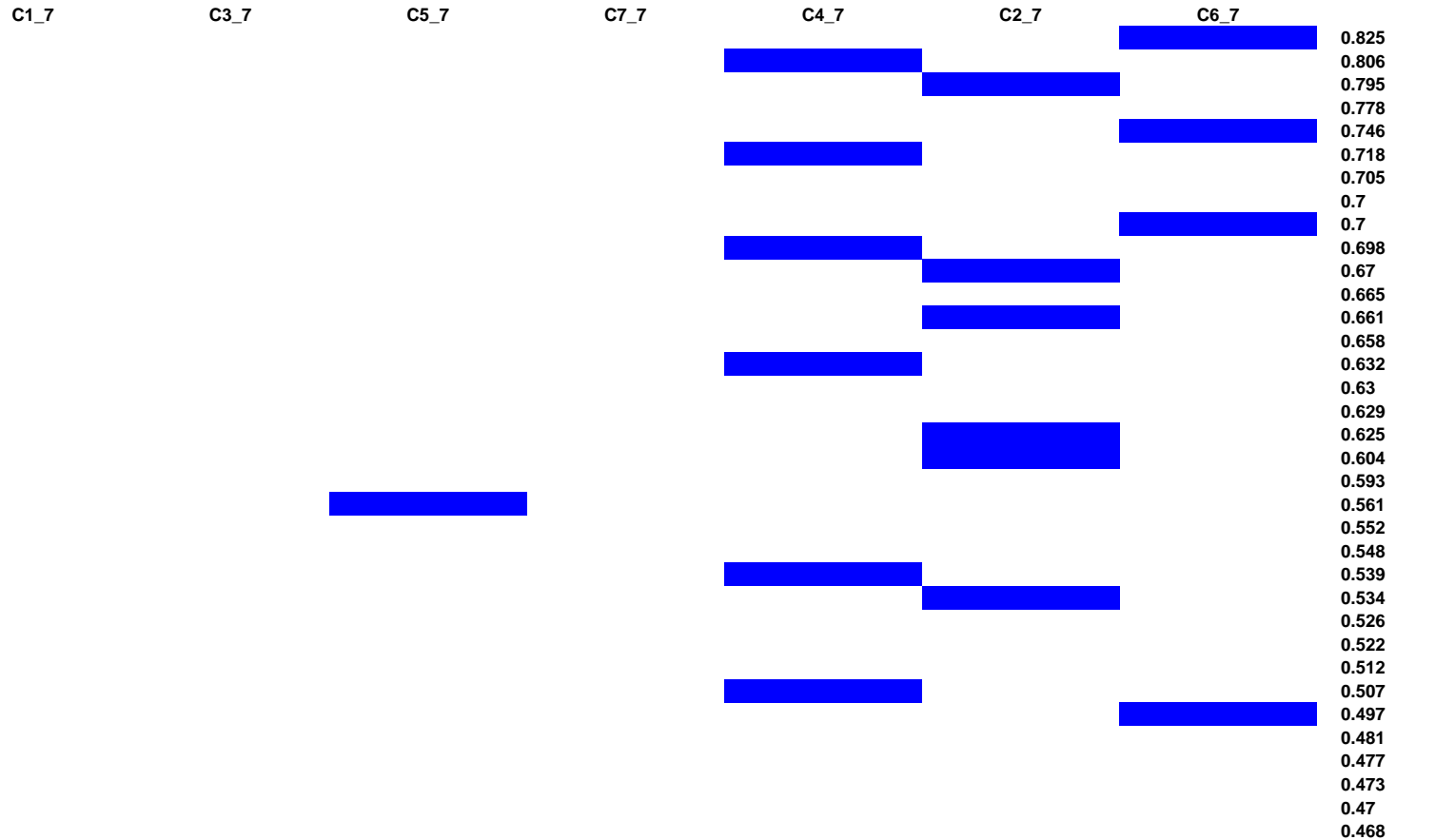
FAS\_OE\_WT.mismatch\_A375 (PUJANA\_BRC)



C2_7	C7_7	C5_7	C3_7	C6_7	C1_7	C4_7	IC (Information Coefficient)
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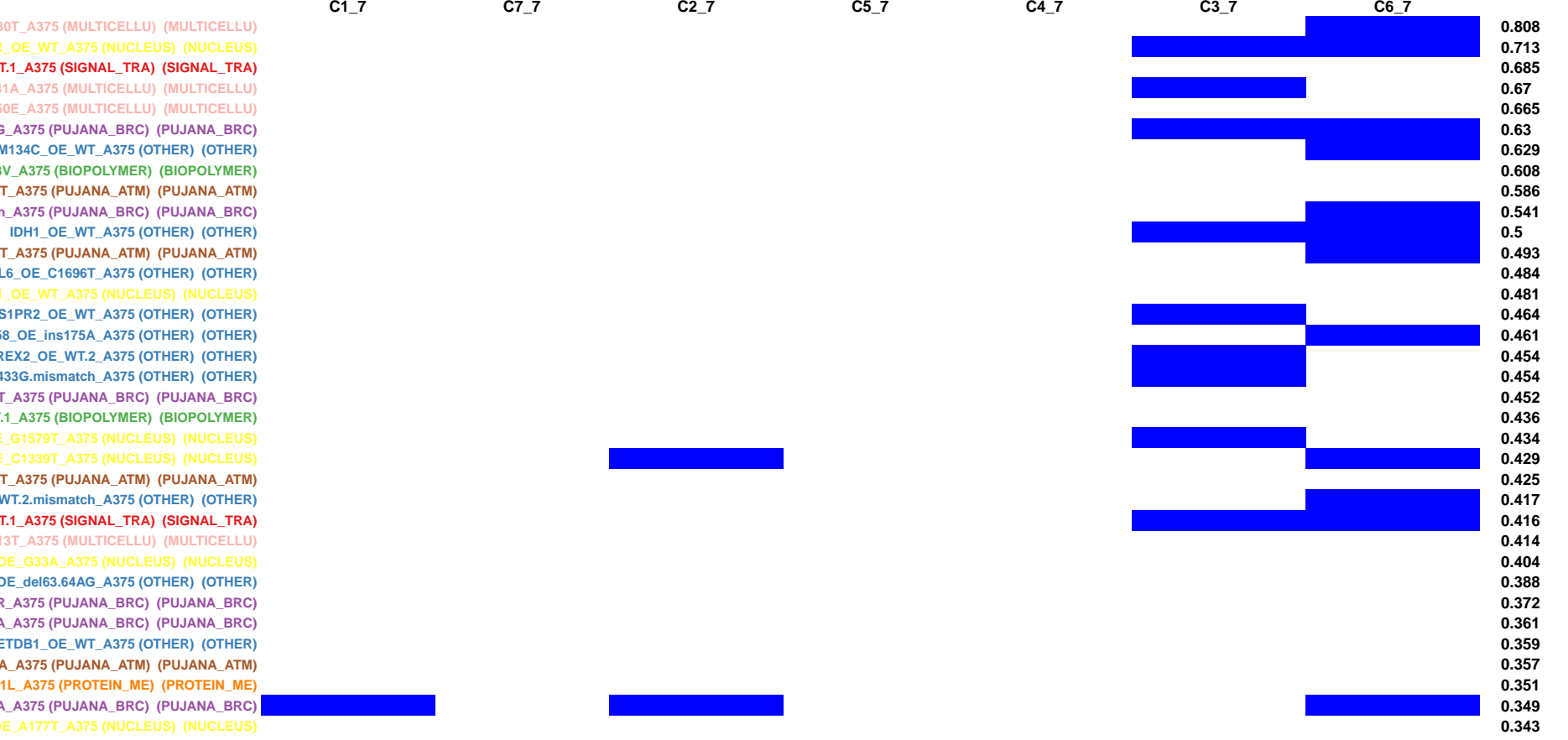
C1_7	C3_7	C5_7	C7_7	C4_7	C2_7	C6_7	IC (Information Coefficient)
0.98	0.95	0.92	0.89	0.86	0.83	0.80	0.77
0.97	0.94	0.91	0.88	0.85	0.82	0.79	0.76
0.96	0.93	0.90	0.87	0.84	0.81	0.78	0.75
0.95	0.92	0.89	0.86	0.83	0.80	0.77	0.74
0.94	0.91	0.88	0.85	0.82	0.79	0.76	0.73
0.93	0.90	0.87	0.84	0.81	0.78	0.75	0.72
0.92	0.89	0.86	0.83	0.80	0.77	0.74	0.71
0.91	0.88	0.85	0.82	0.79	0.76	0.73	0.70
0.90	0.87	0.84	0.81	0.78	0.75	0.72	0.69
0.89	0.86	0.83	0.80	0.77	0.74	0.71	0.68
0.88	0.85	0.82	0.79	0.76	0.73	0.70	0.67
0.87	0.84	0.81	0.78	0.75	0.72	0.69	0.66
0.86	0.83	0.80	0.77	0.74	0.71	0.68	0.65
0.85	0.82	0.79	0.76	0.73	0.70	0.67	0.64
0.84	0.81	0.78	0.75	0.72	0.69	0.66	0.63
0.83	0.80	0.77	0.74	0.71	0.68	0.65	0.62
0.82	0.79	0.76	0.73	0.70	0.67	0.64	0.61
0.81	0.78	0.75	0.72	0.69	0.66	0.63	0.60
0.80	0.77	0.74	0.71	0.68	0.65	0.62	0.59
0.79	0.76	0.73	0.70	0.67	0.64	0.61	0.58
0.78	0.75	0.72	0.69	0.66	0.63	0.60	0.57
0.77	0.74	0.71	0.68	0.65	0.62	0.59	0.56
0.76	0.73	0.70	0.67	0.64	0.61	0.58	0.55
0.75	0.72	0.69	0.66	0.63	0.60	0.57	0.54
0.74	0.71	0.68	0.65	0.62	0.59	0.56	0.53
0.73	0.70	0.67	0.64	0.61	0.58	0.55	0.52
0.72	0.69	0.66	0.63	0.60	0.57	0.54	0.51
0.71	0.68	0.65	0.62	0.59	0.56	0.53	0.50
0.70	0.67	0.64	0.61	0.58	0.55	0.52	0.49
0.69	0.66	0.63	0.60	0.57	0.54	0.51	0.48
0.68	0.65	0.62	0.59	0.56	0.53	0.50	0.47
0.67	0.64	0.61	0.58	0.55	0.52	0.49	0.46
0.66	0.63	0.60	0.57	0.54	0.51	0.48	0.45
0.65	0.62	0.59	0.56	0.53	0.50	0.47	0.44
0.64	0.61	0.58	0.55	0.52	0.49	0.46	0.43
0.63	0.60	0.57	0.54	0.51	0.48	0.45	0.42
0.62	0.59	0.56	0.53	0.50	0.47	0.44	0.41
0.61	0.58	0.55	0.52	0.49	0.46	0.43	0.40
0.60	0.57	0.54	0.51	0.48	0.45	0.42	0.39
0.59	0.56	0.53	0.50	0.47	0.44	0.41	0.38
0.58	0.55	0.52	0.49	0.46	0.43	0.40	0.37
0.57	0.54	0.51	0.48	0.45	0.42	0.39	0.36
0.56	0.53	0.50	0.47	0.44	0.41	0.38	0.35
0.55	0.52	0.49	0.46	0.43	0.40	0.37	0.34
0.54	0.51	0.48	0.45	0.42	0.39	0.36	0.33
0.53	0.50	0.47	0.44	0.41	0.38	0.35	0.32
0.52	0.49	0.46	0.43	0.40	0.37	0.34	0.31
0.51	0.48	0.45	0.42	0.39	0.36	0.33	0.30
0.50	0.47	0.44	0.41	0.38	0.35	0.32	0.29
0.49	0.46	0.43	0.40	0.37	0.34	0.31	0.28
0.48	0.45	0.42	0.39	0.36	0.33	0.30	0.27
0.47	0.44	0.41</					



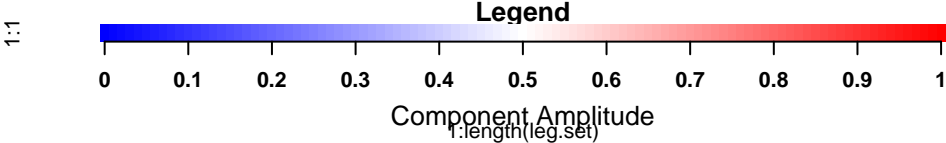
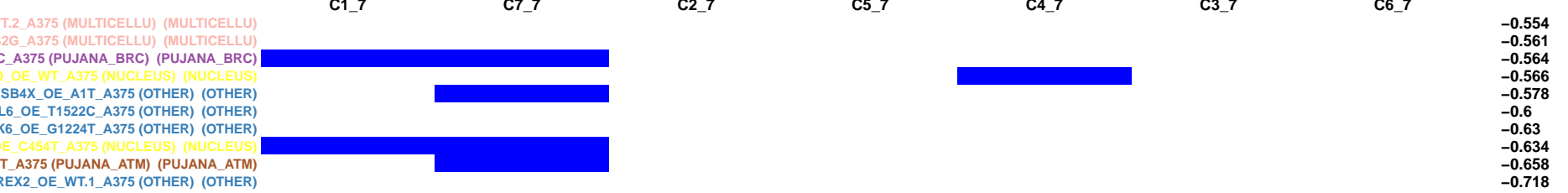
LUCIFERASE\_OE\_CTRL\_A375 (OTHER)



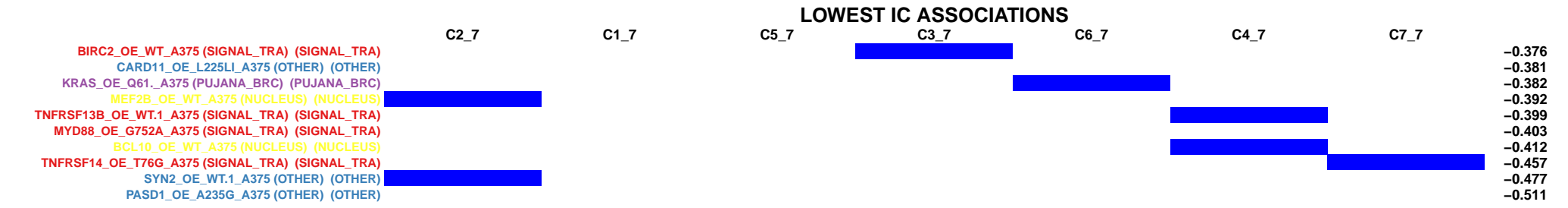
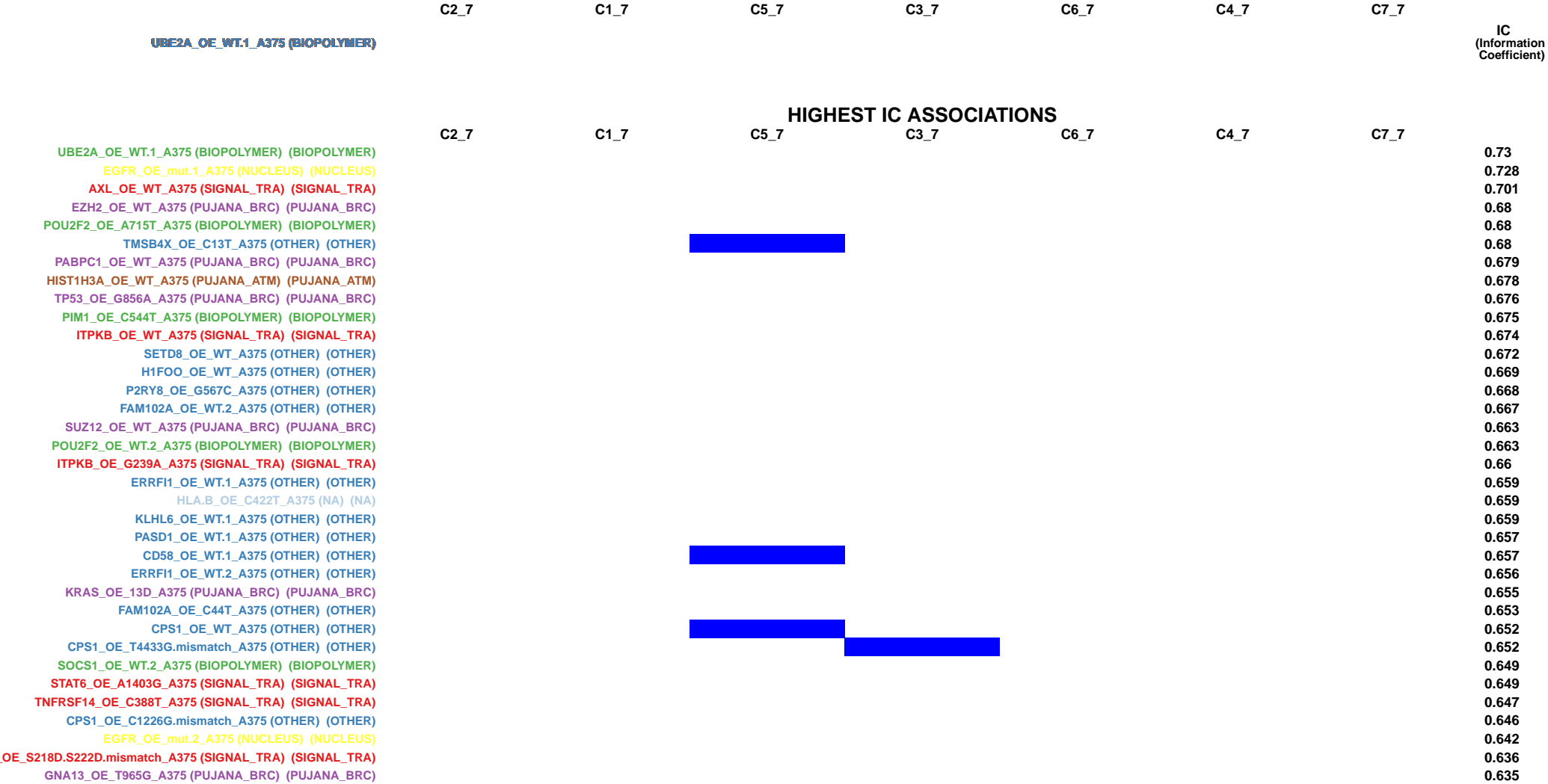
HIGHEST IC ASSOCIATIONS



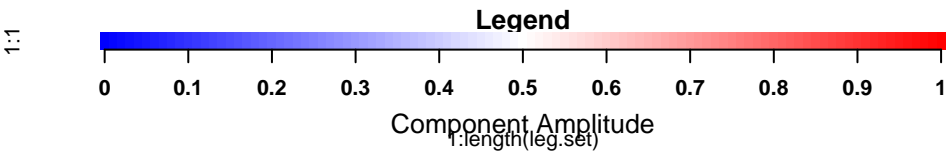
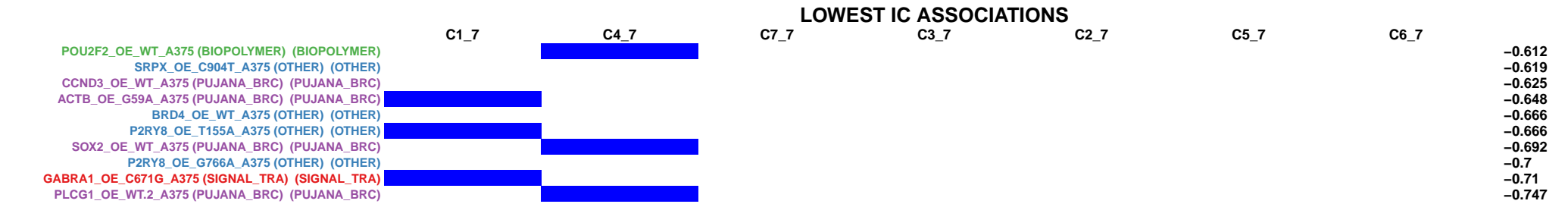
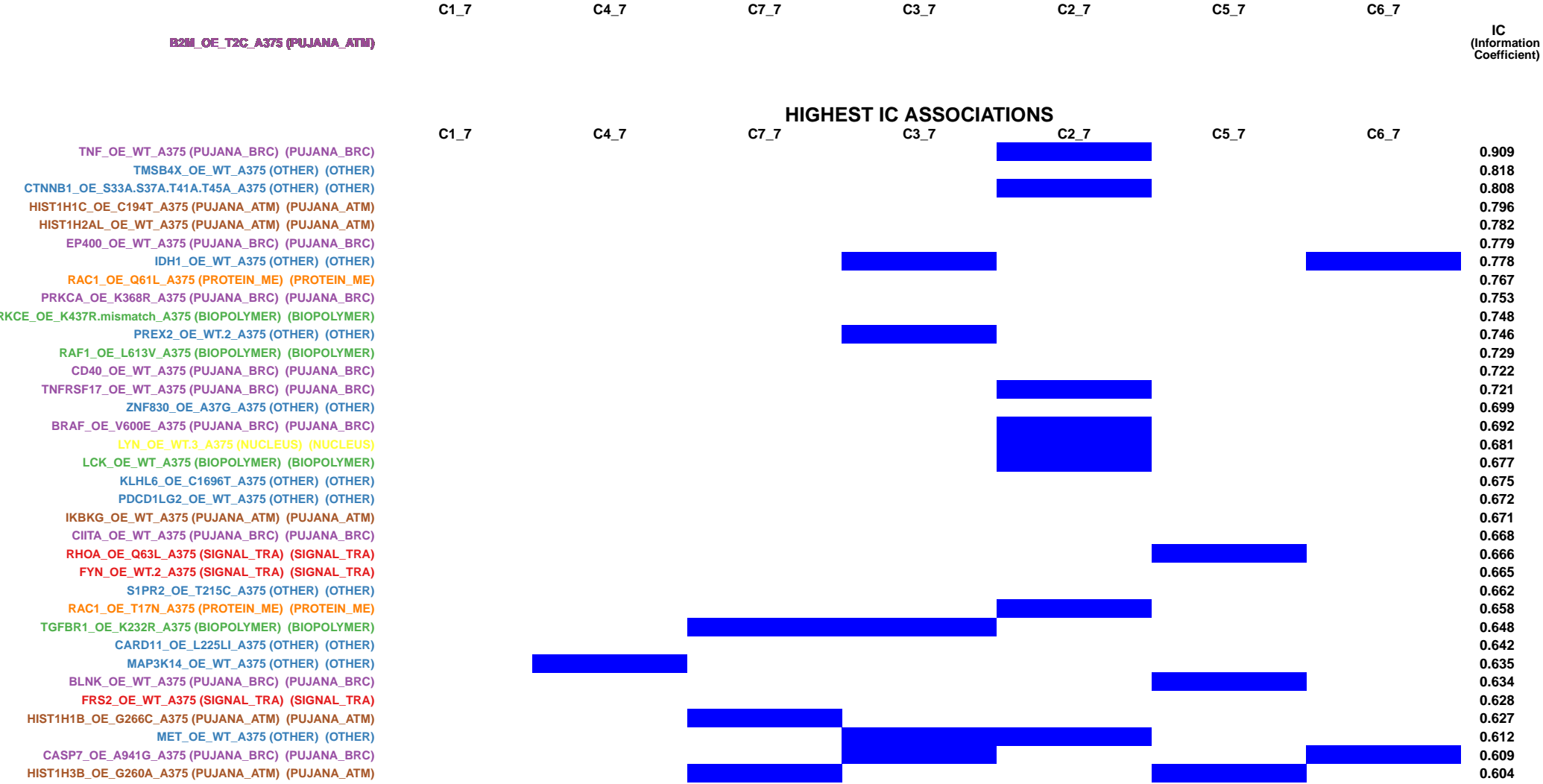
LOWEST IC ASSOCIATIONS



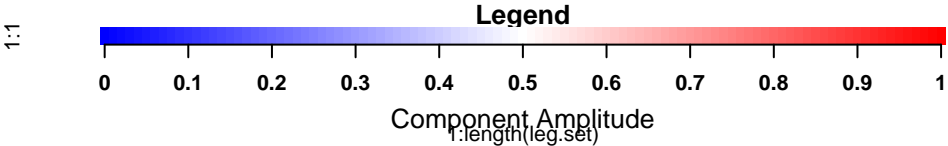
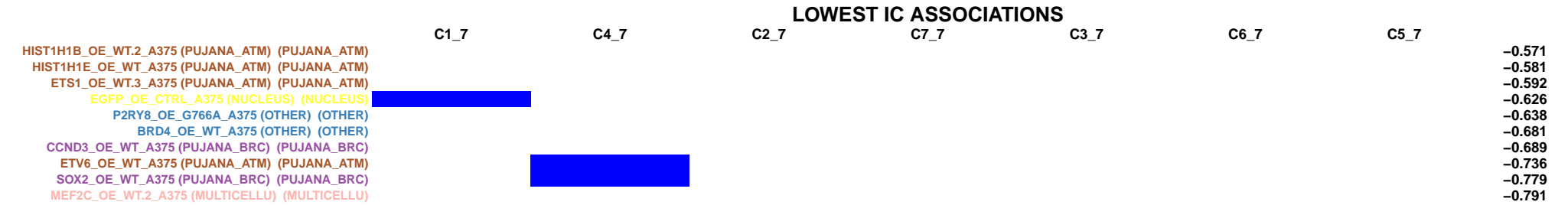
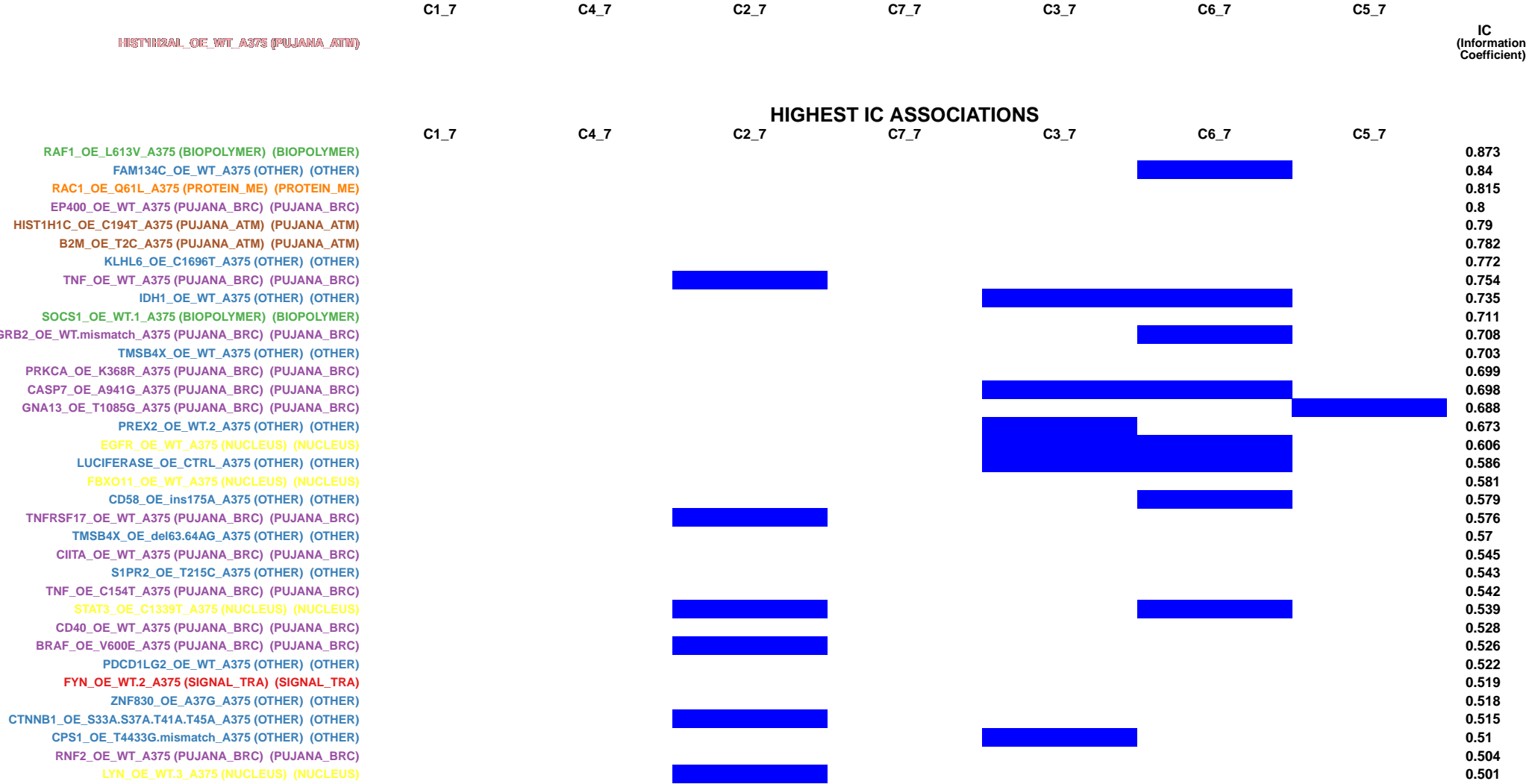
UBE2A\_OE\_WT.1\_A375 (BIOPOLYMER)



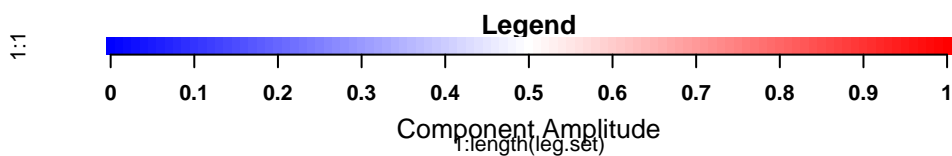
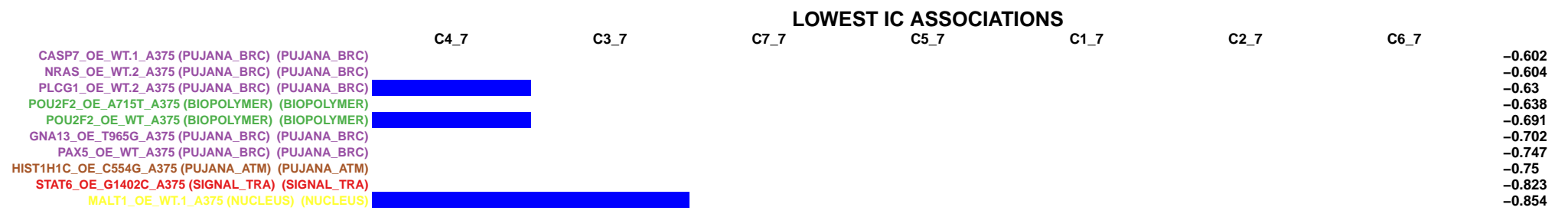
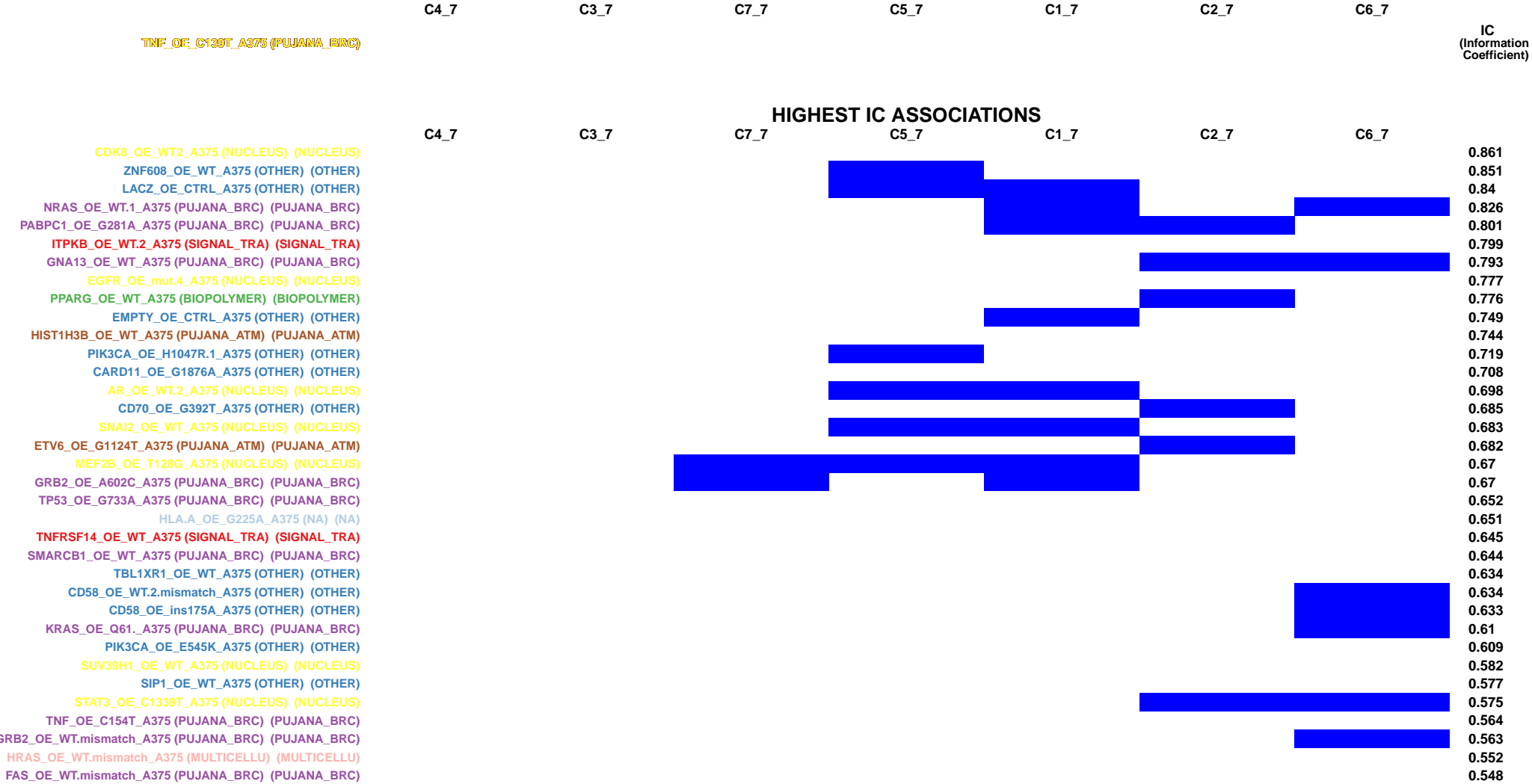
B2M\_OE\_T2C\_A375 (PUJANA\_ATM)



HIST1H2AL\_OE\_WT\_A375 (PUJANA\_ATM)

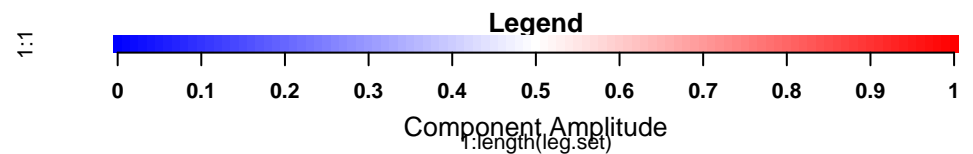
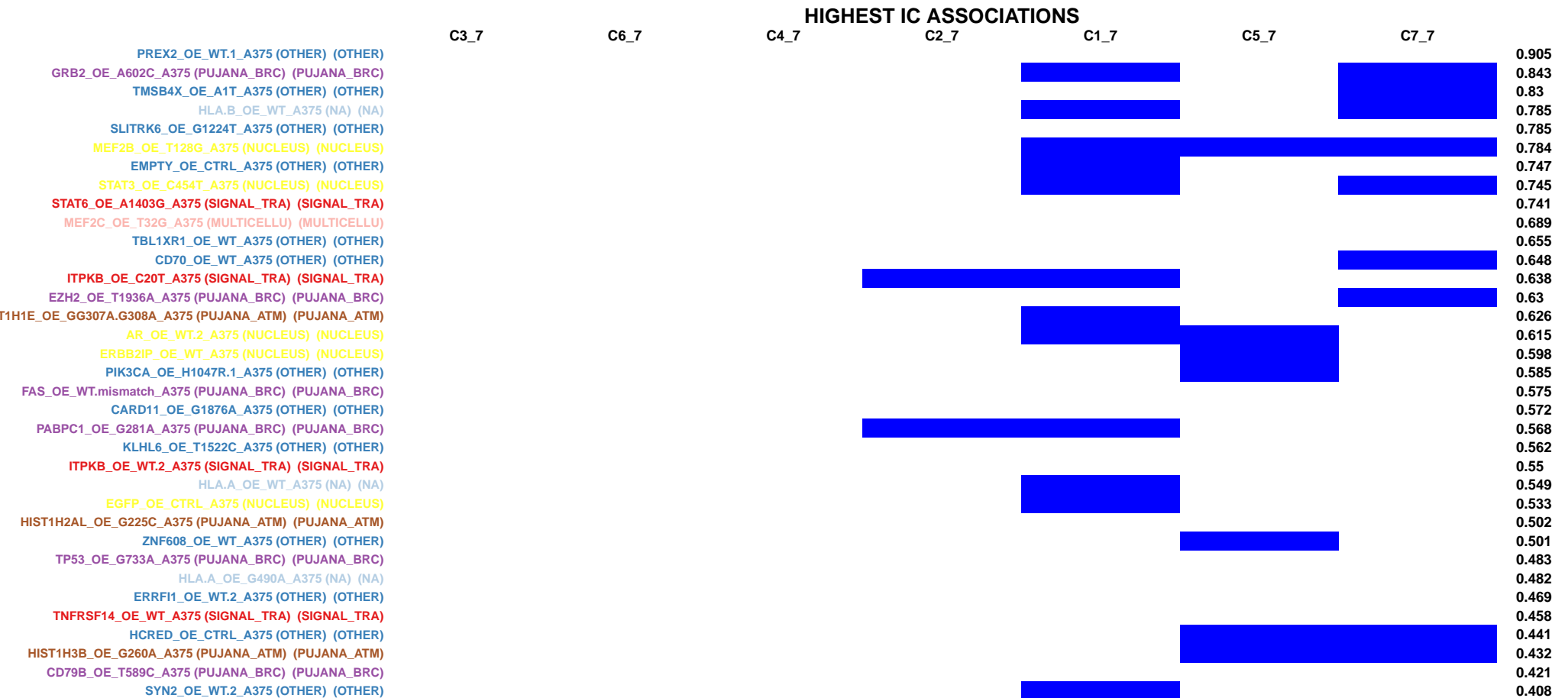


TNF\_OE\_C139T\_A375 (PUJANA\_BRC)

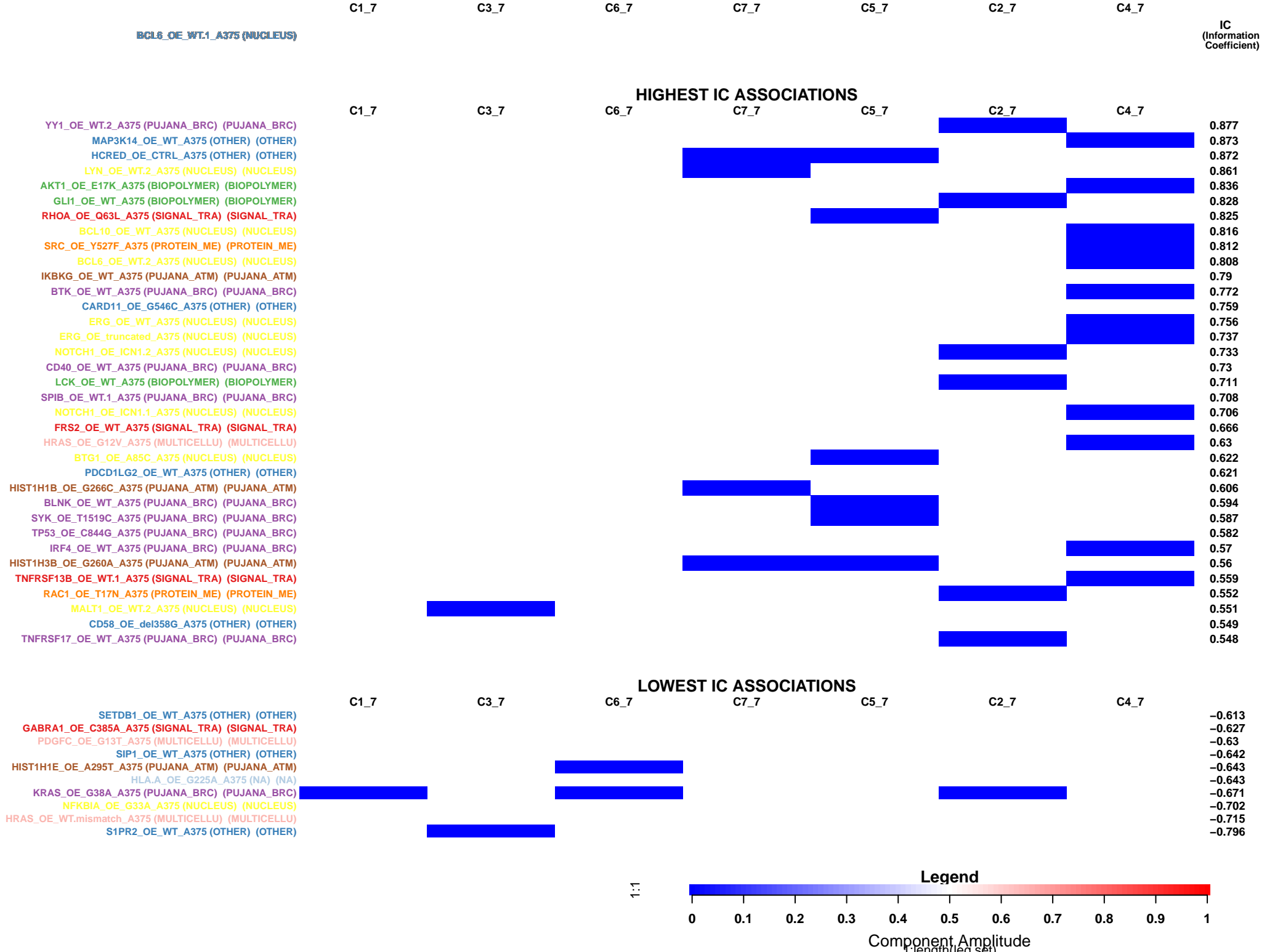




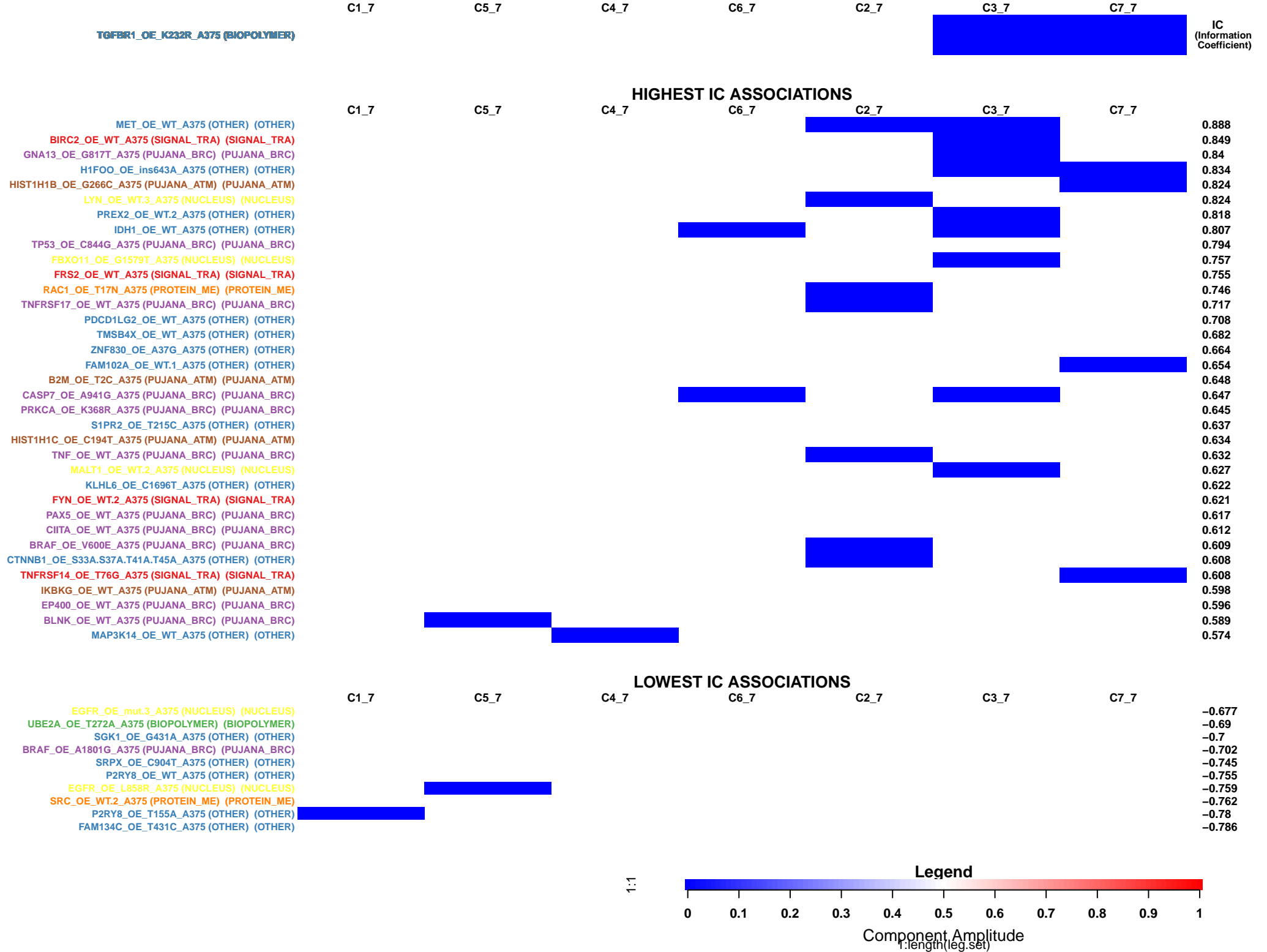
C3_7	C6_7	C4_7	C2_7	C1_7	C5_7	C7_7	IC (Information Coefficient)
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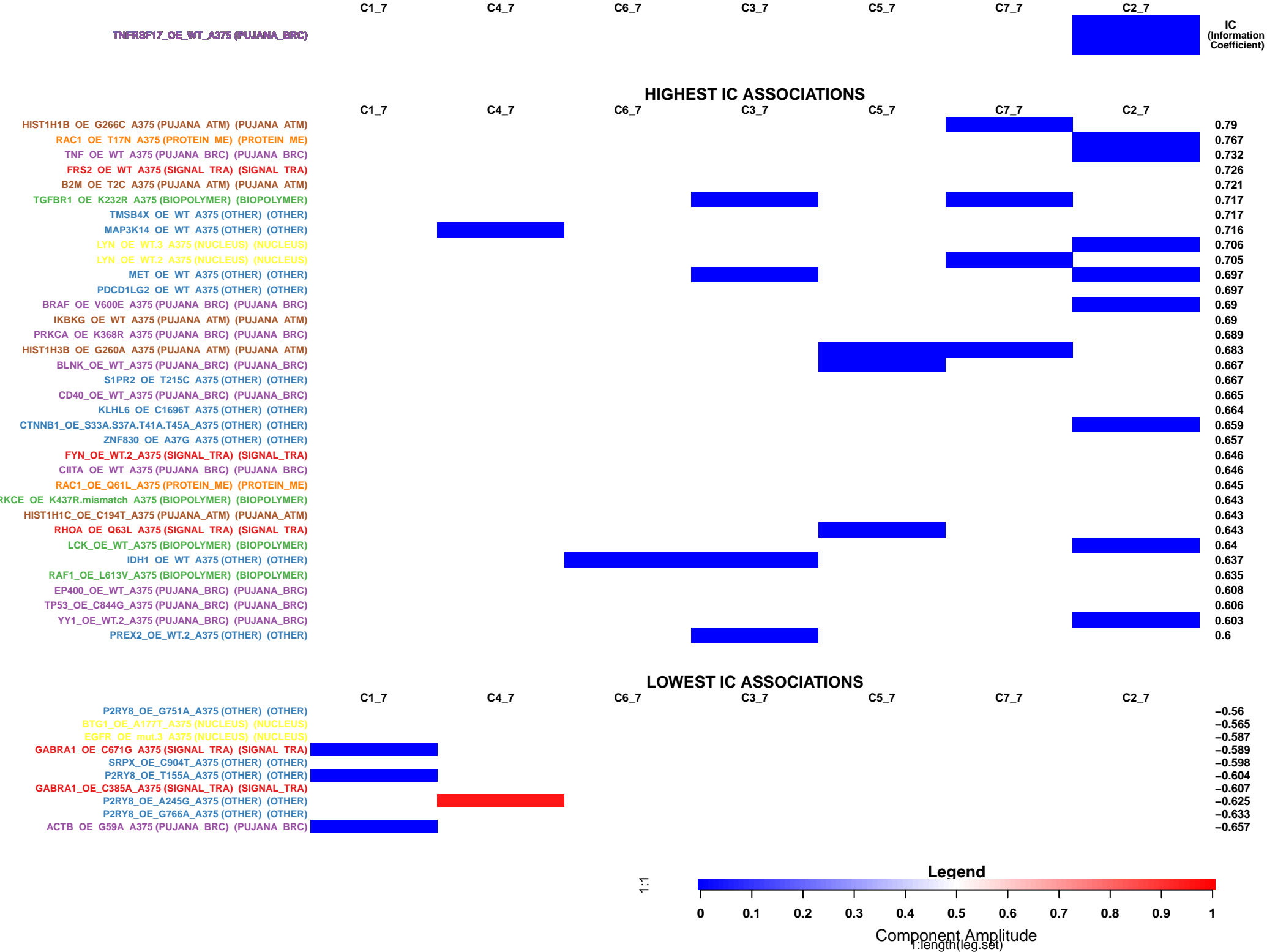
BCL6\_OE\_WT.1\_A375 (NUCLEUS)



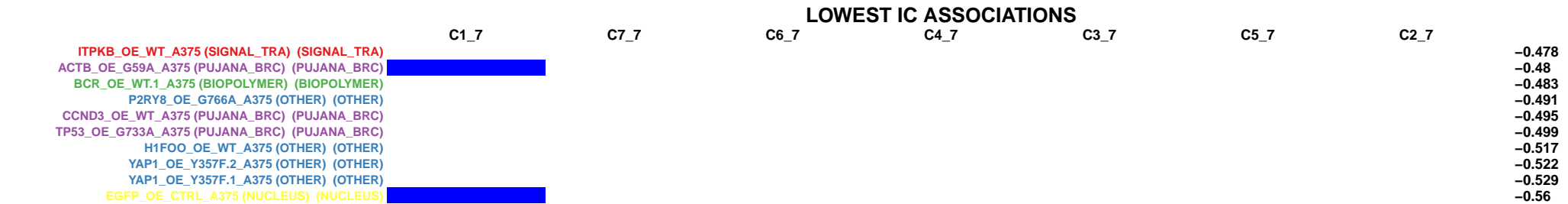
TGFBR1\_OE\_K232R\_A375 (BIOPOLYMER)



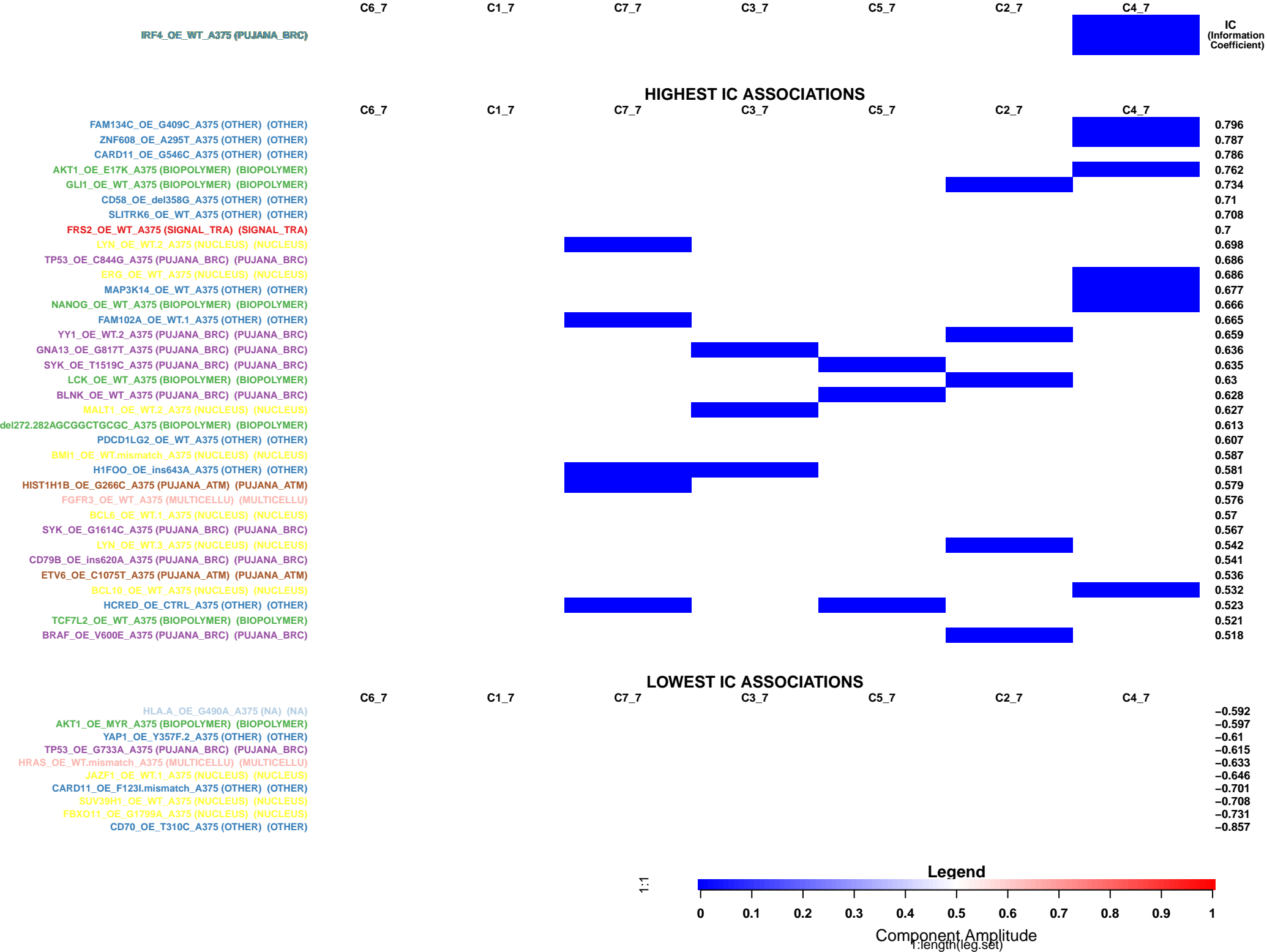
TNFRSF17\_OE\_WT\_A375 (PUJANA\_BRC)



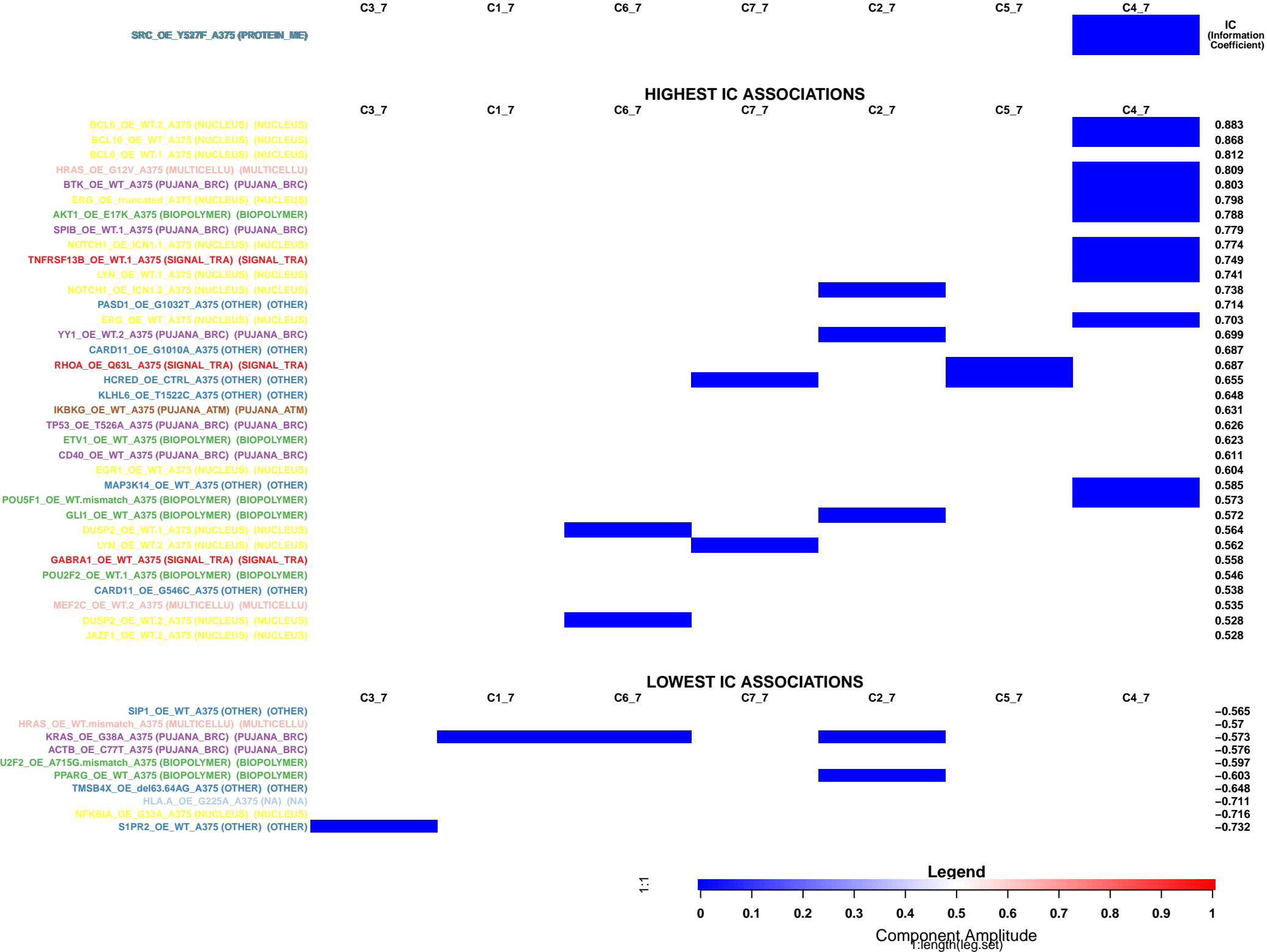
C1_7	C7_7	C6_7	C4_7	C3_7	C5_7	C2_7	IC (Information Coefficient)
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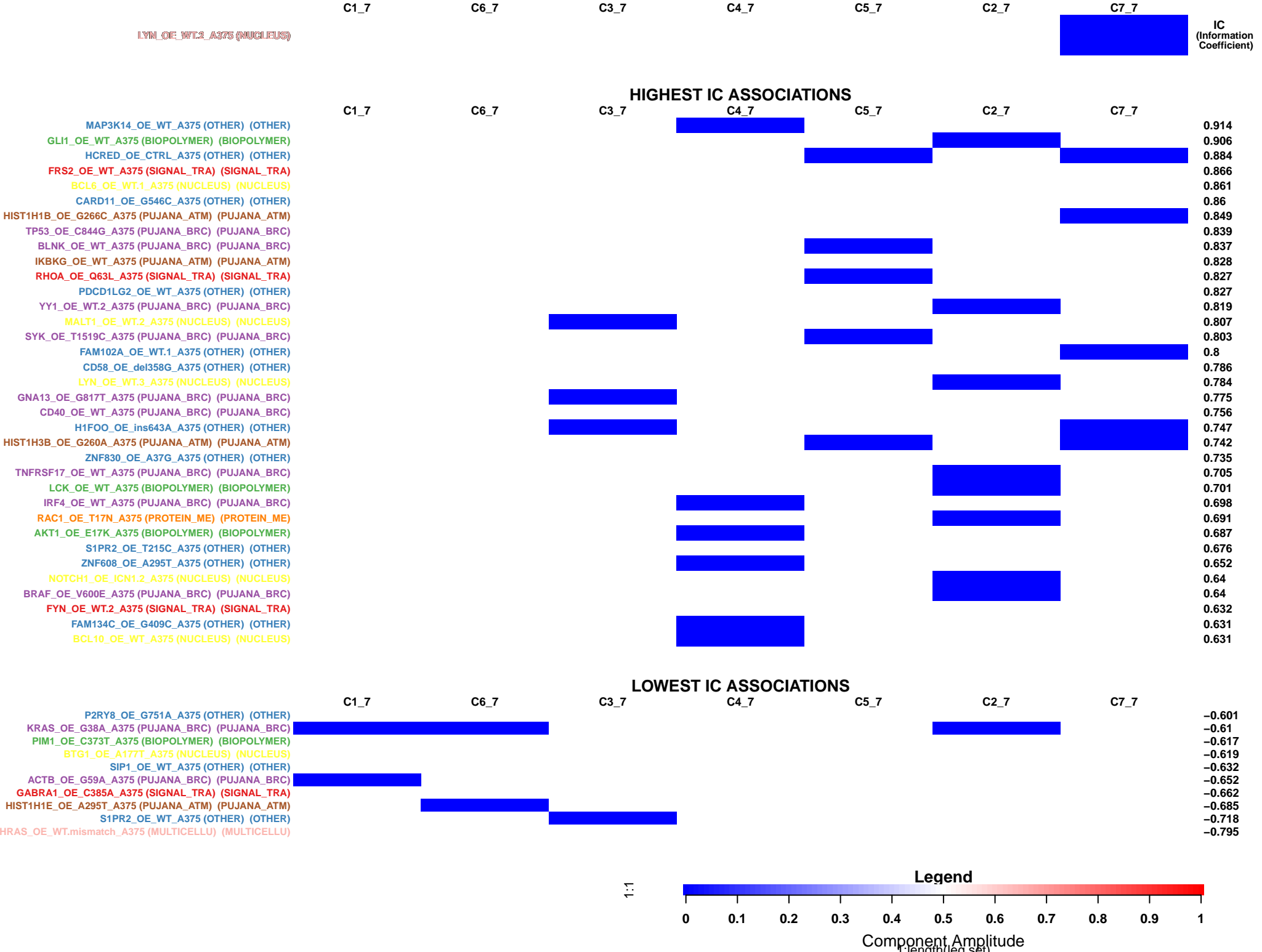
IRF4\_OE\_WT\_A375 (PUJANA\_BRC)



SRC\_OE\_Y527F\_A375 (PROTEIN\_ME)

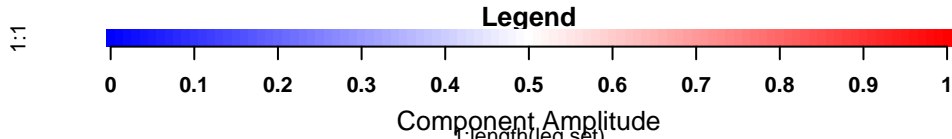
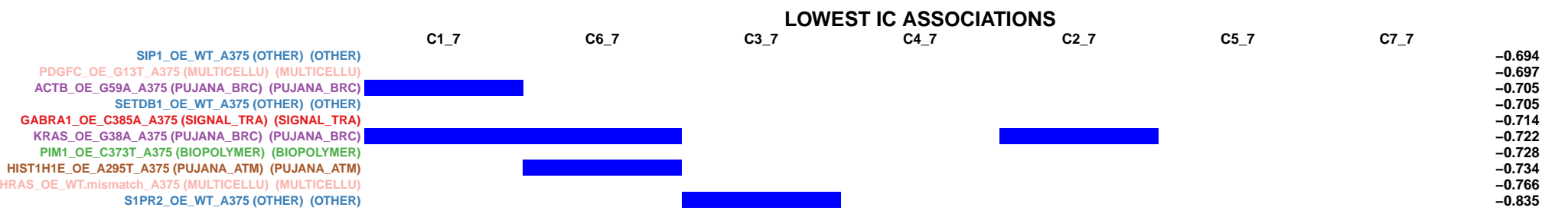
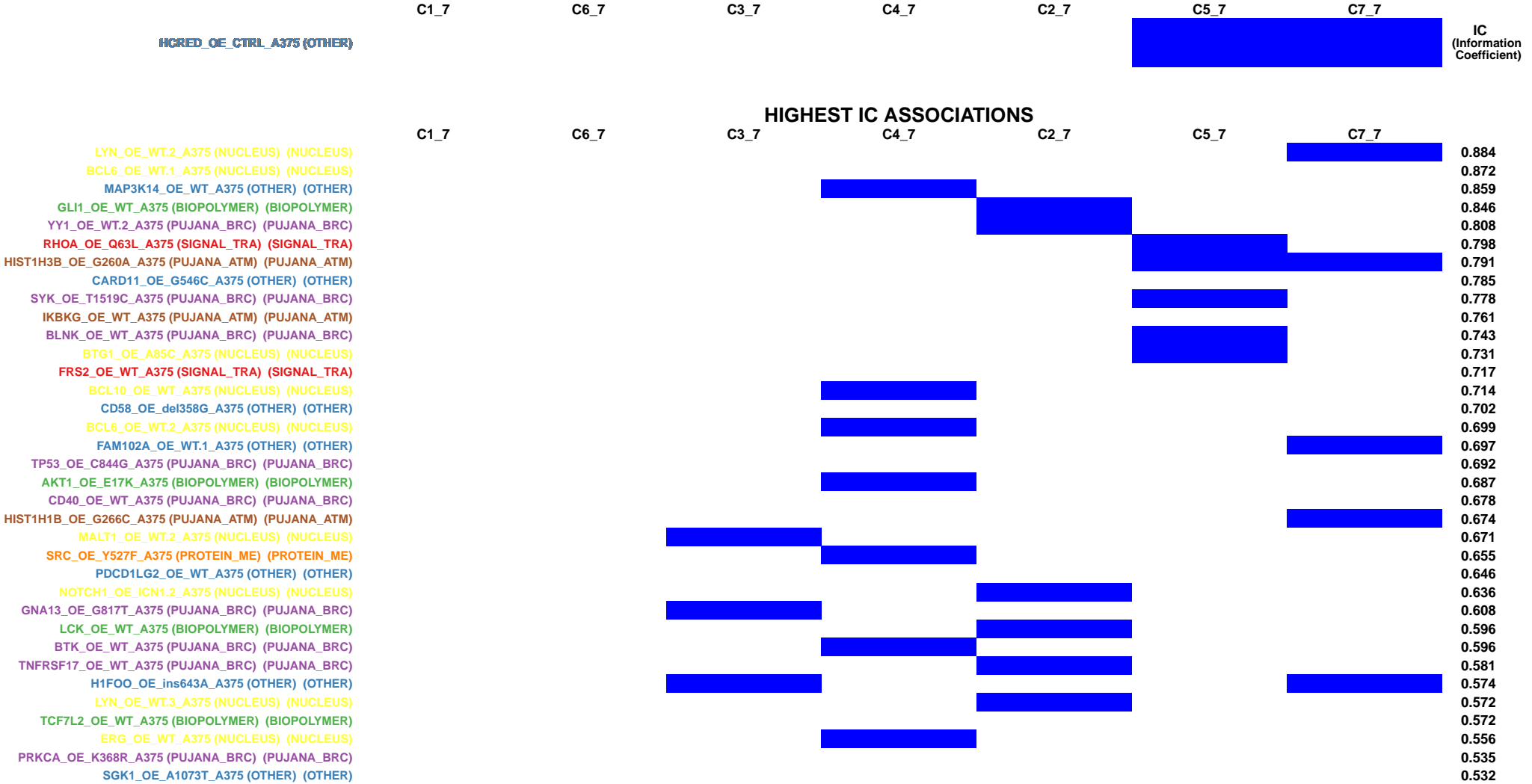


LYN\_OE\_WT.2\_A375 (NUCLEUS)

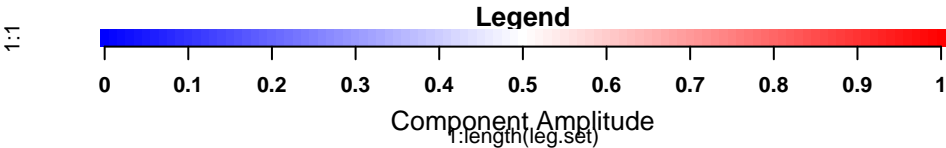
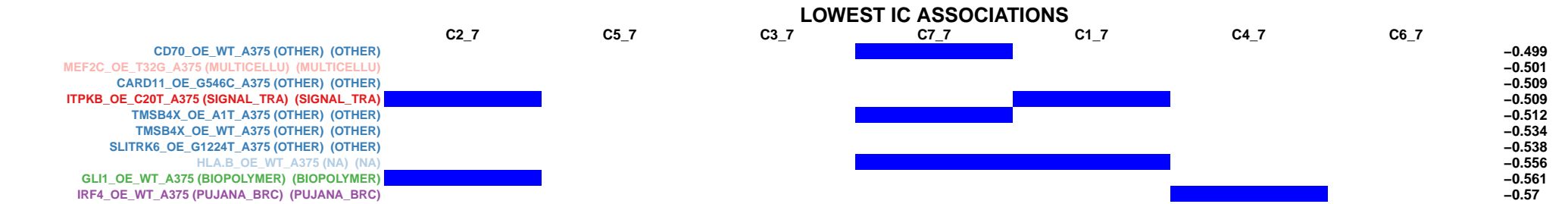
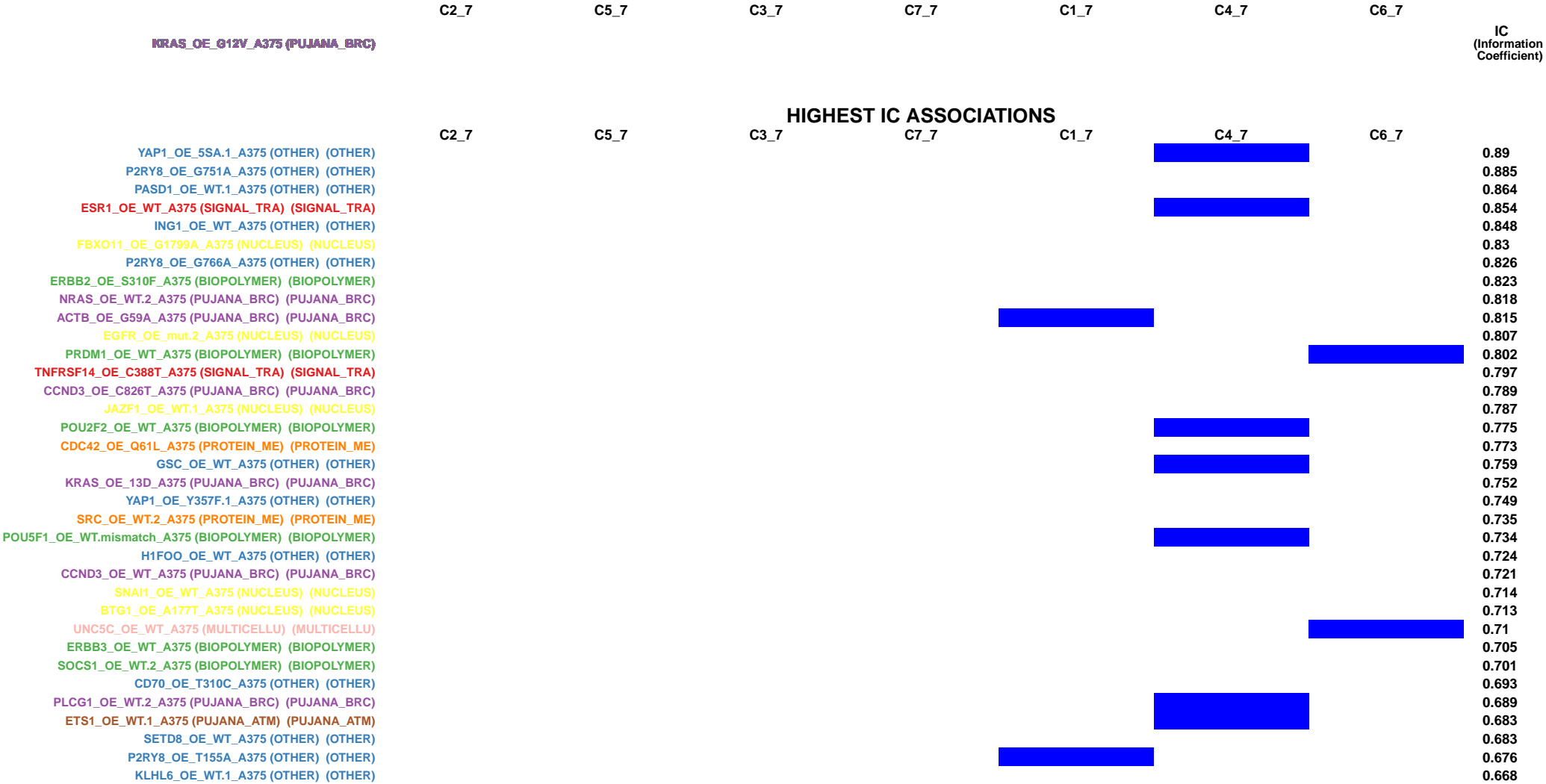




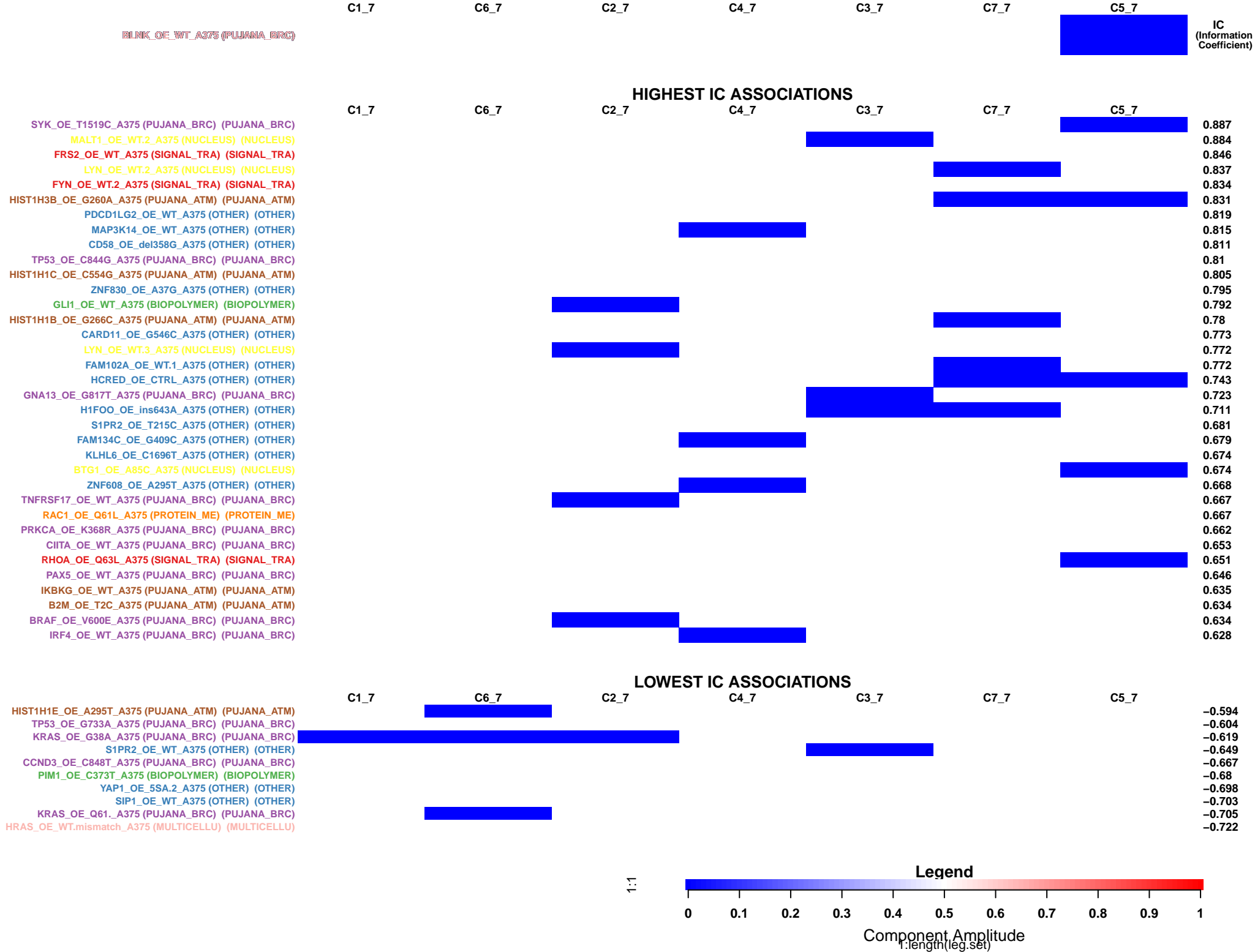
HCRED\_OE\_CTRL\_A375 (OTHER)



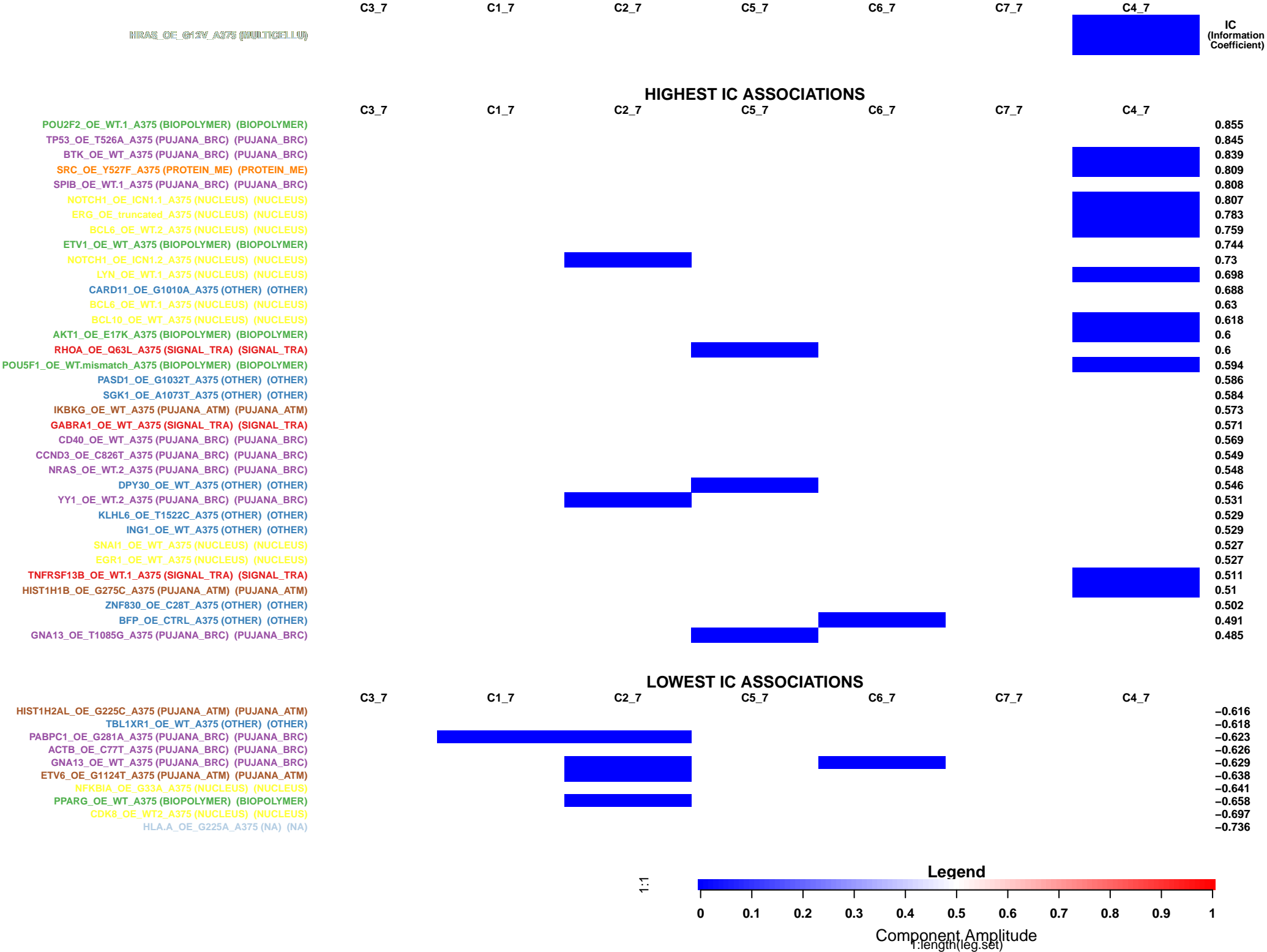
KRAS\_OE\_G12V\_A375 (PUJANA\_BRC)



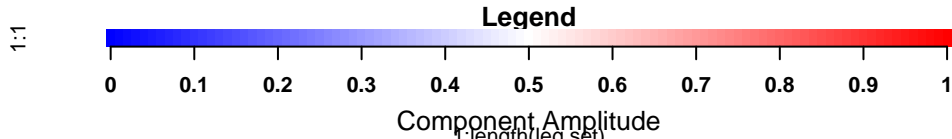
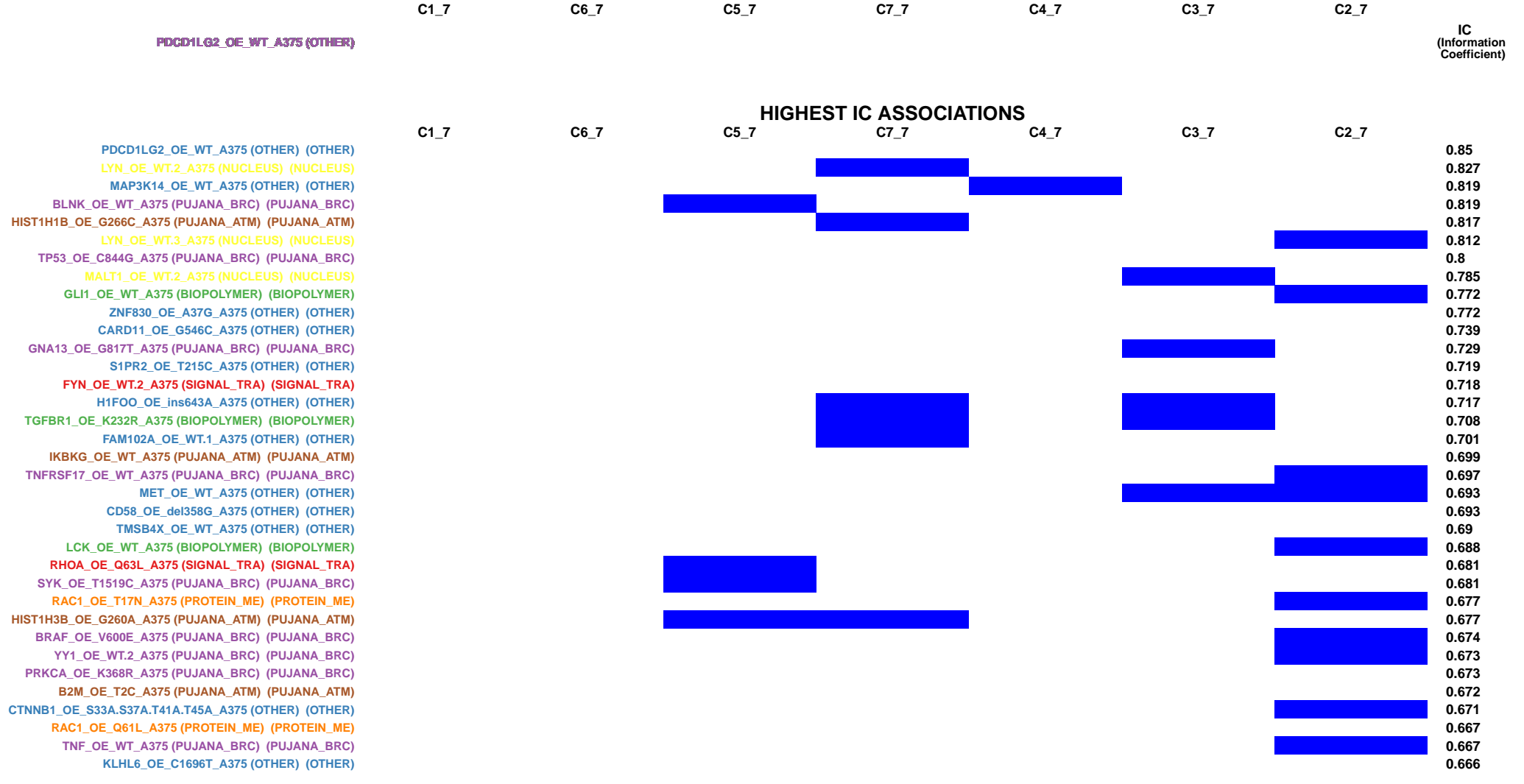
BLNK\_OE\_WT\_A375 (PUJANA\_BRC)



HRAS\_OE\_G12V\_A375 (MULTICELLU)



PDCD1LG2\_OE\_WT\_A375 (OTHER)



MAP2K1\_OE\_S218D.S222D.mismatch\_A375 (SIGNAL\_TRA)

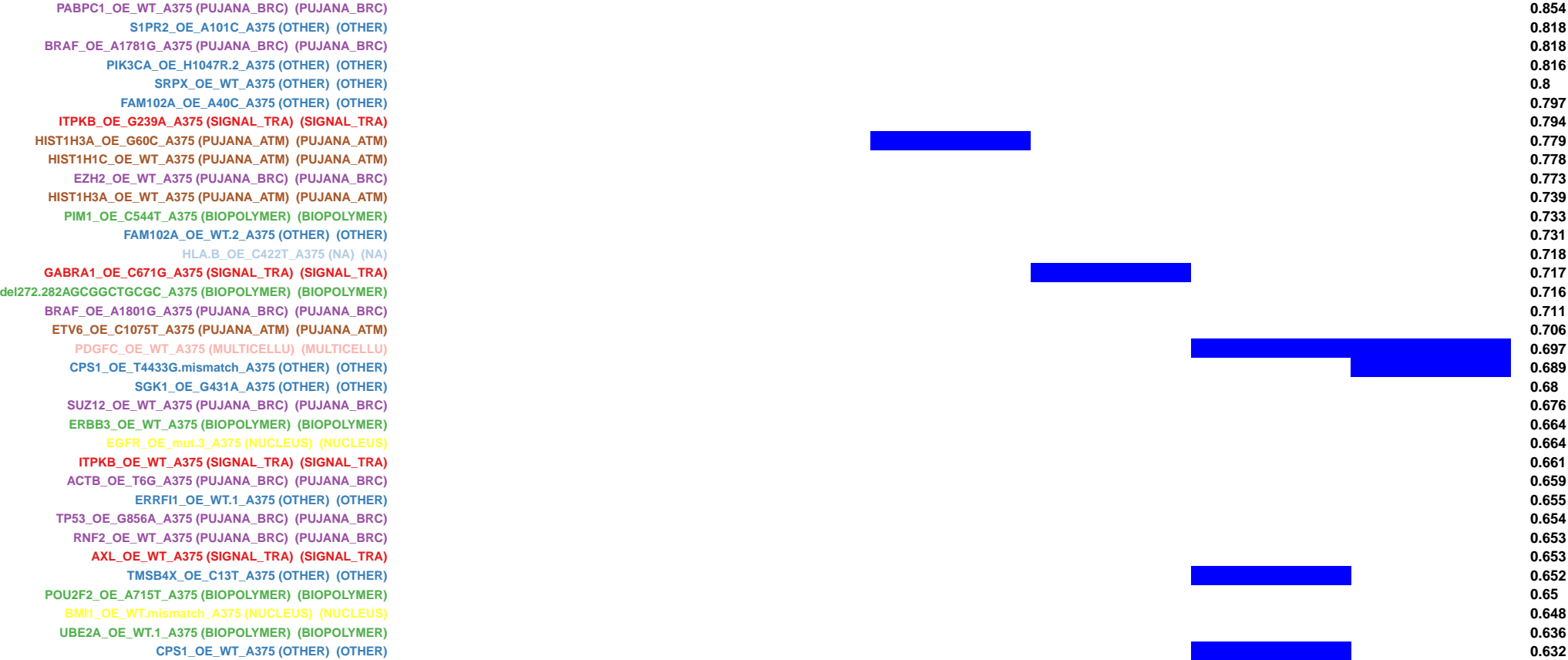
C2\_7C6\_7C7\_7C4\_7C1\_7C5\_7C3\_7

MAP2K1\_OE\_S218D.S222D.mismatch\_A375 (SIGNAL\_TRA)

IC  
(Information  
Coefficient)

HIGHEST IC ASSOCIATIONS

C2\_7C6\_7C7\_7C4\_7C1\_7C5\_7C3\_7



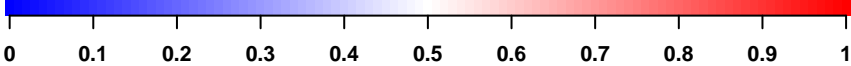
LOWEST IC ASSOCIATIONS

C2\_7C6\_7C7\_7C4\_7C1\_7C5\_7C3\_7



1:1

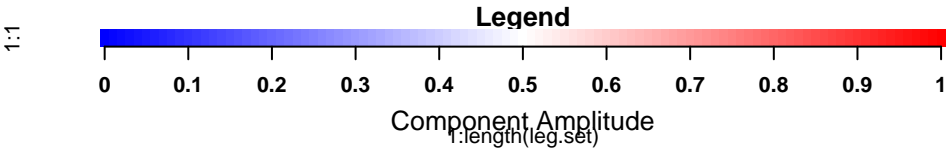
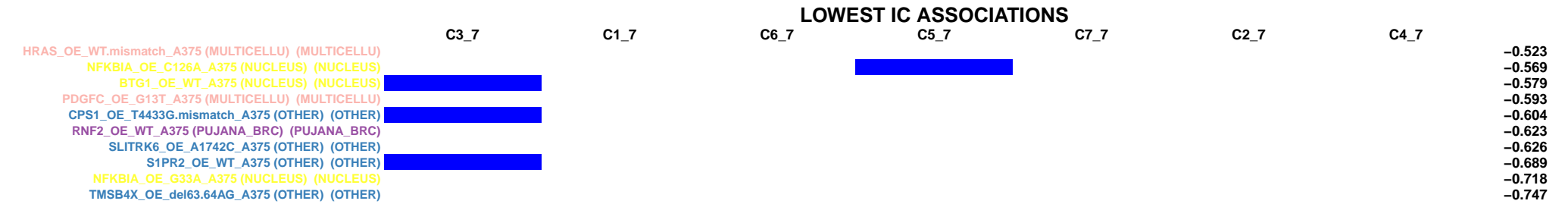
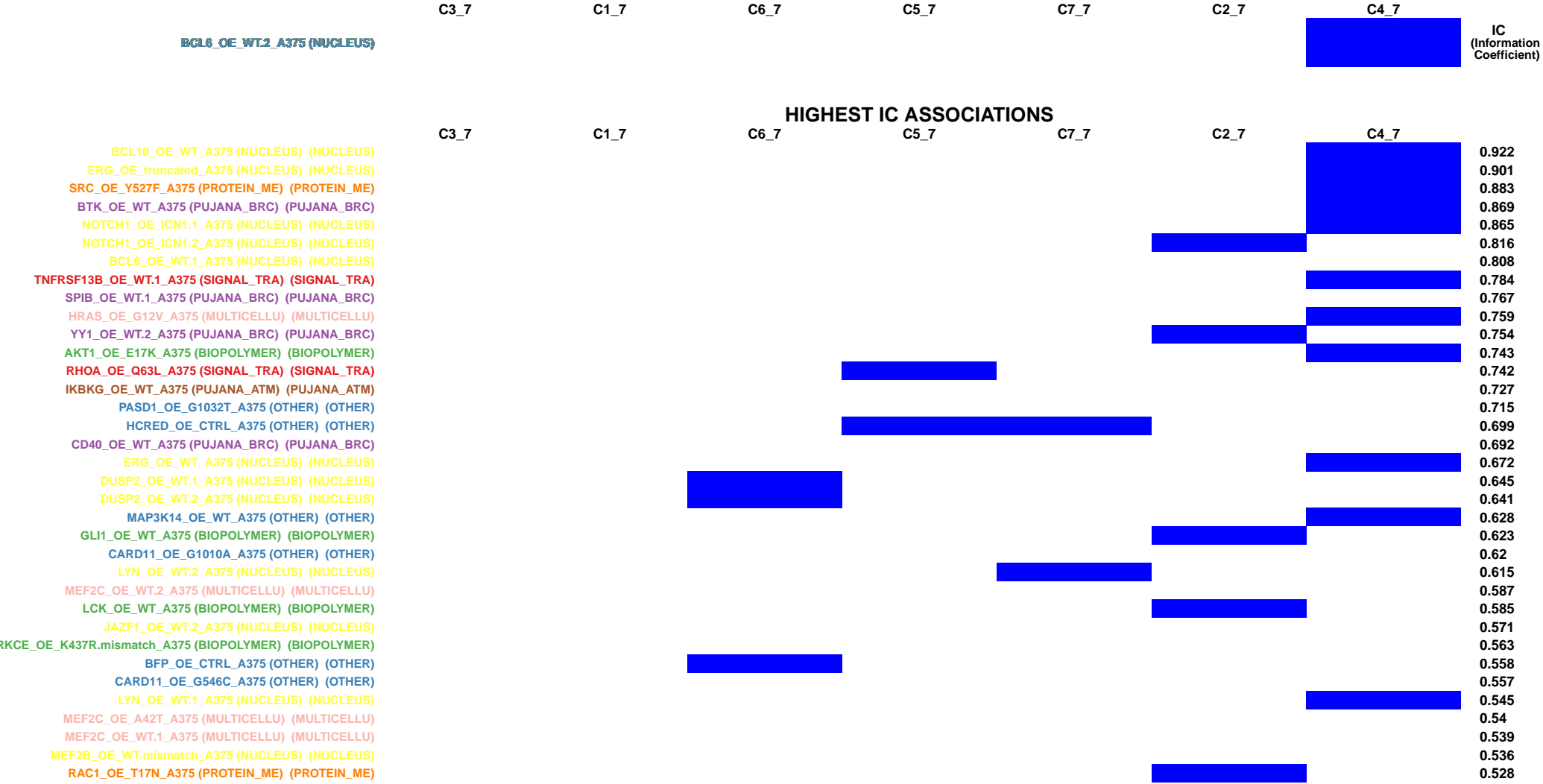
Legend



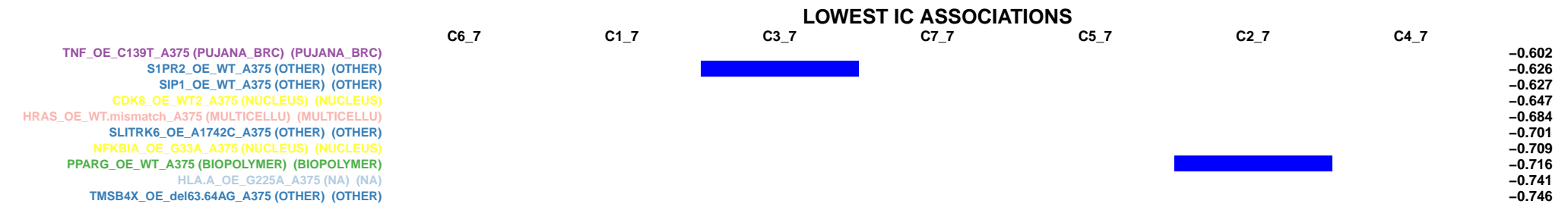
Component Amplitude

1:length(leg.set)

BCL6\_OE\_WT.2\_A375 (NUCLEUS)

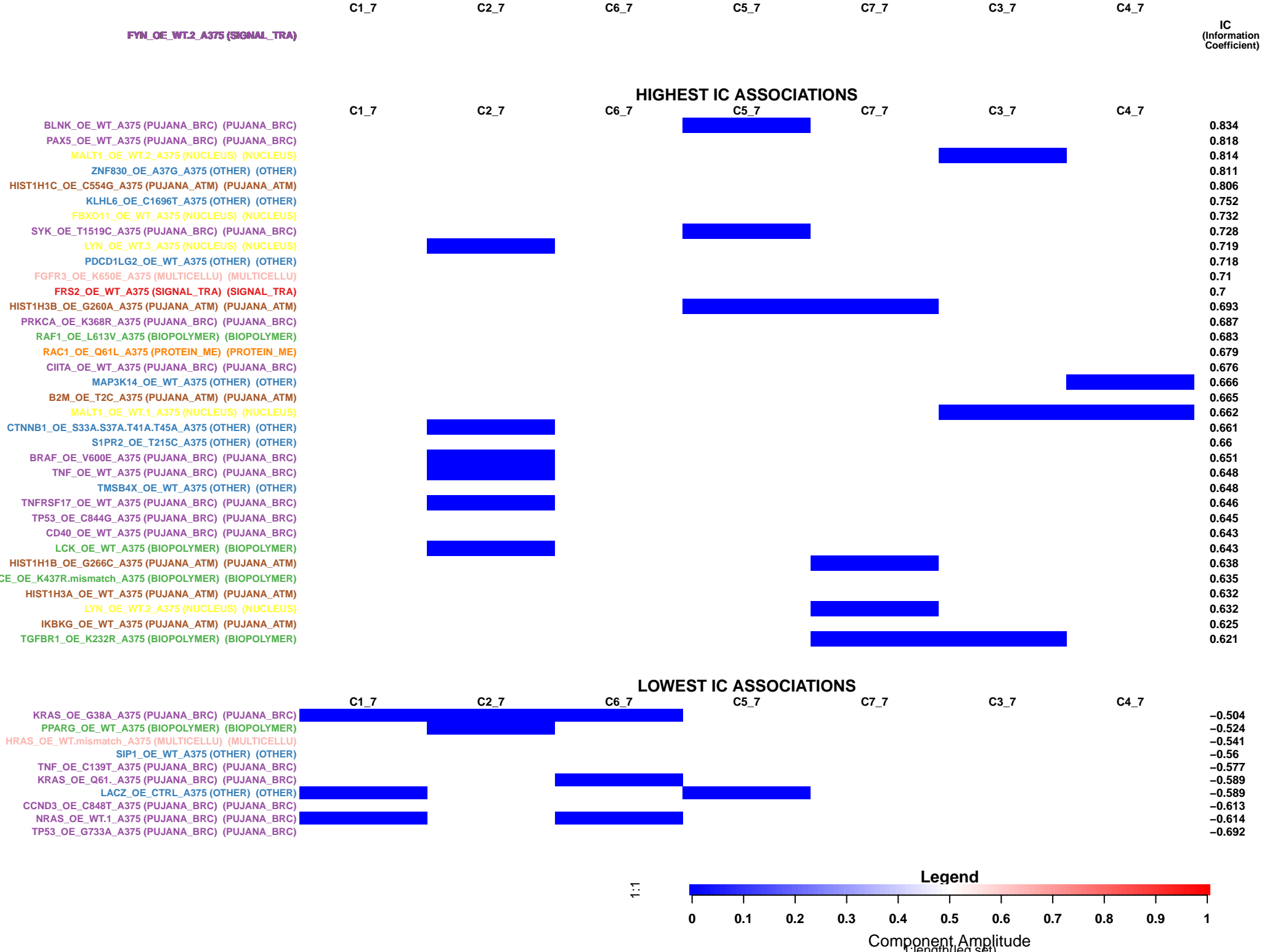


C6_7	C1_7	C3_7	C7_7	C5_7	C2_7	C4_7	IC (Information Coefficient)
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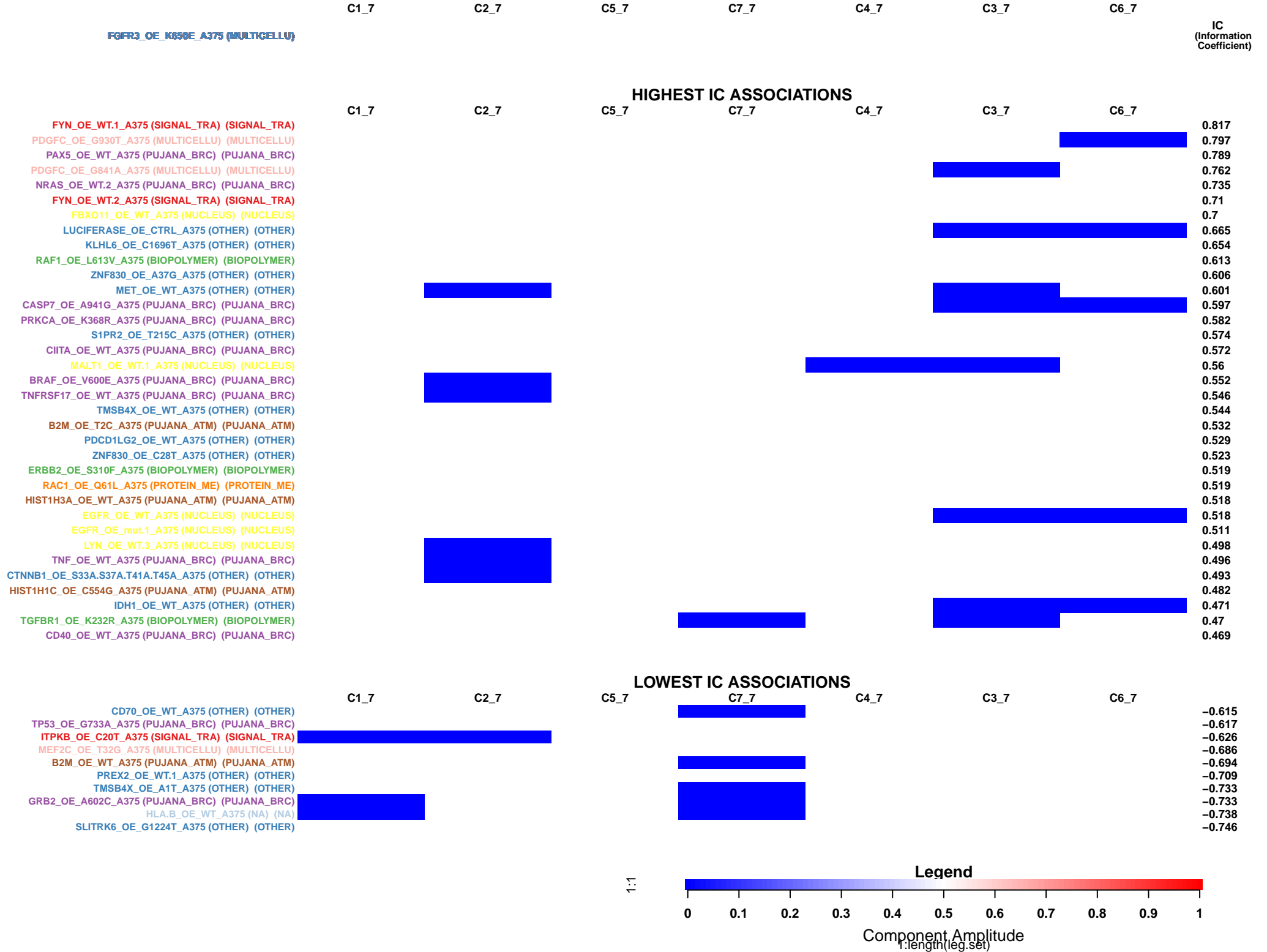




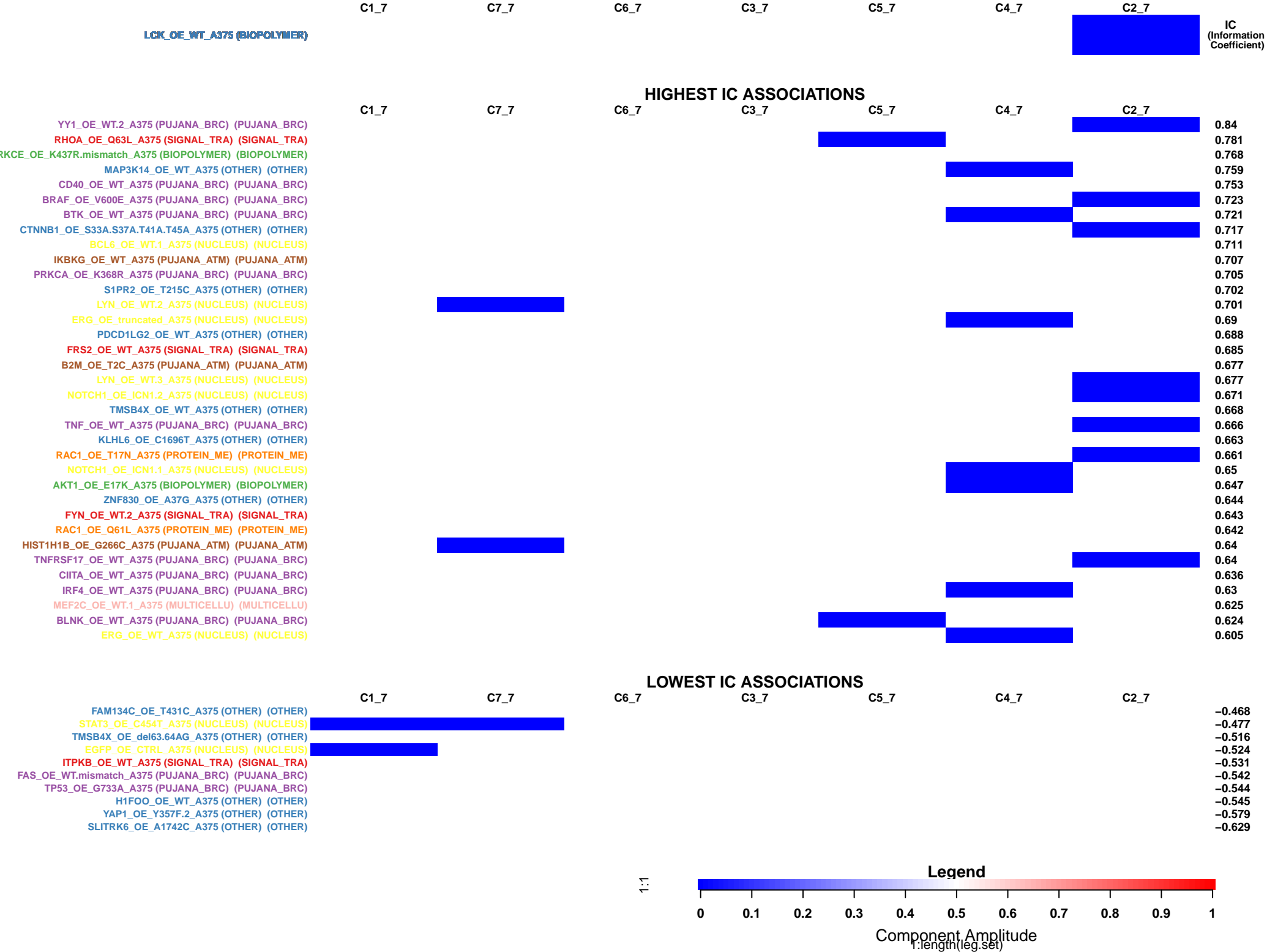
FYN\_OE\_WT.2\_A375 (SIGNAL\_TRA)



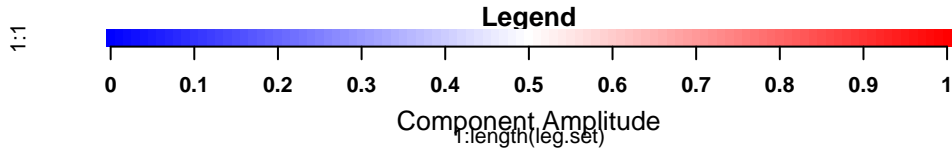
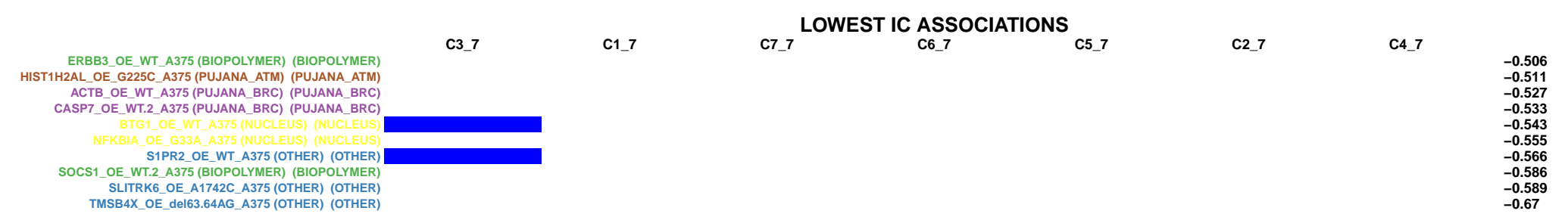
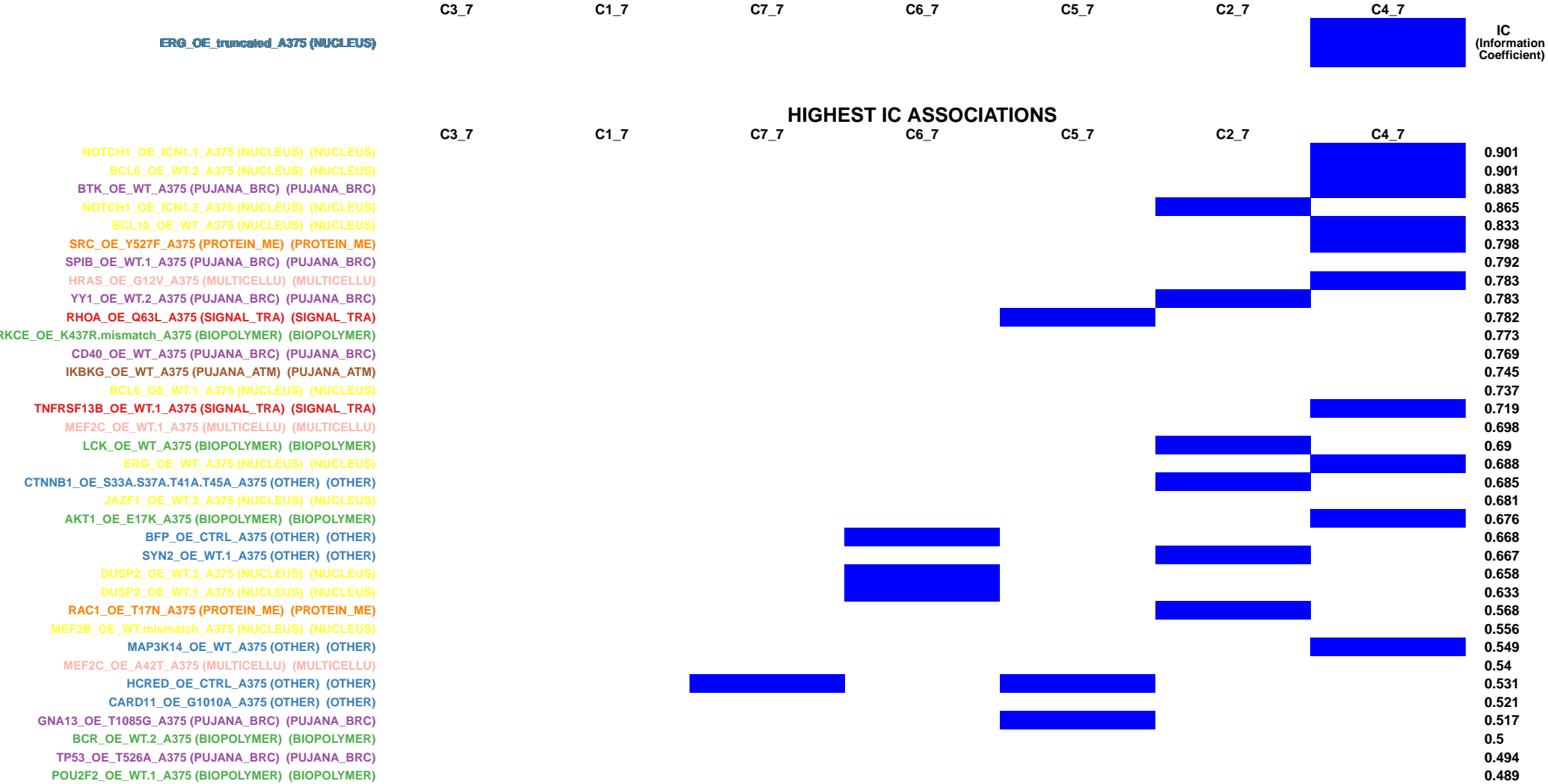
FGFR3\_OE\_K650E\_A375 (MULTICELLU)



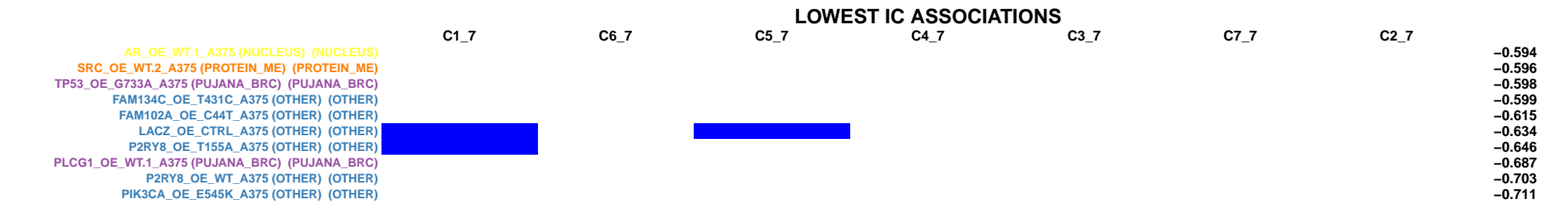
LCK\_OE\_WT\_A375 (BIOPOLYMER)



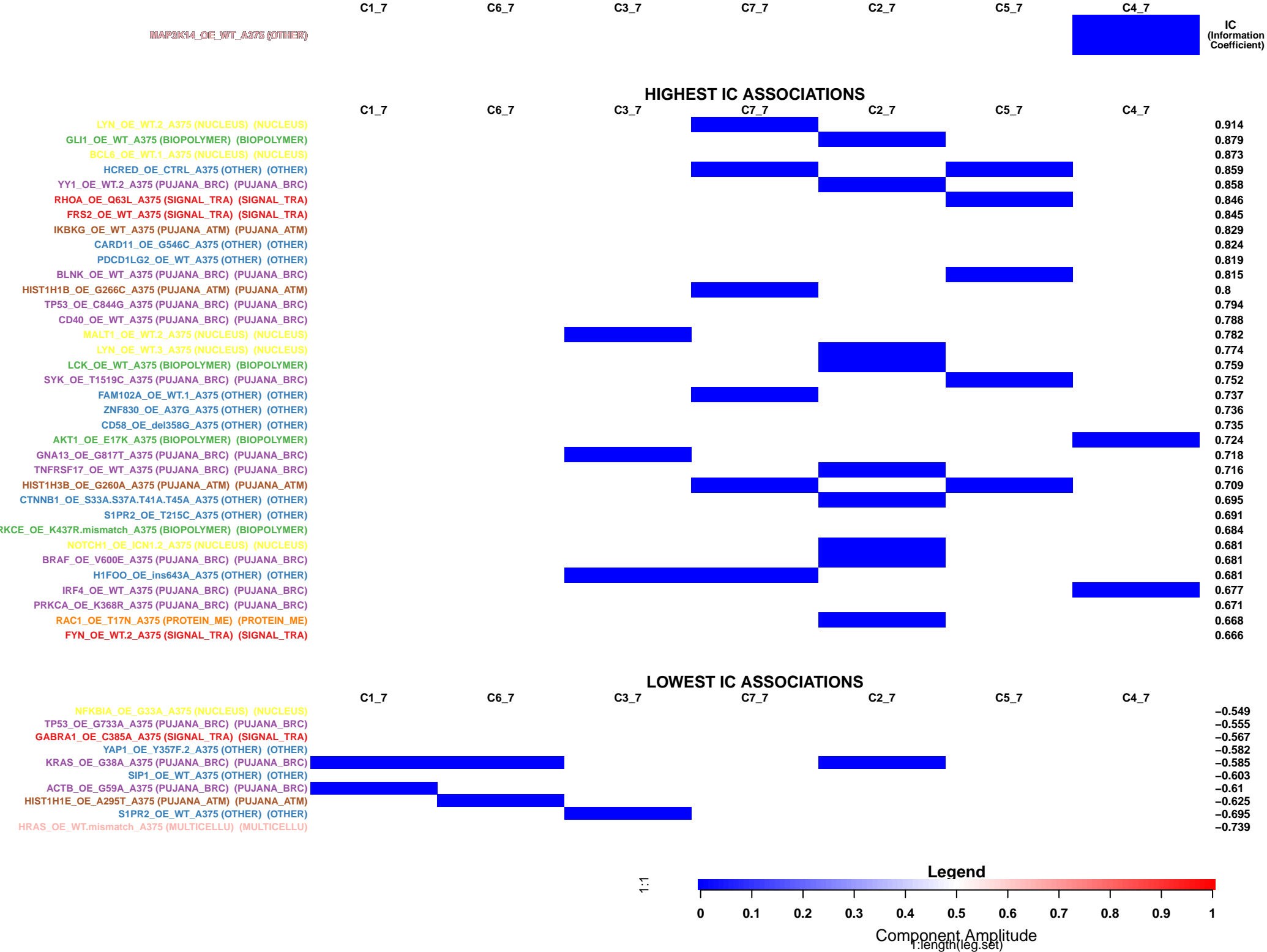
ERG\_OE\_truncated\_A375 (NUCLEUS)



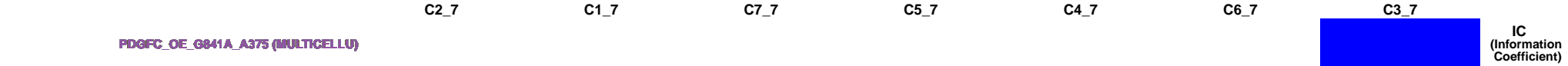
C1_7	C6_7	C5_7	C4_7	C3_7	C7_7	C2_7	IC (Information Coefficient)
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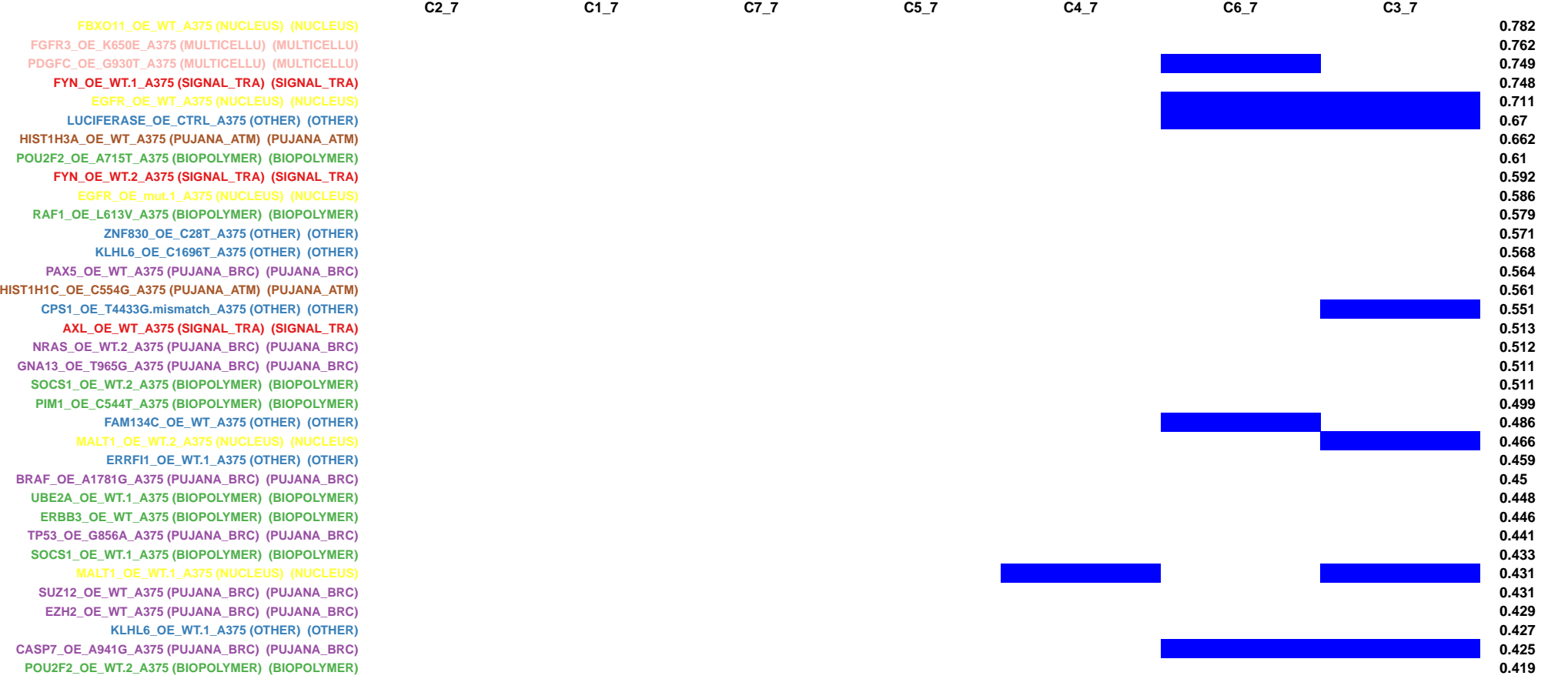
MAP3K14\_OE\_WT\_A375 (OTHER)



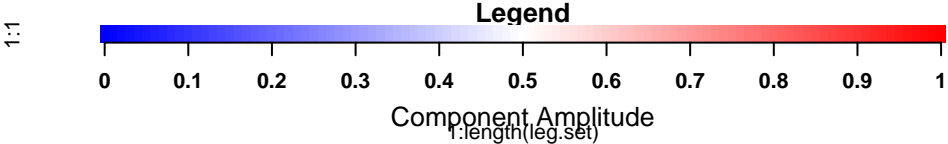
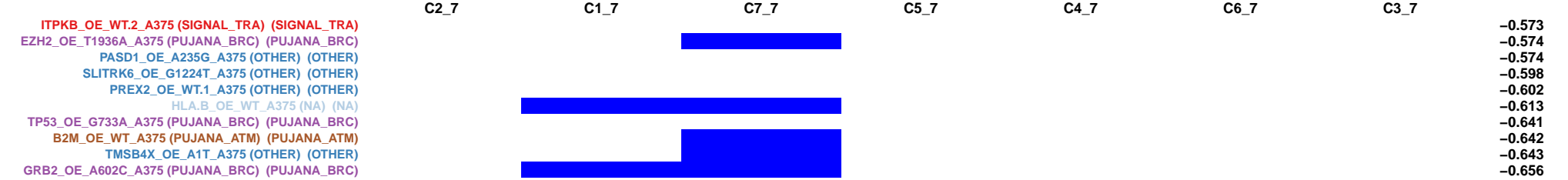
PDGFC\_OE\_G841A\_A375 (MULTICELLU)



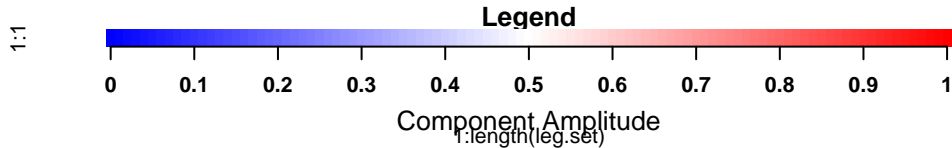
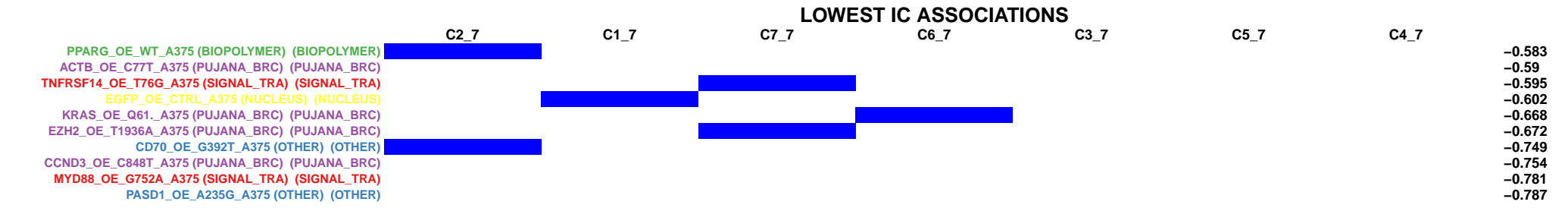
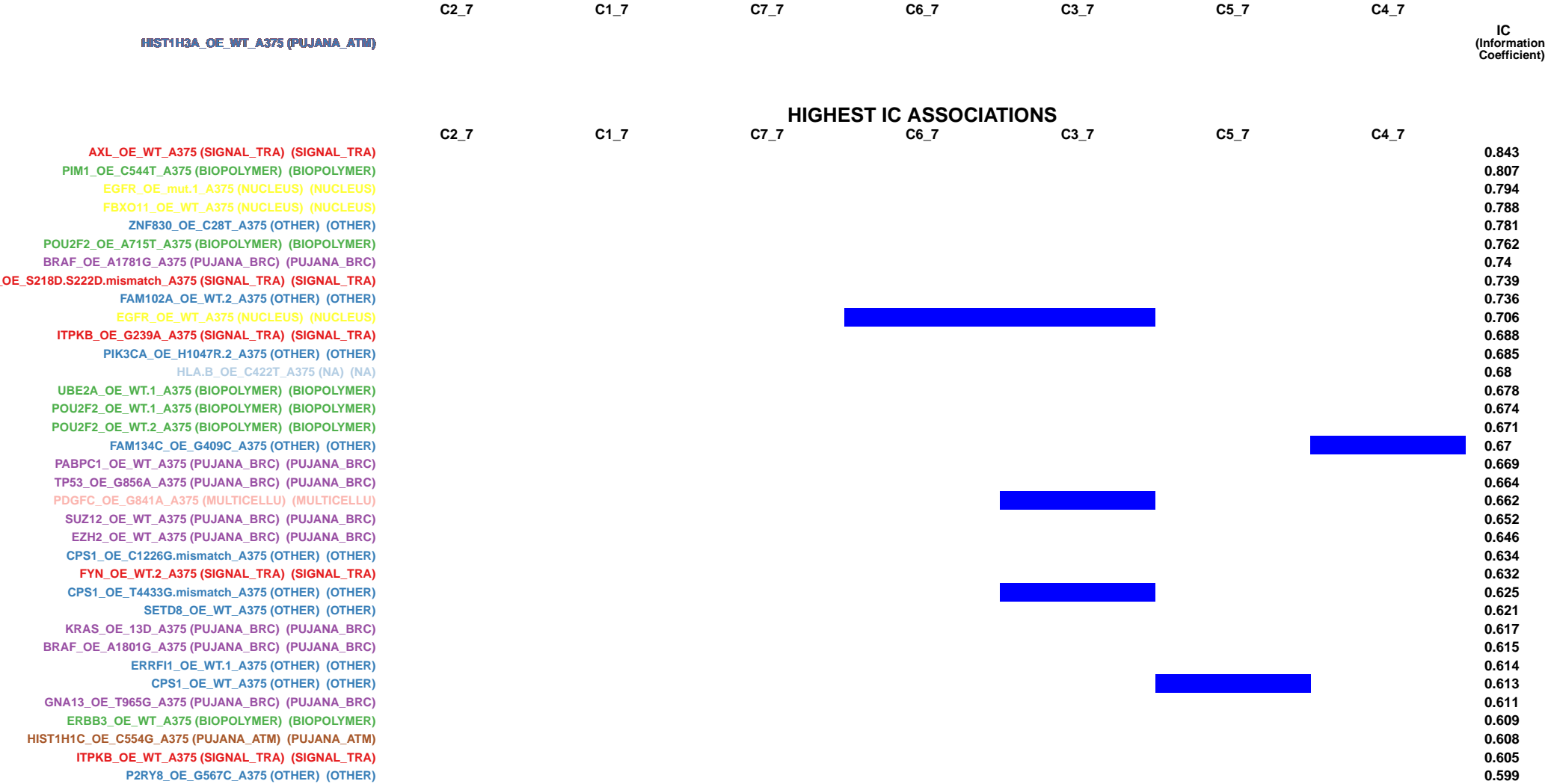
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS

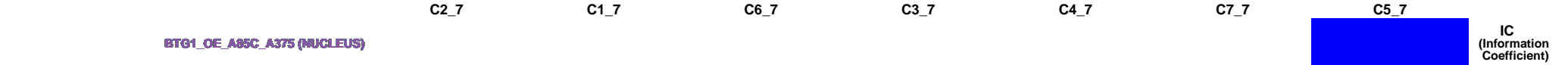


HIST1H3A\_OE\_WT\_A375 (PUJANA\_ATM)

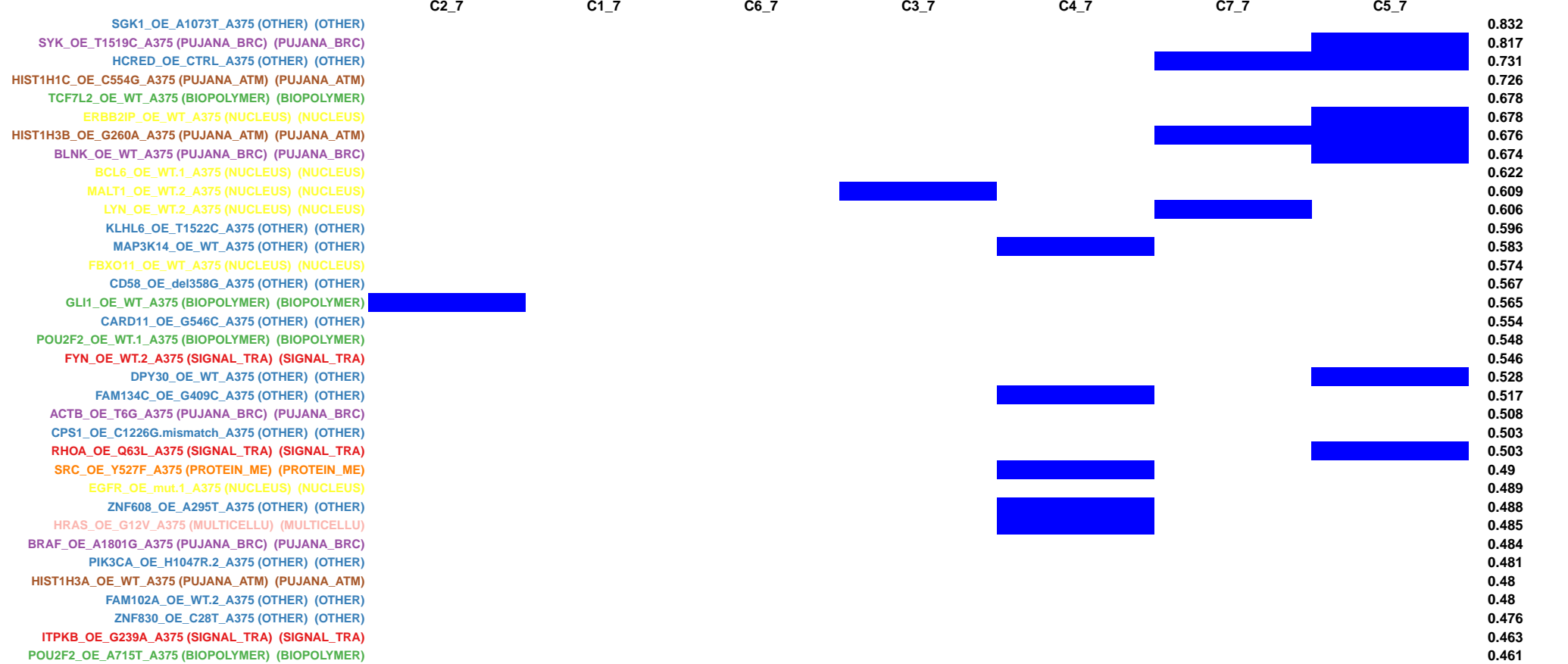




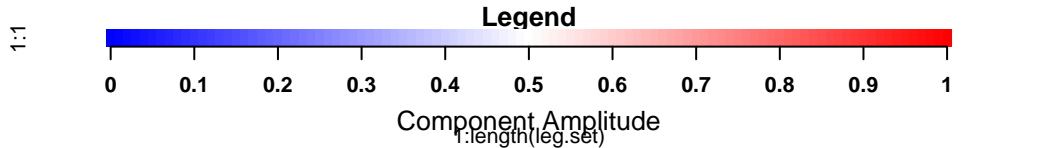
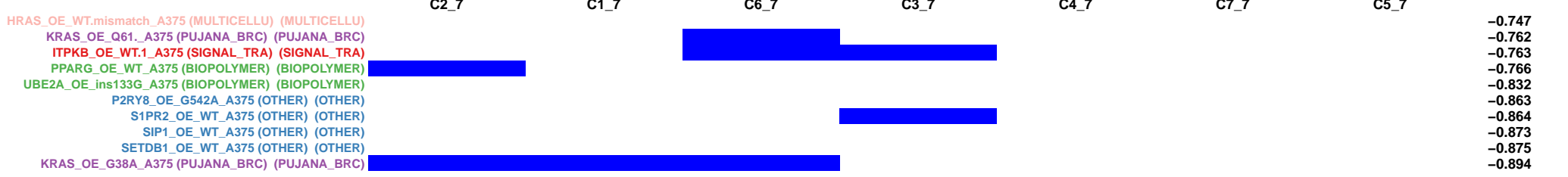
BTG1\_OE\_A85C\_A375 (NUCLEUS)



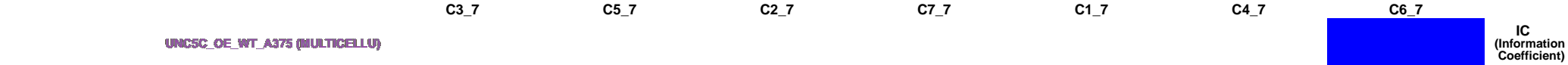
HIGHEST IC ASSOCIATIONS



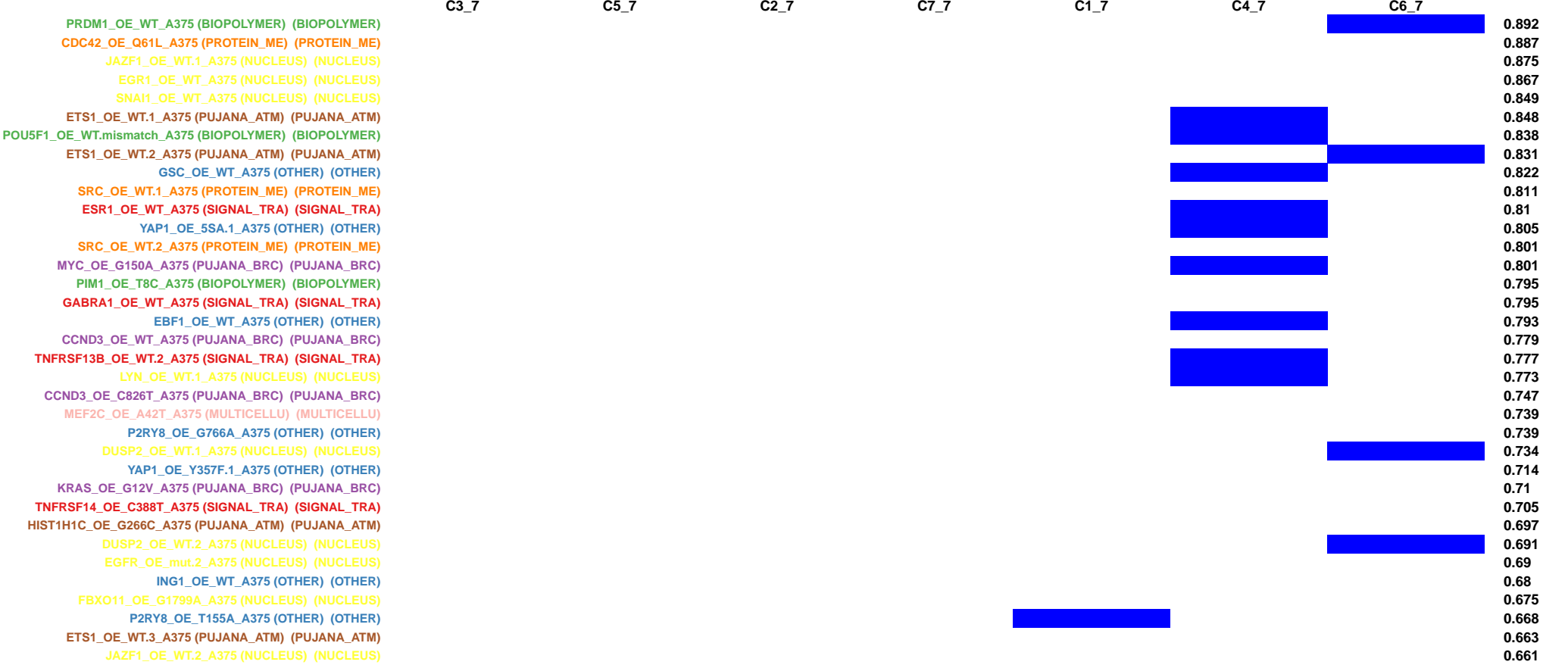
LOWEST IC ASSOCIATIONS



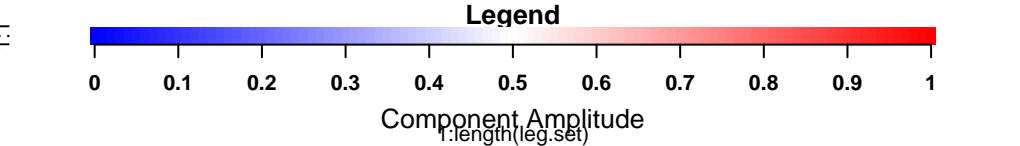
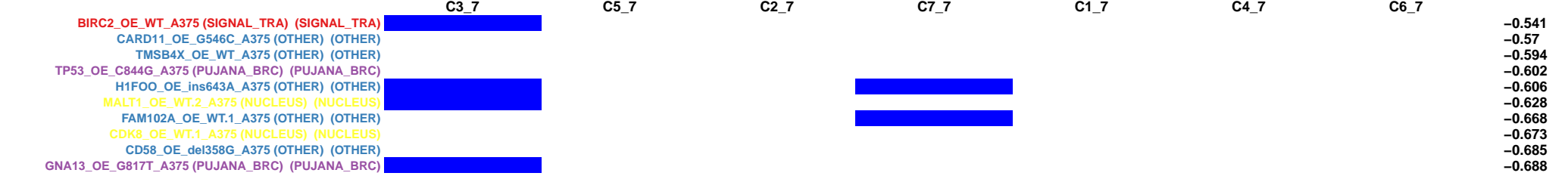
UNC5C\_OE\_WT\_A375 (MULTICELLU)



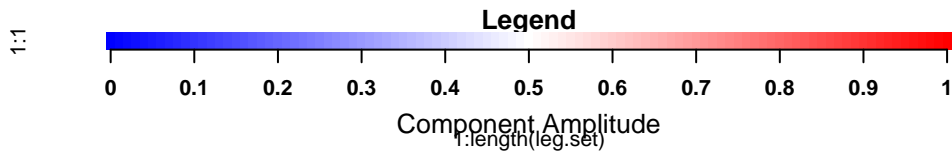
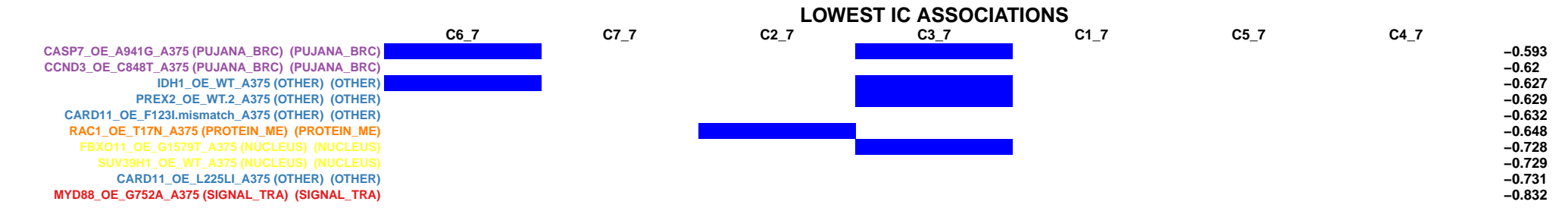
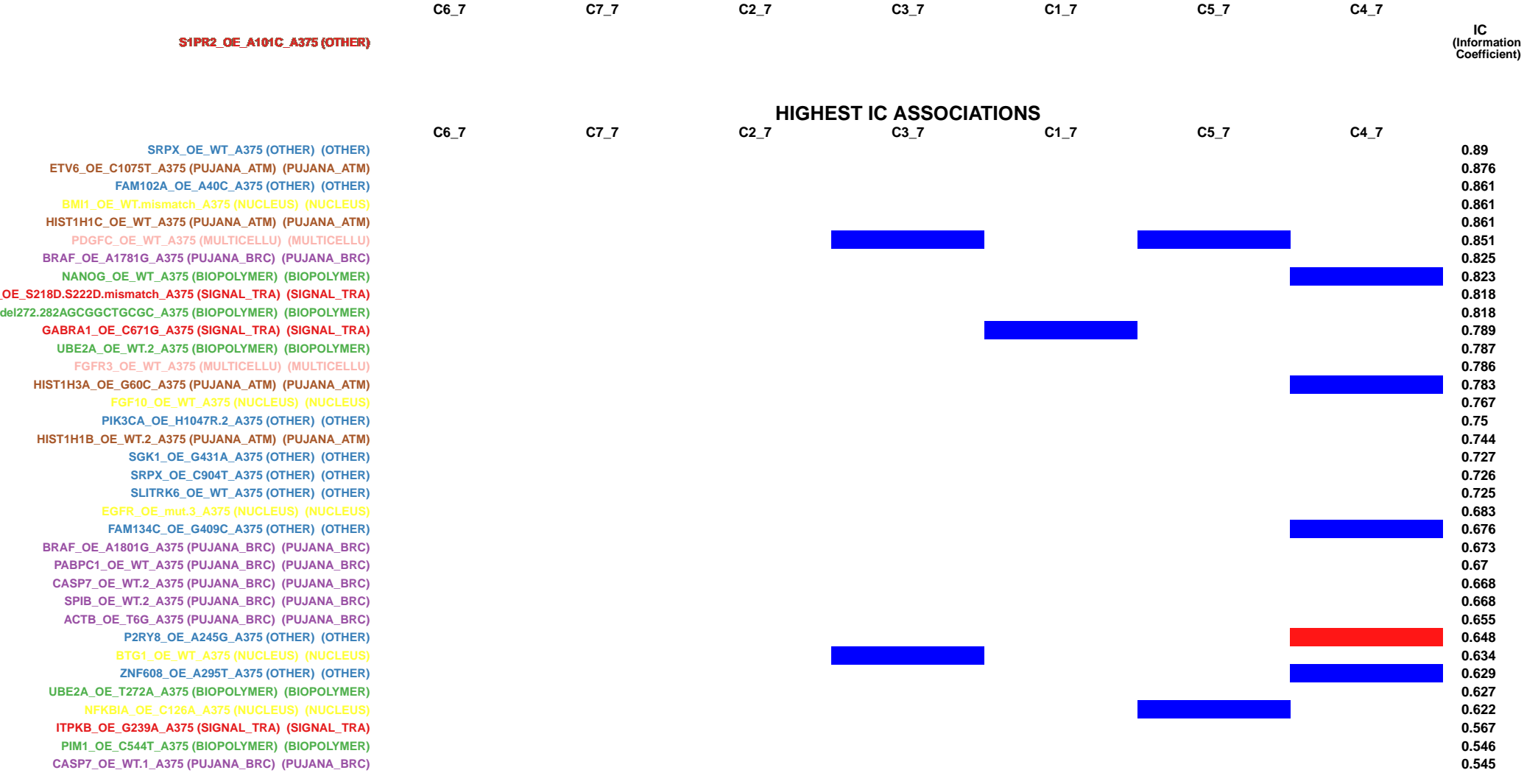
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS



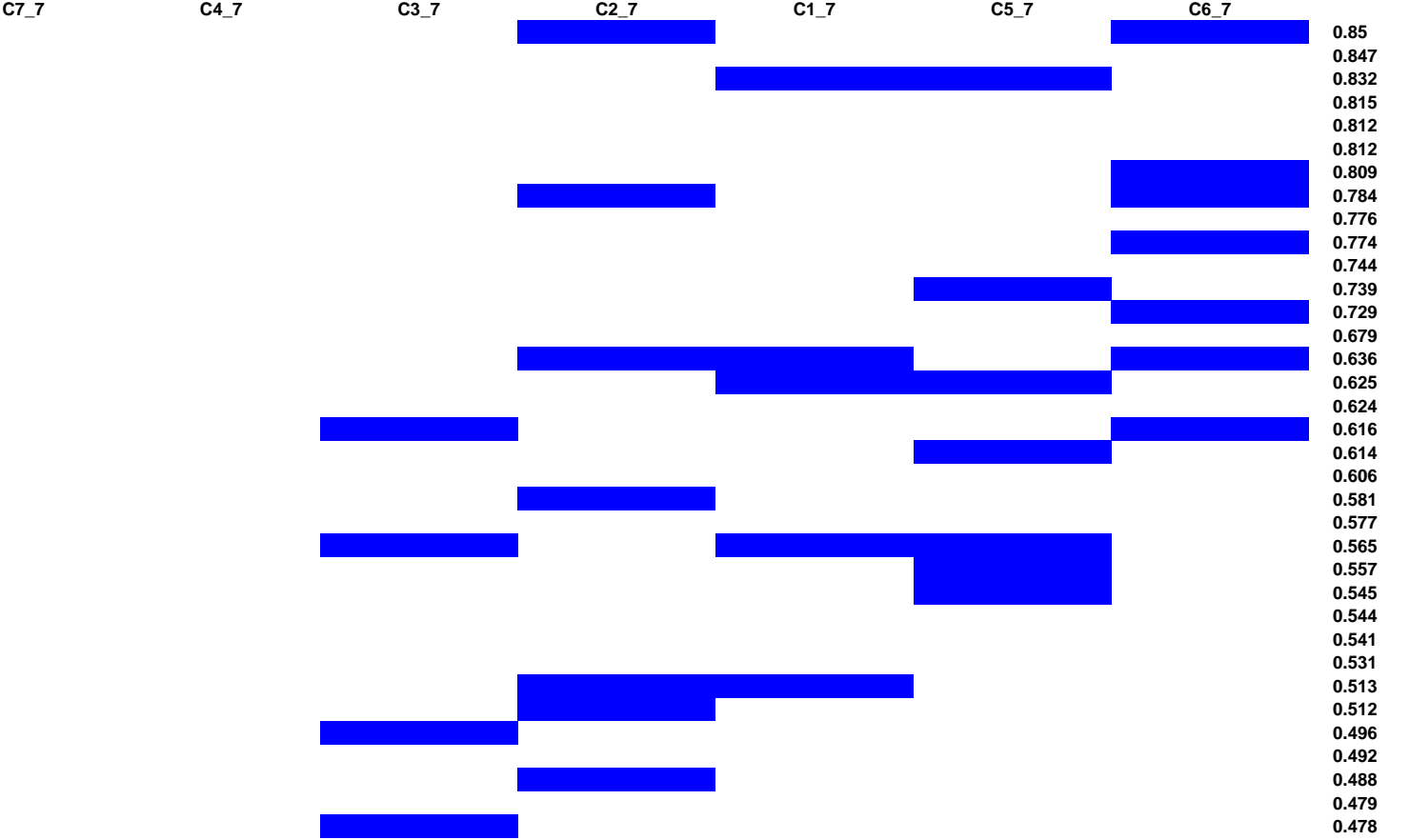
S1PR2\_OE\_A101C\_A375 (OTHER)



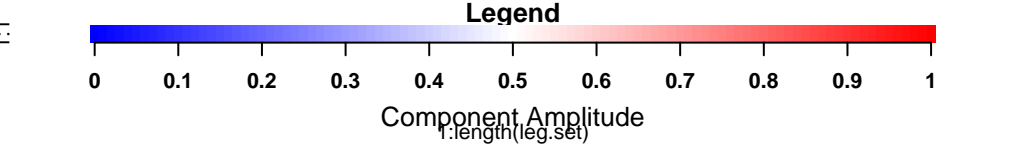
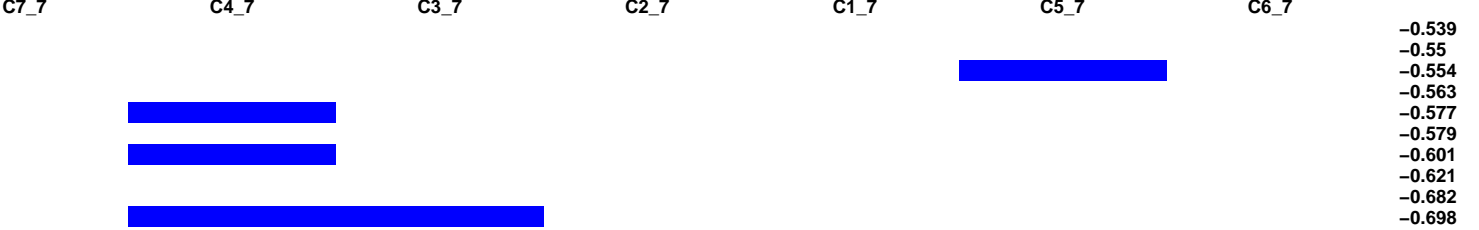
HIST1H3B\_OE\_WT\_A375 (PUJANA\_ATM)



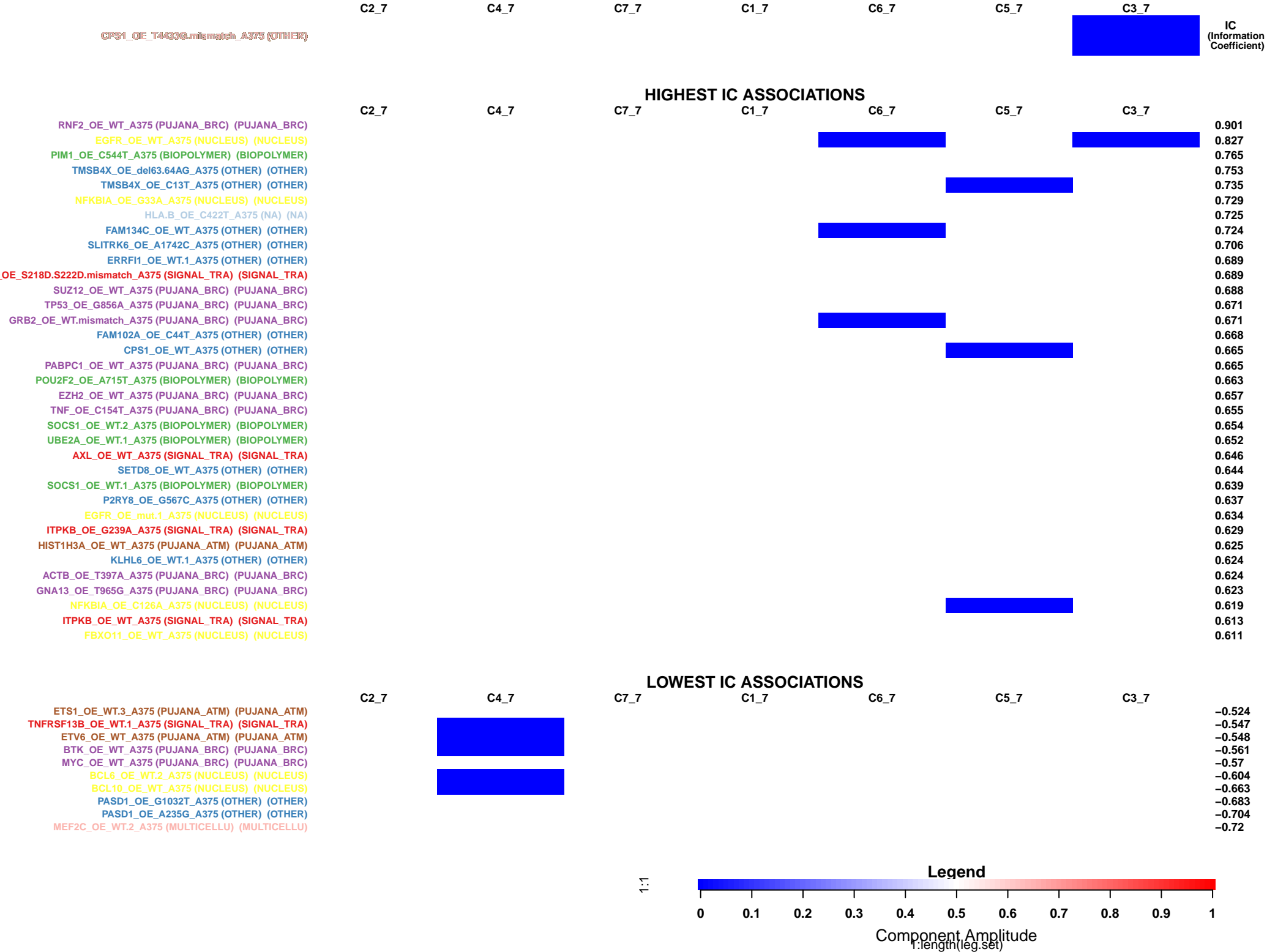
HIGHEST IC ASSOCIATIONS



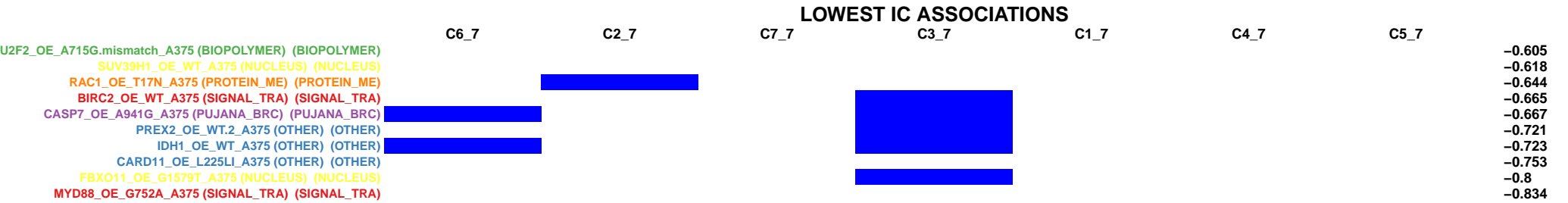
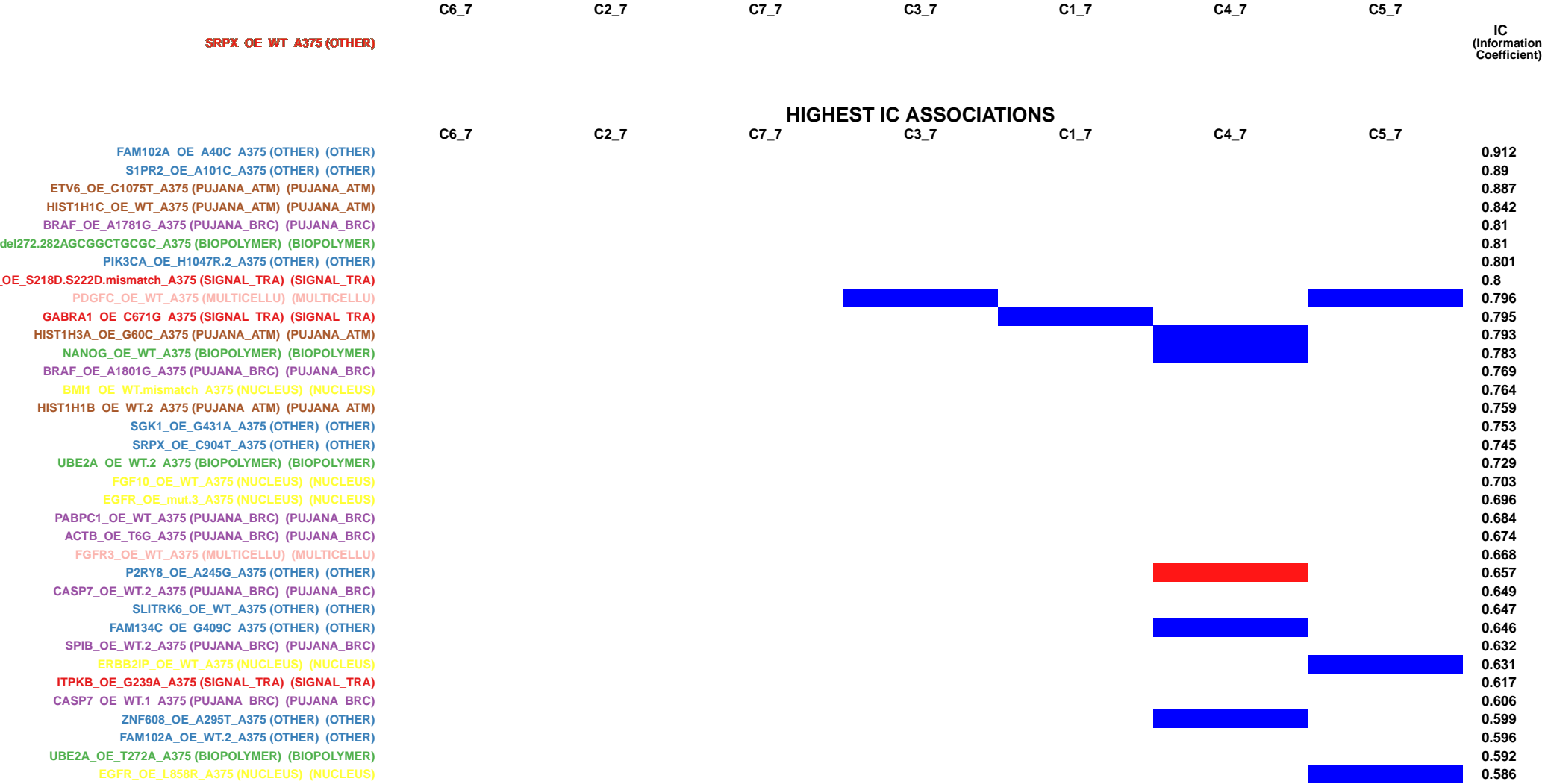
LOWEST IC ASSOCIATIONS



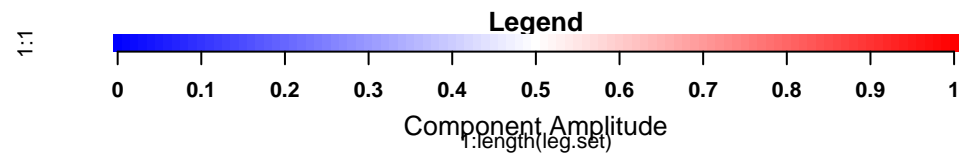
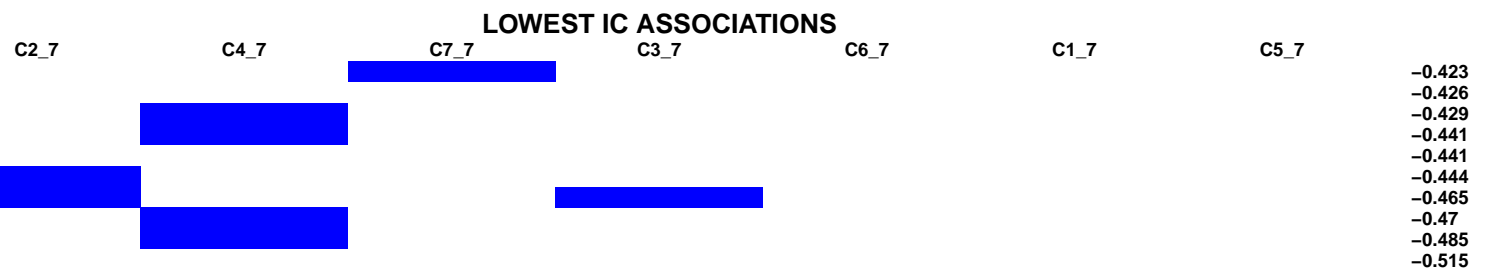
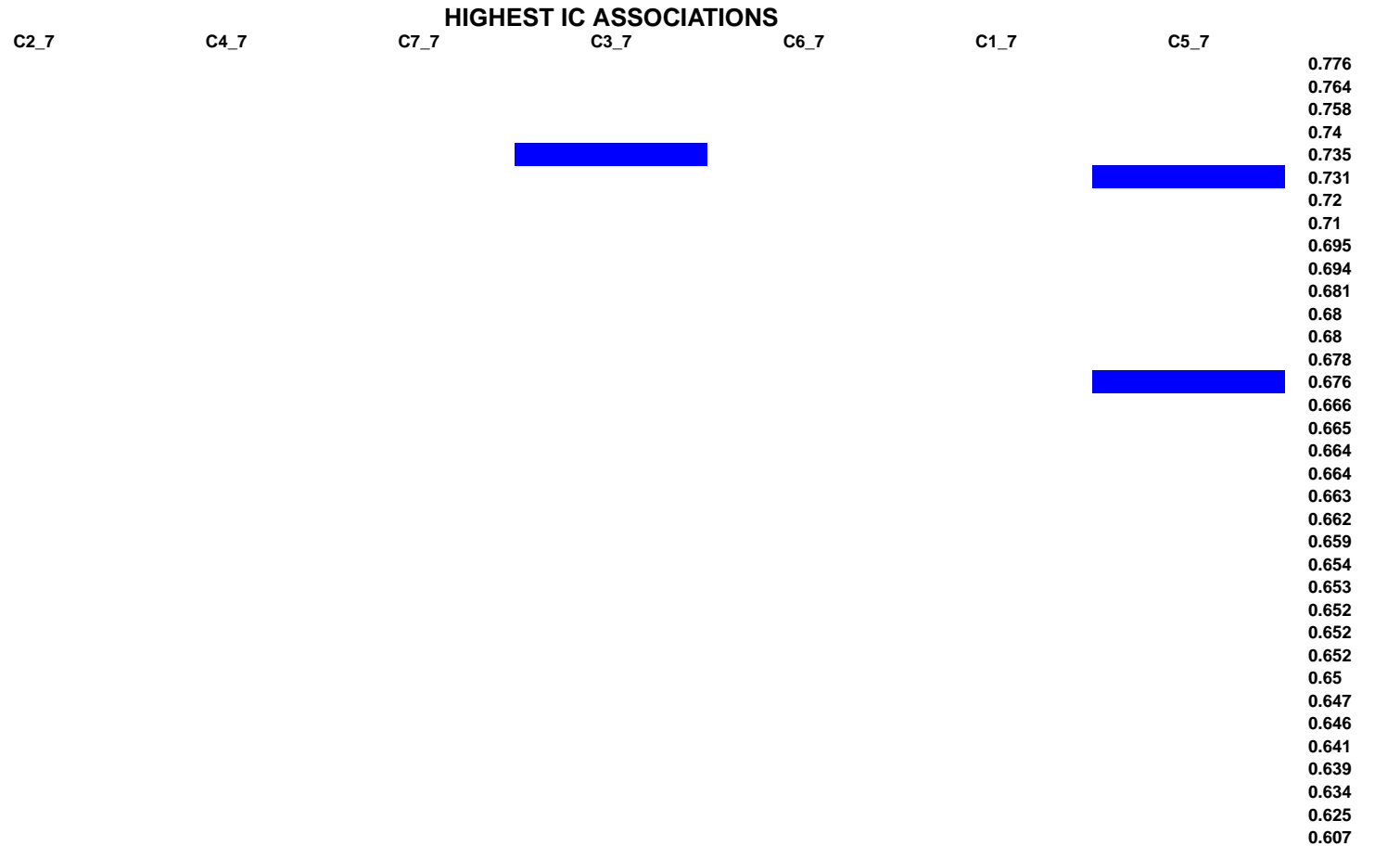
CPS1\_OE\_T4433G.mismatch\_A375 (OTHER)



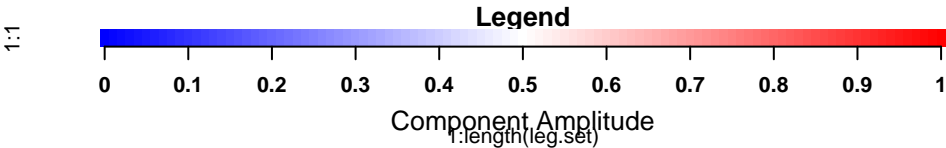
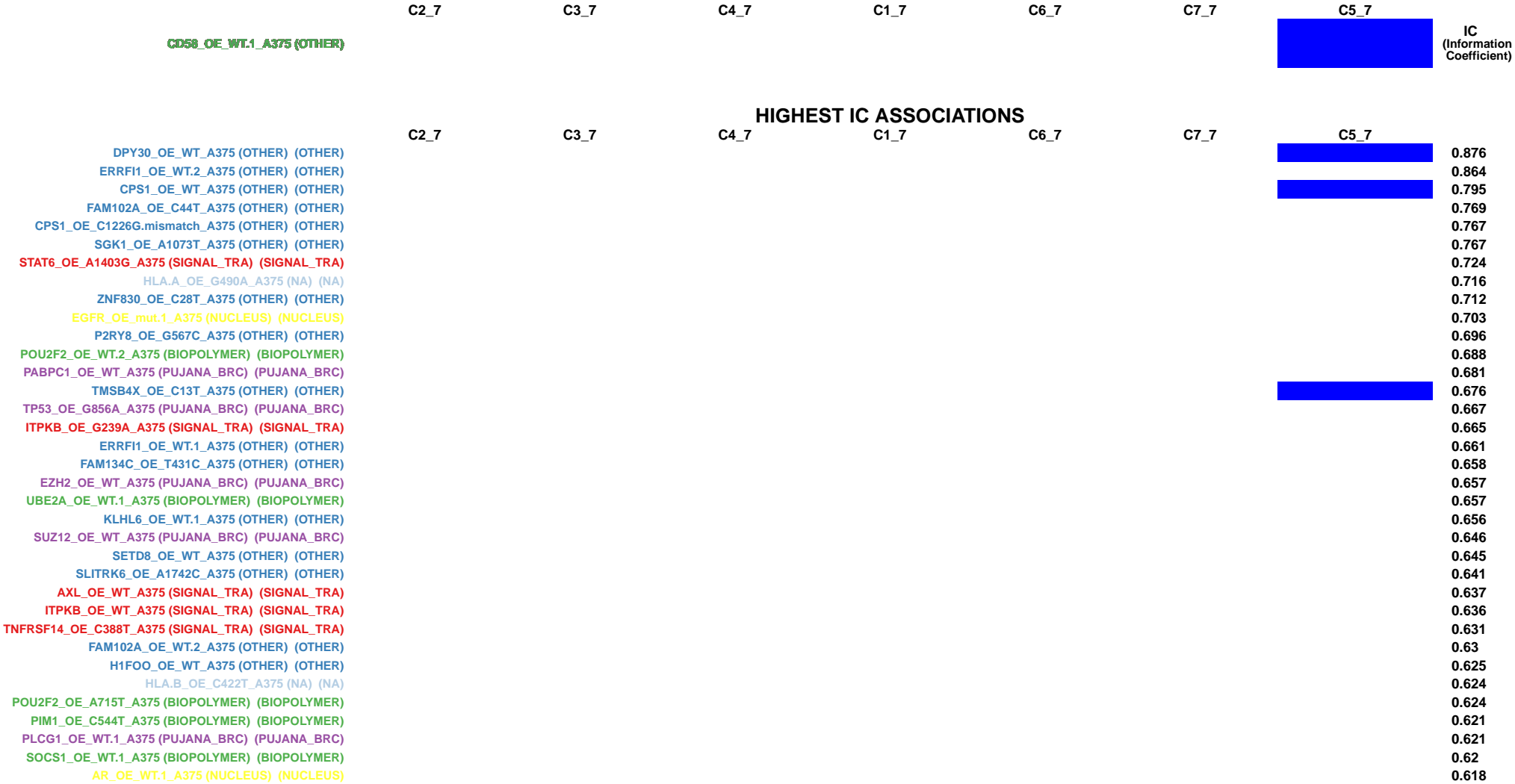
SRPX\_OE\_WT\_A375 (OTHER)



C2_7	C4_7	C7_7	C3_7	C6_7	C1_7	C5_7	IC (Information Coefficient)



CD58\_OE\_WT.1\_A375 (OTHER)



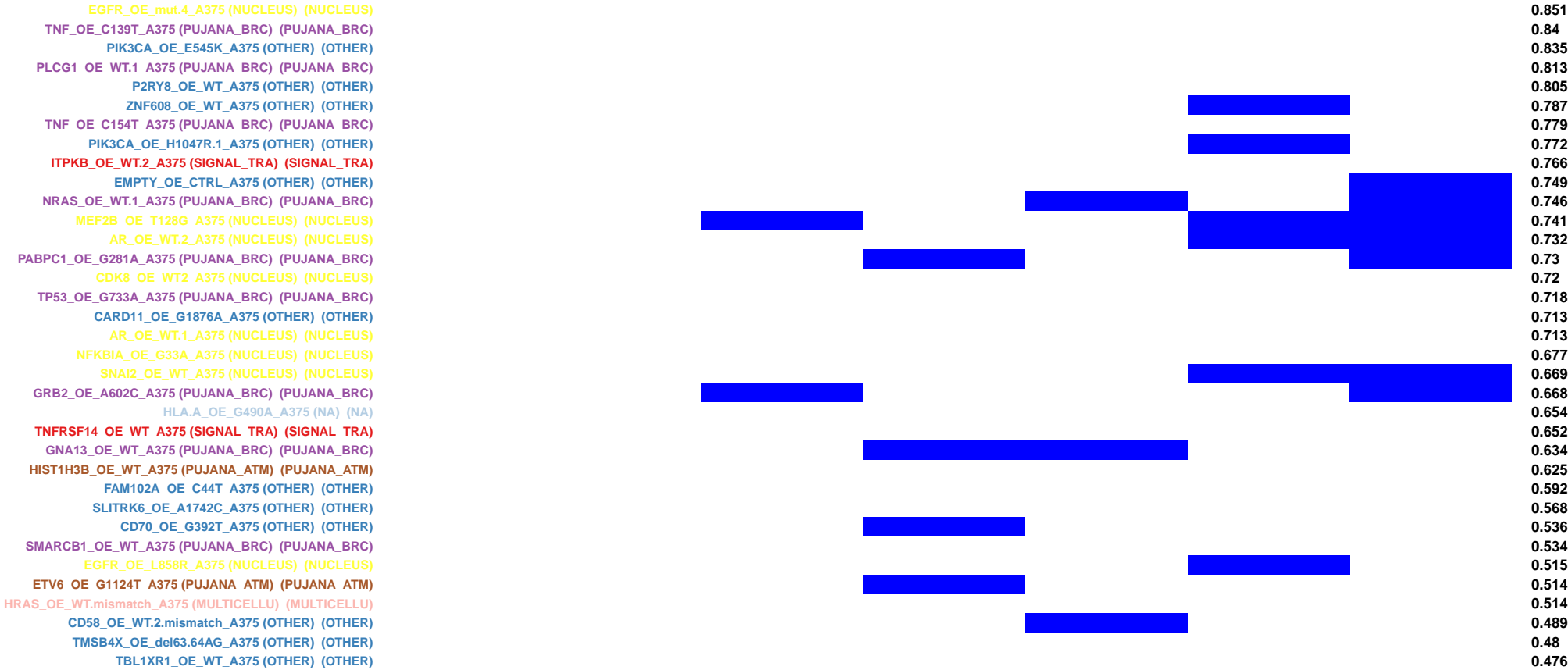


LACZ\_OE\_CTRL\_A375 (OTHER)

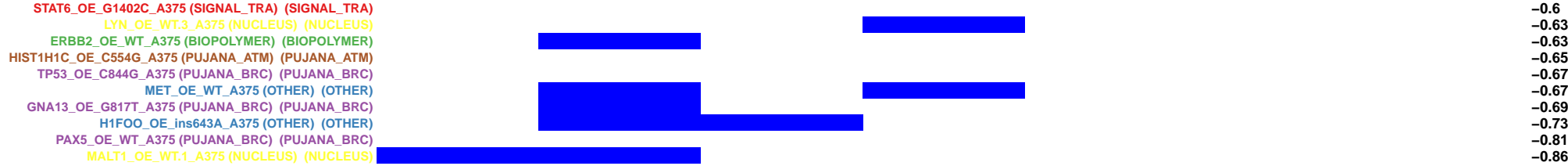
LACZ\_OE\_CTRL\_A375 (OTHER)

IC  
(Information  
Coefficient)

HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS



1:1

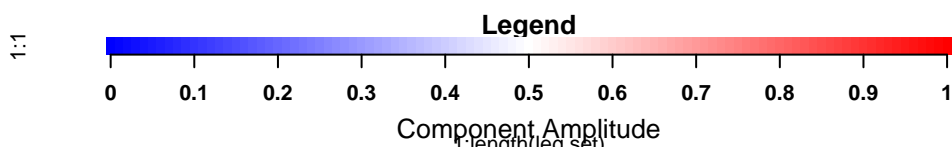
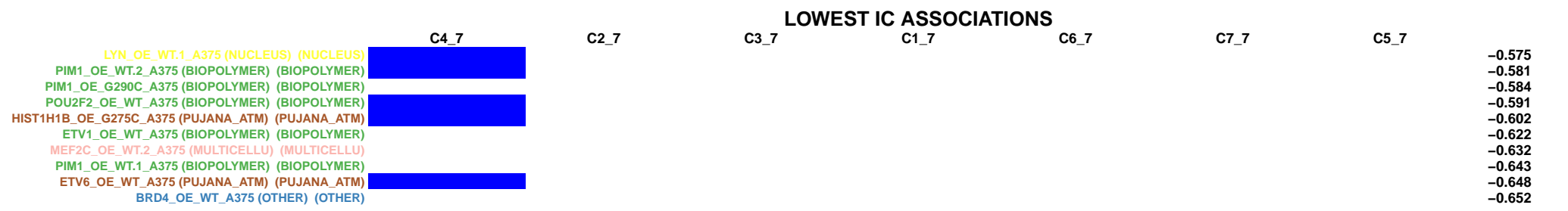
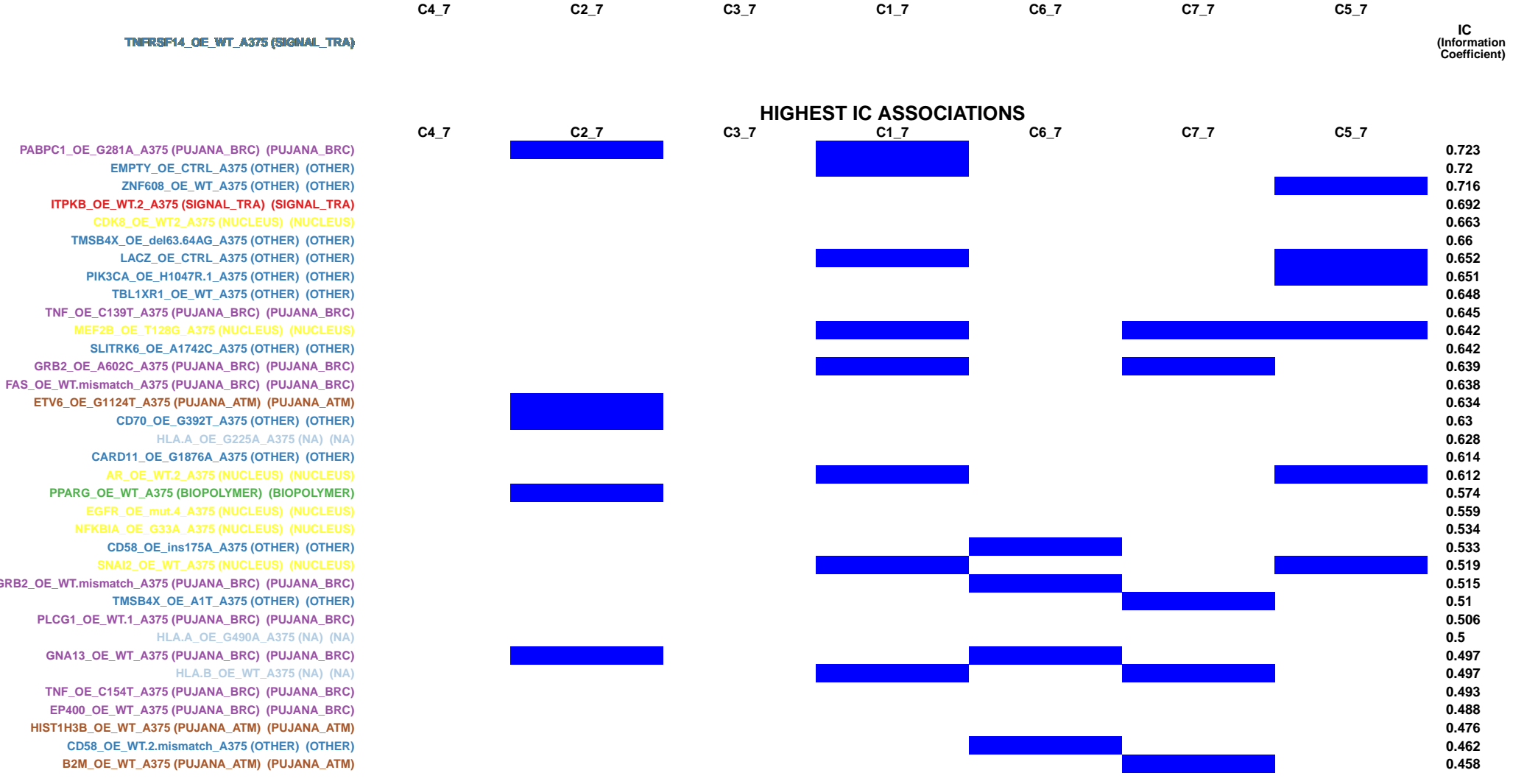
Legend

0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1

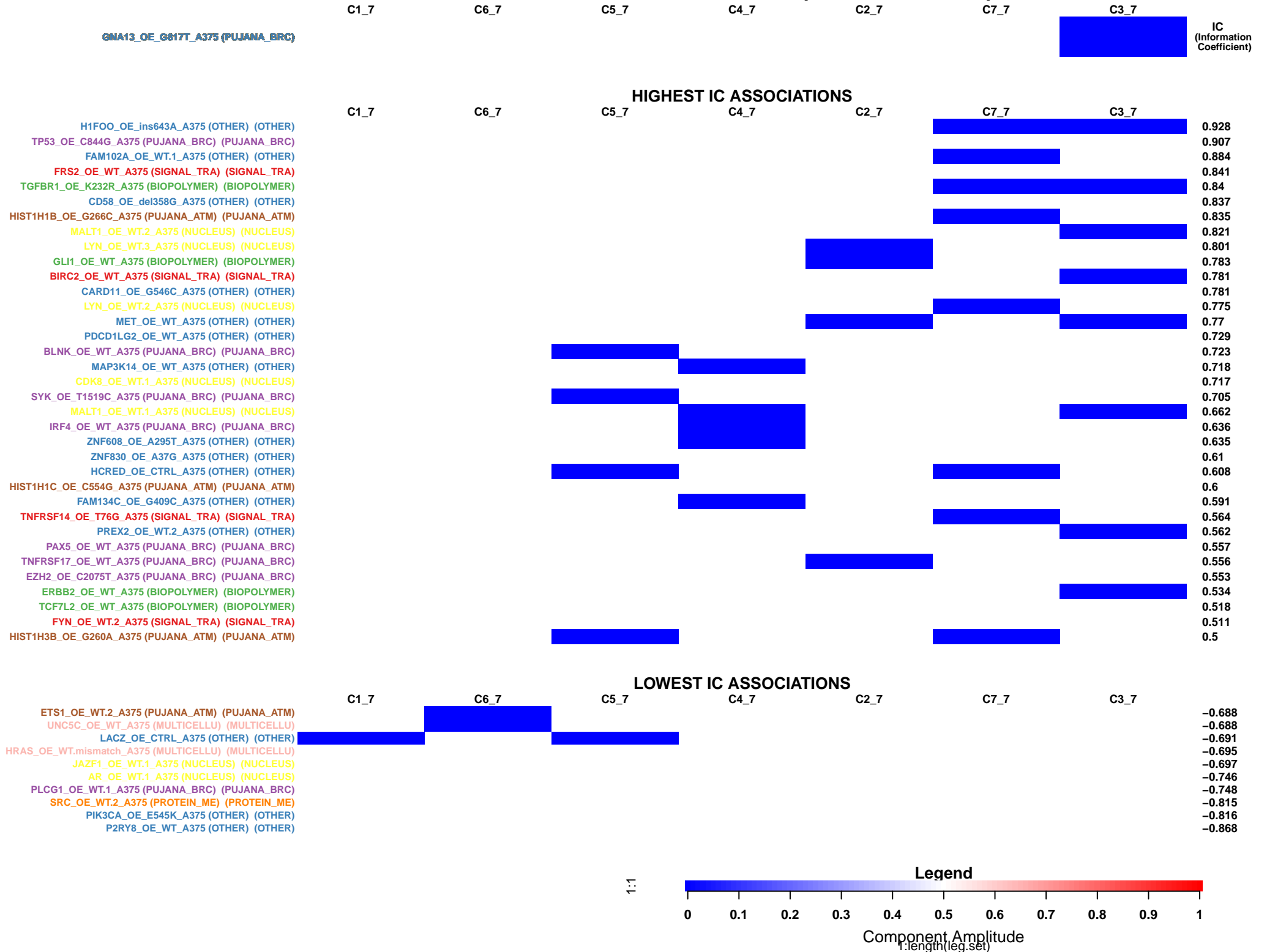
Component Amplitude

1:length(leg.set)

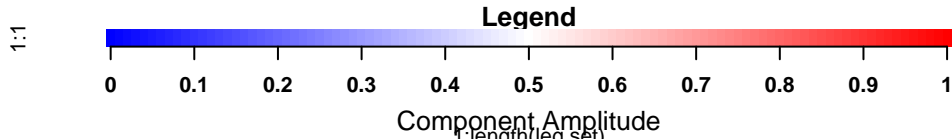
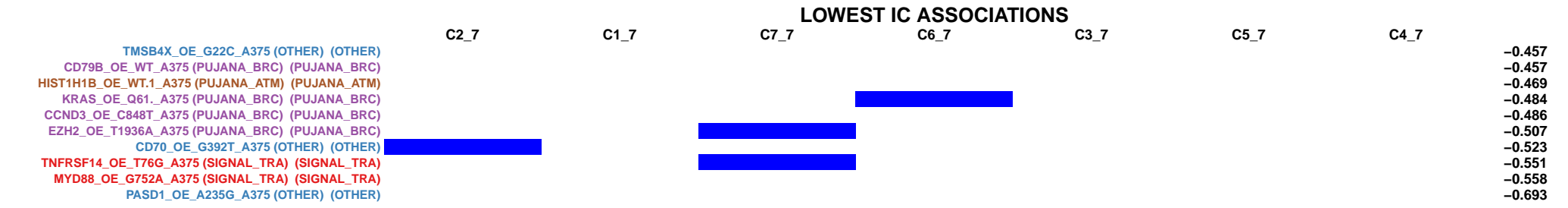
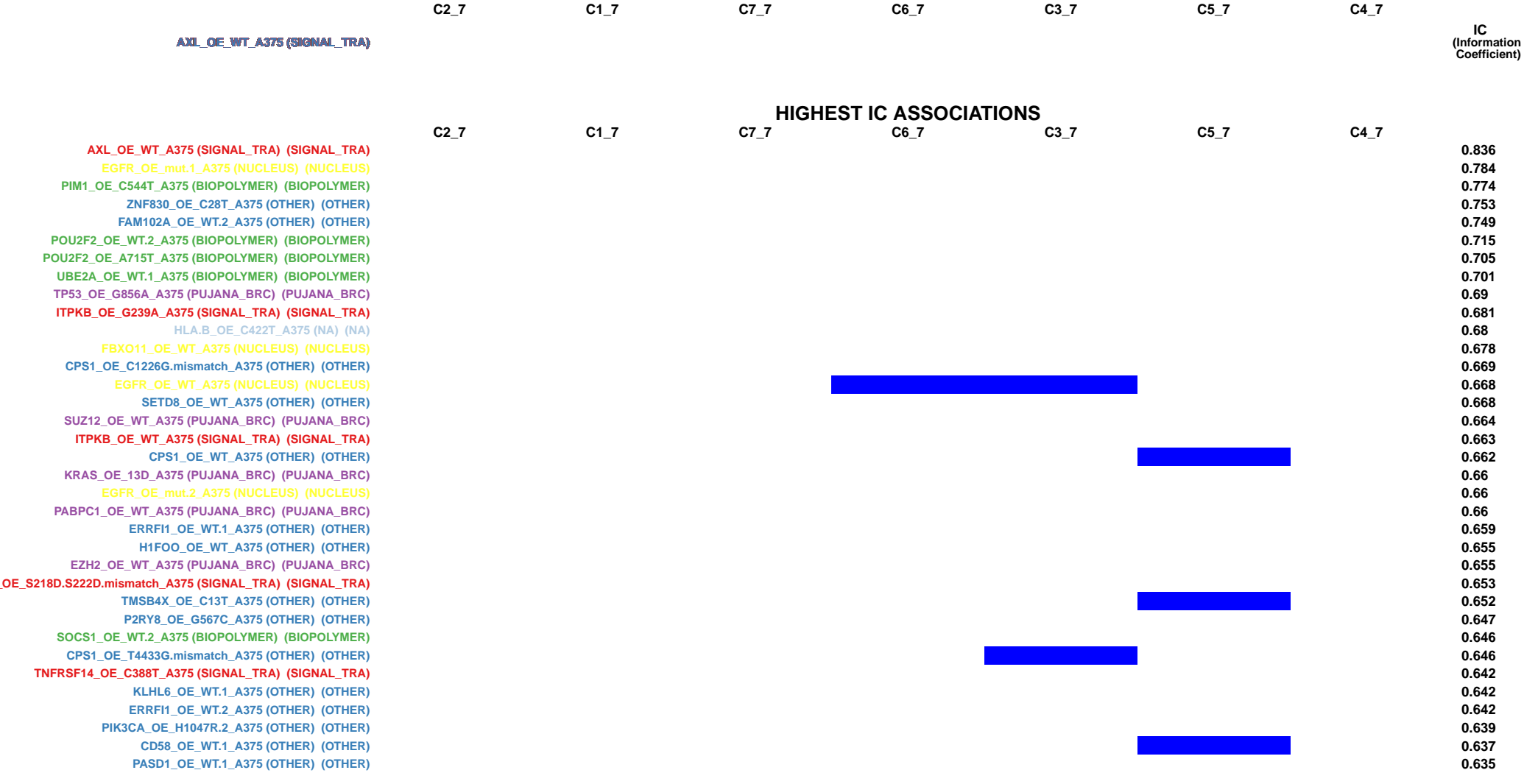
TNFRSF14\_OE\_WT\_A375 (SIGNAL\_TRA)



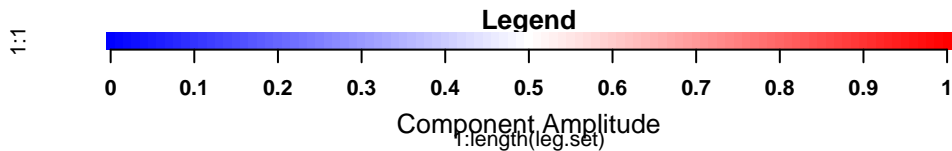
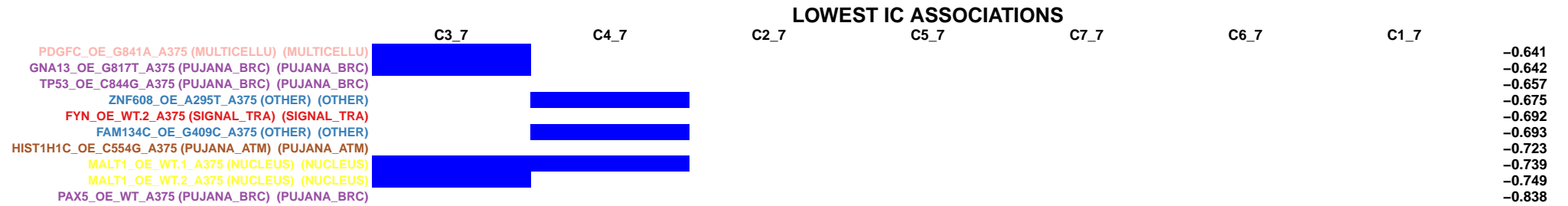
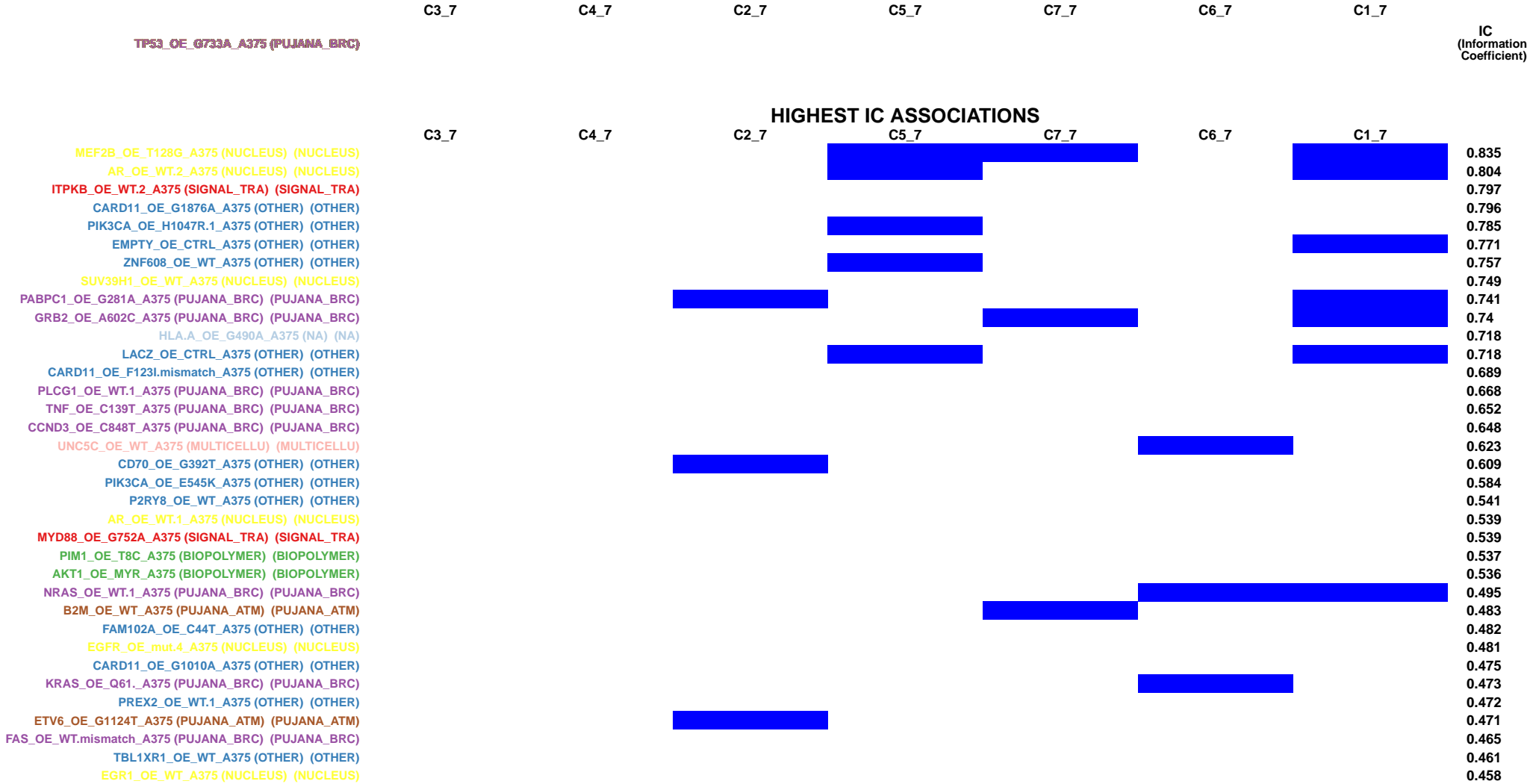
GNA13\_OE\_G817T\_A375 (PUJANA\_BRC)



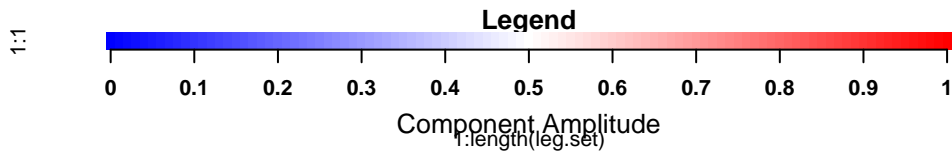
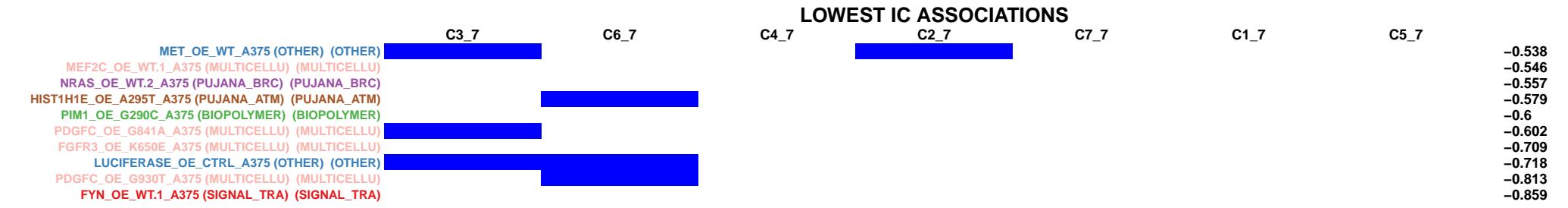
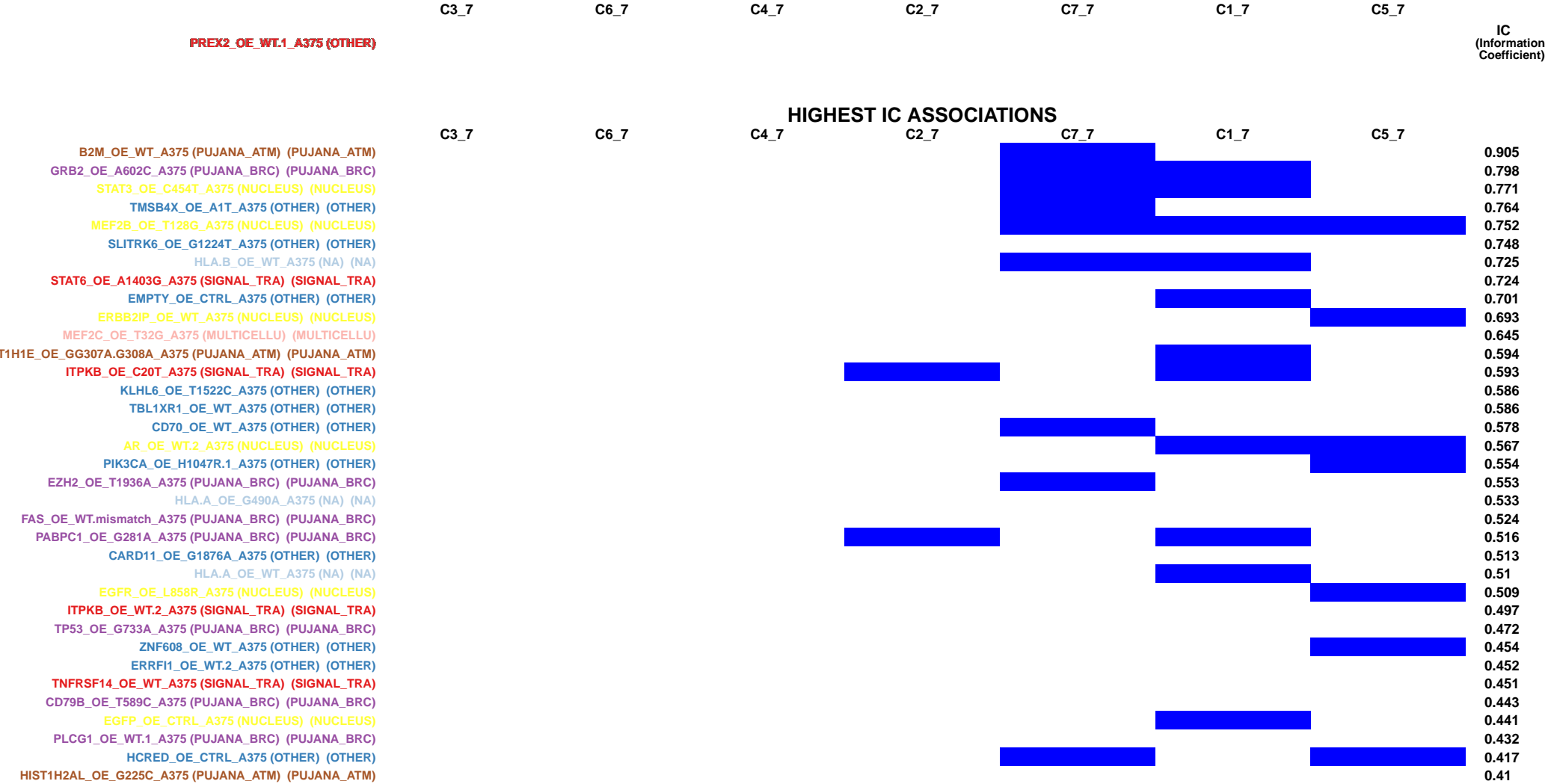
AXL\_OE\_WT\_A375 (SIGNAL\_TRA)



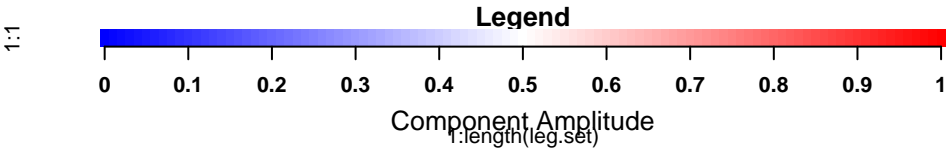
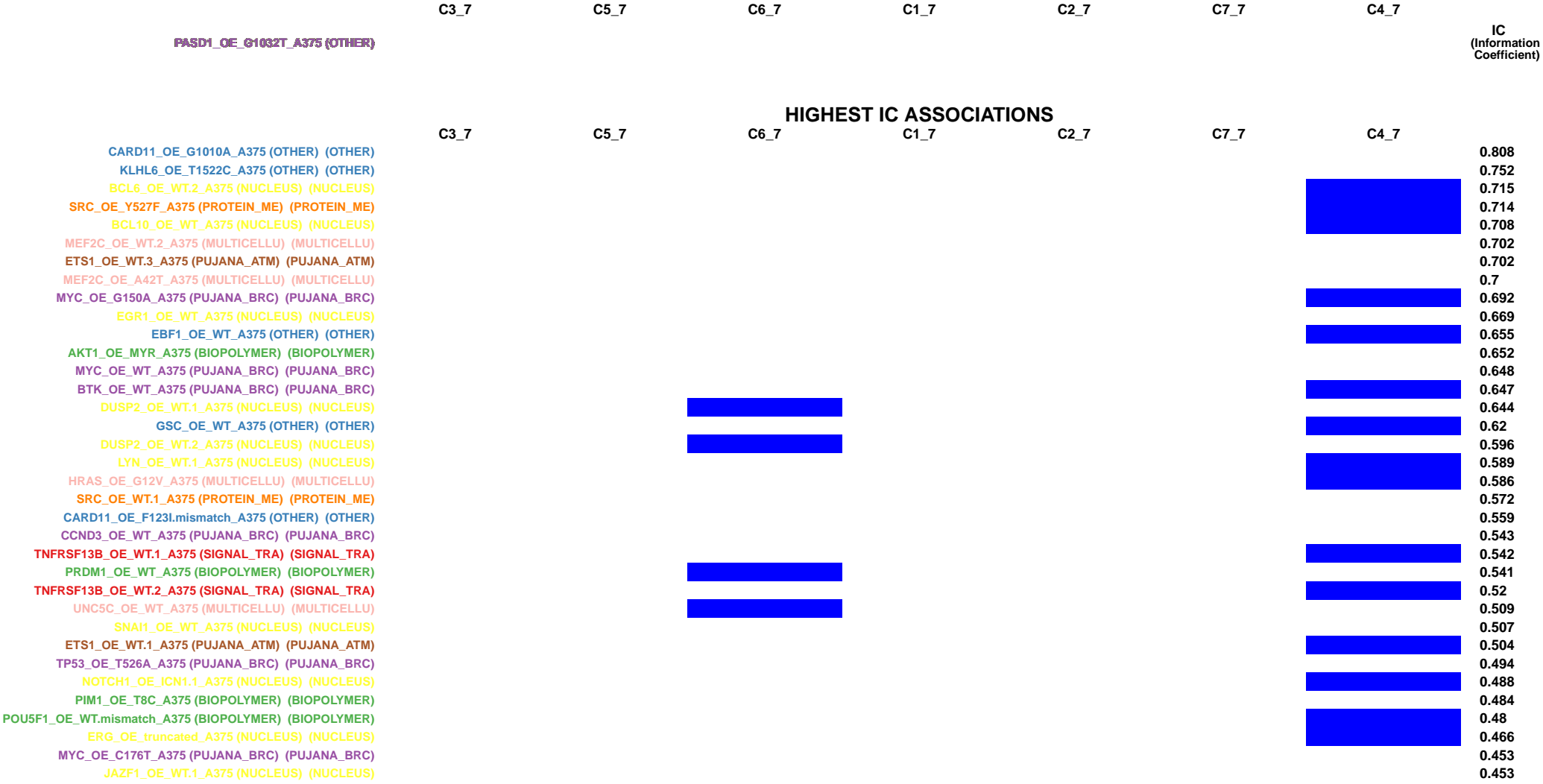
TP53\_OE\_G733A\_A375 (PUJANA\_BRC)



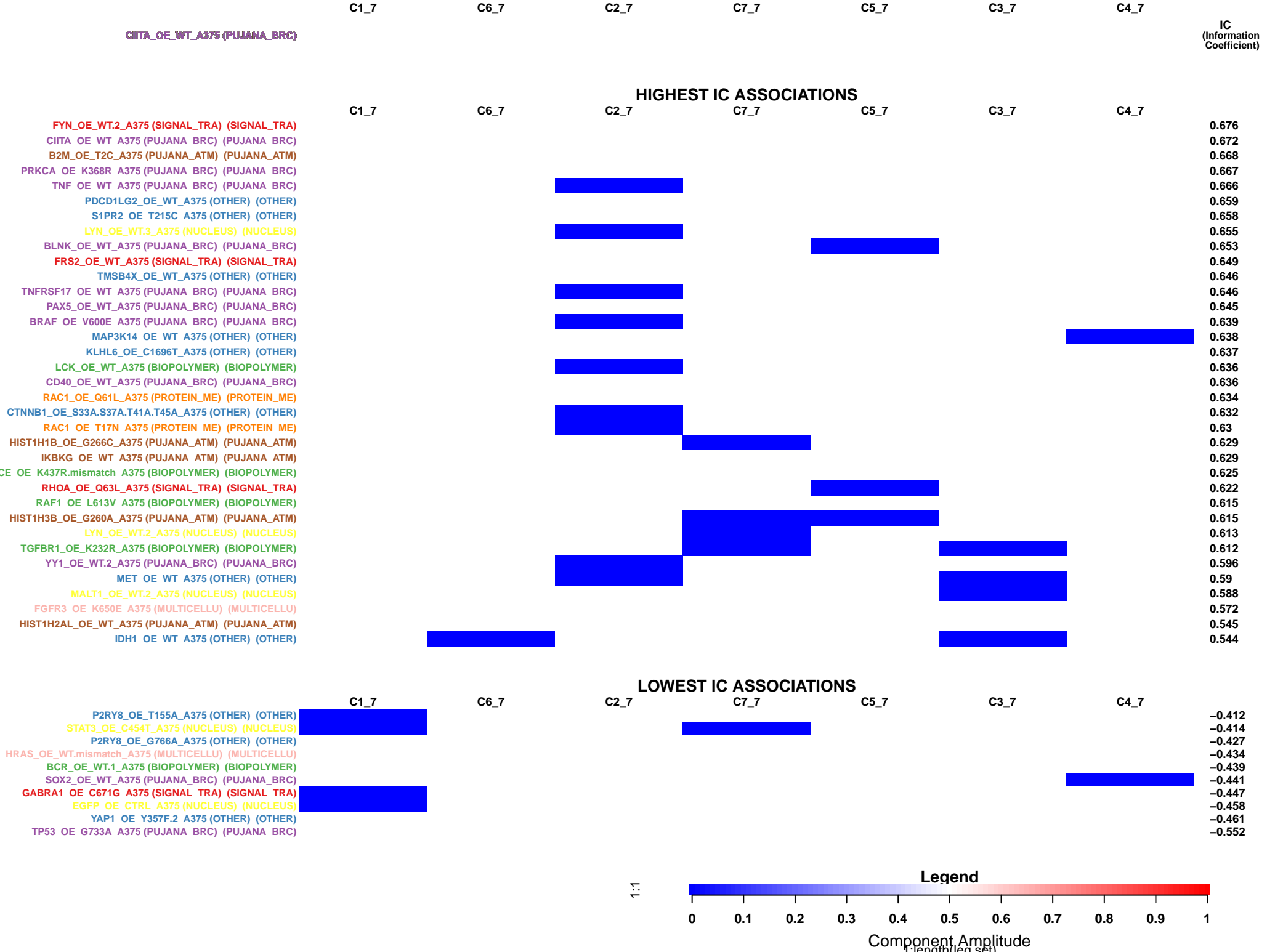
PREX2\_OE\_WT.1\_A375 (OTHER)



PASD1\_OE\_G1032T\_A375 (OTHER)

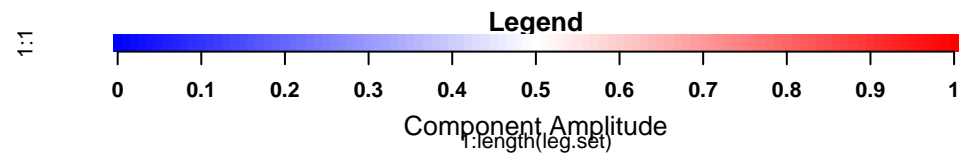
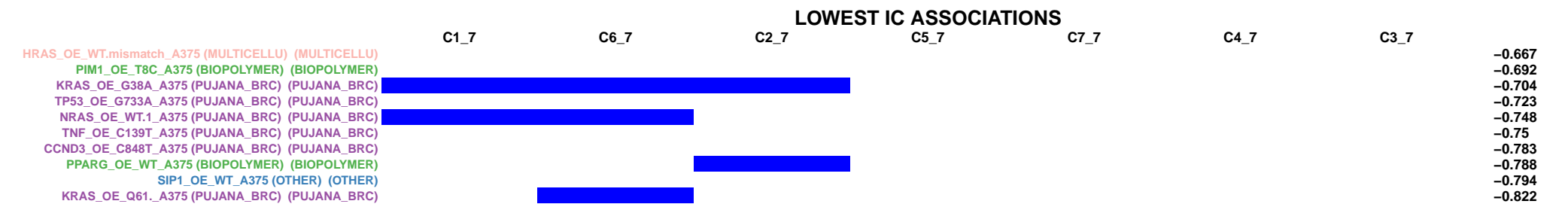
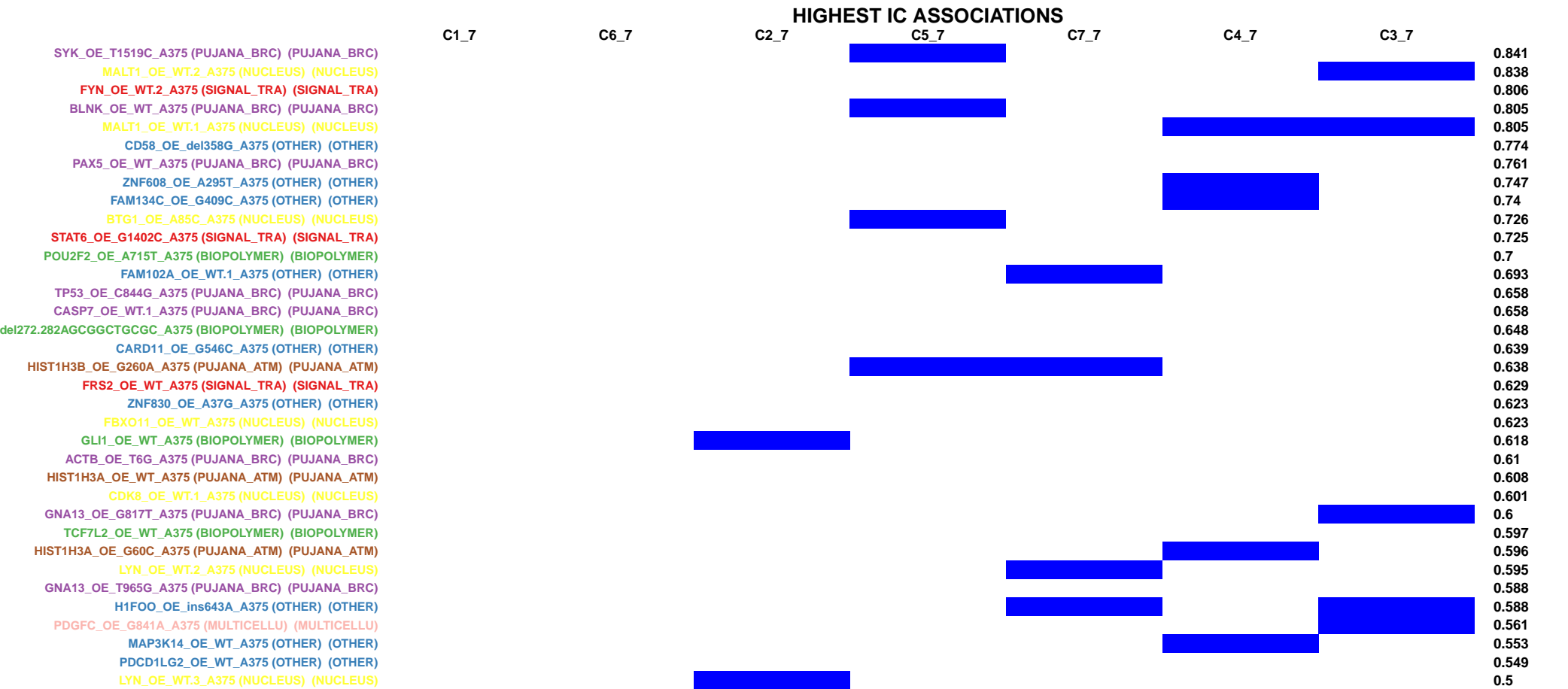


CIITA\_OE\_WT\_A375 (PUJANA\_BRC)

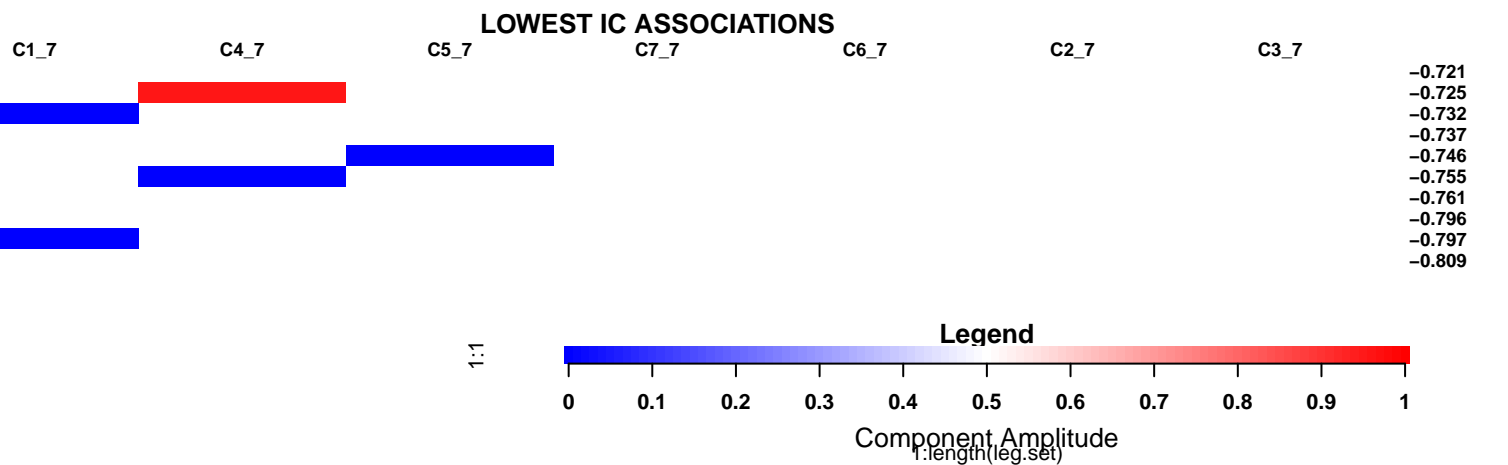
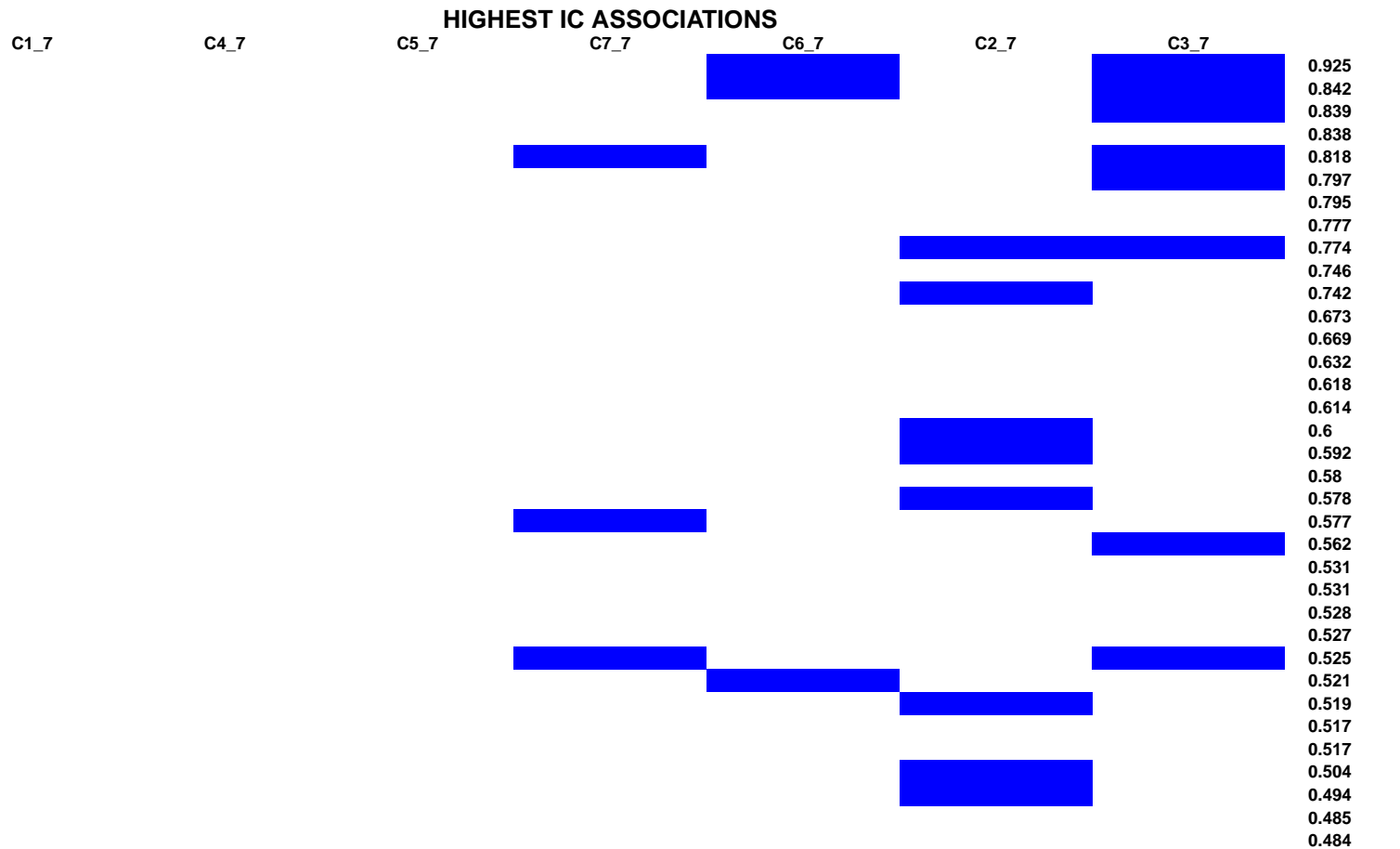




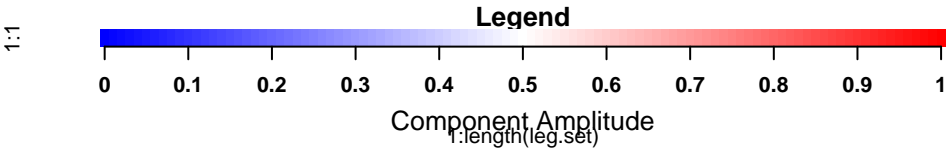
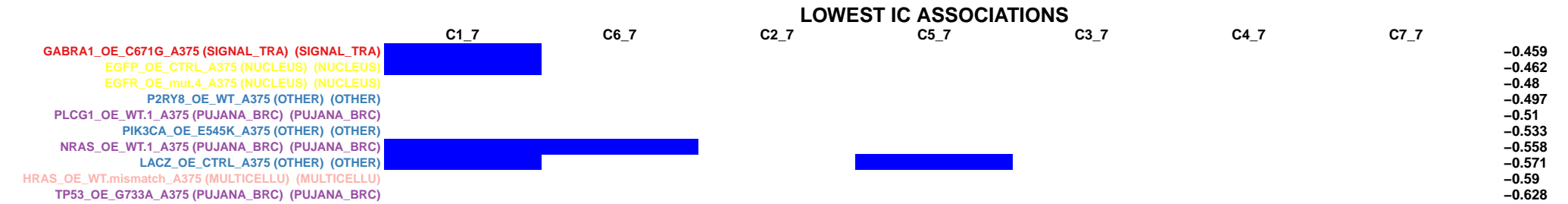
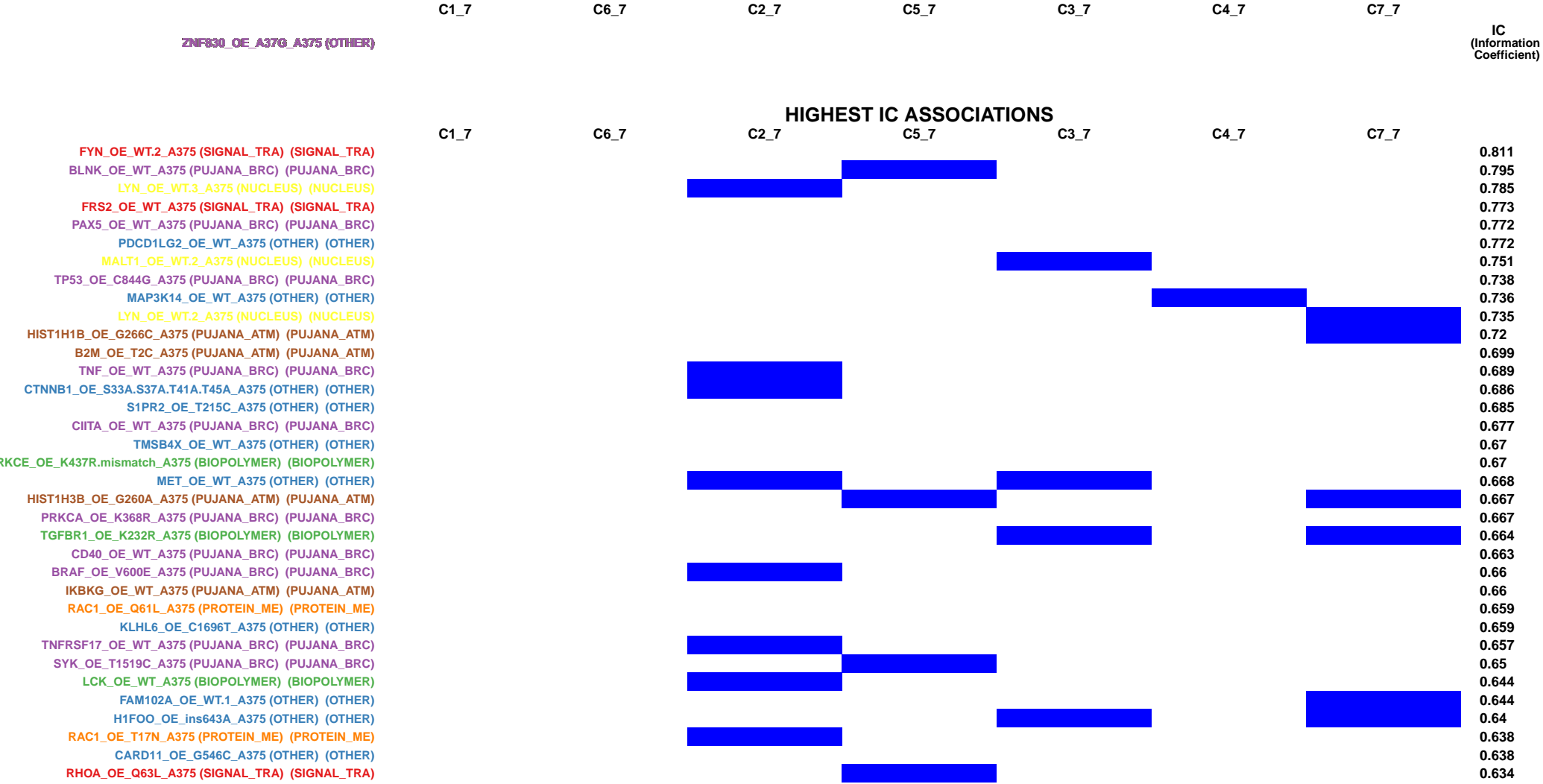
C1_7	C6_7	C2_7	C5_7	C7_7	C4_7	C3_7	IC (Information Coefficient)
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C1_7	C4_7	C5_7	C7_7	C6_7	C2_7	C3_7	IC (Information Coefficient)



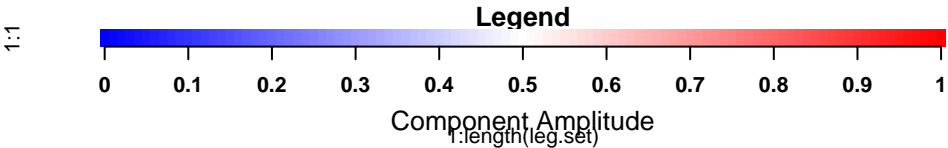
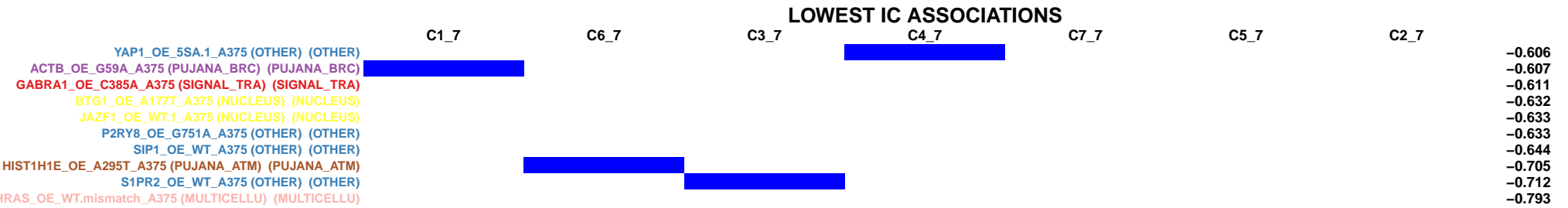
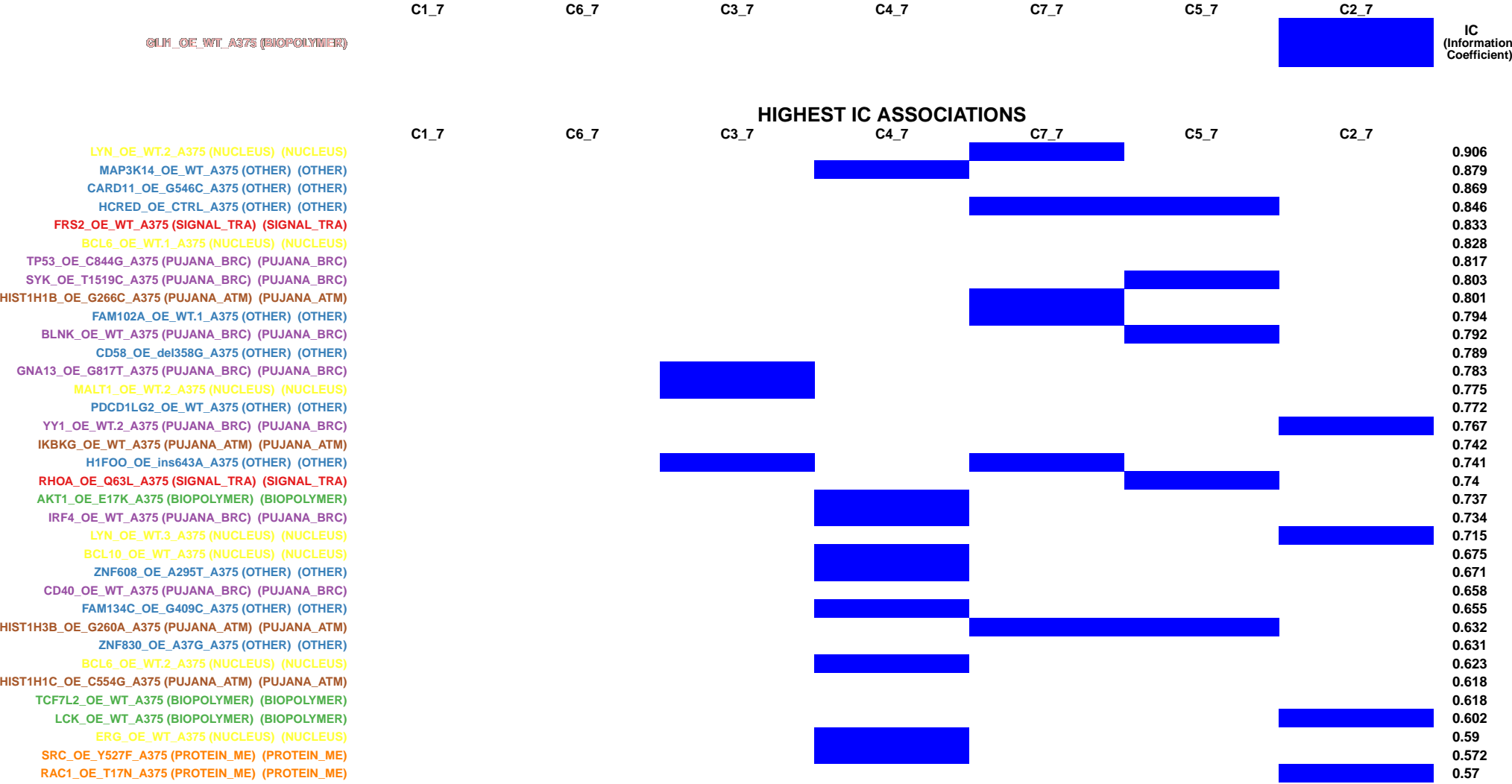
ZNF830\_OE\_A37G\_A375 (OTHER)



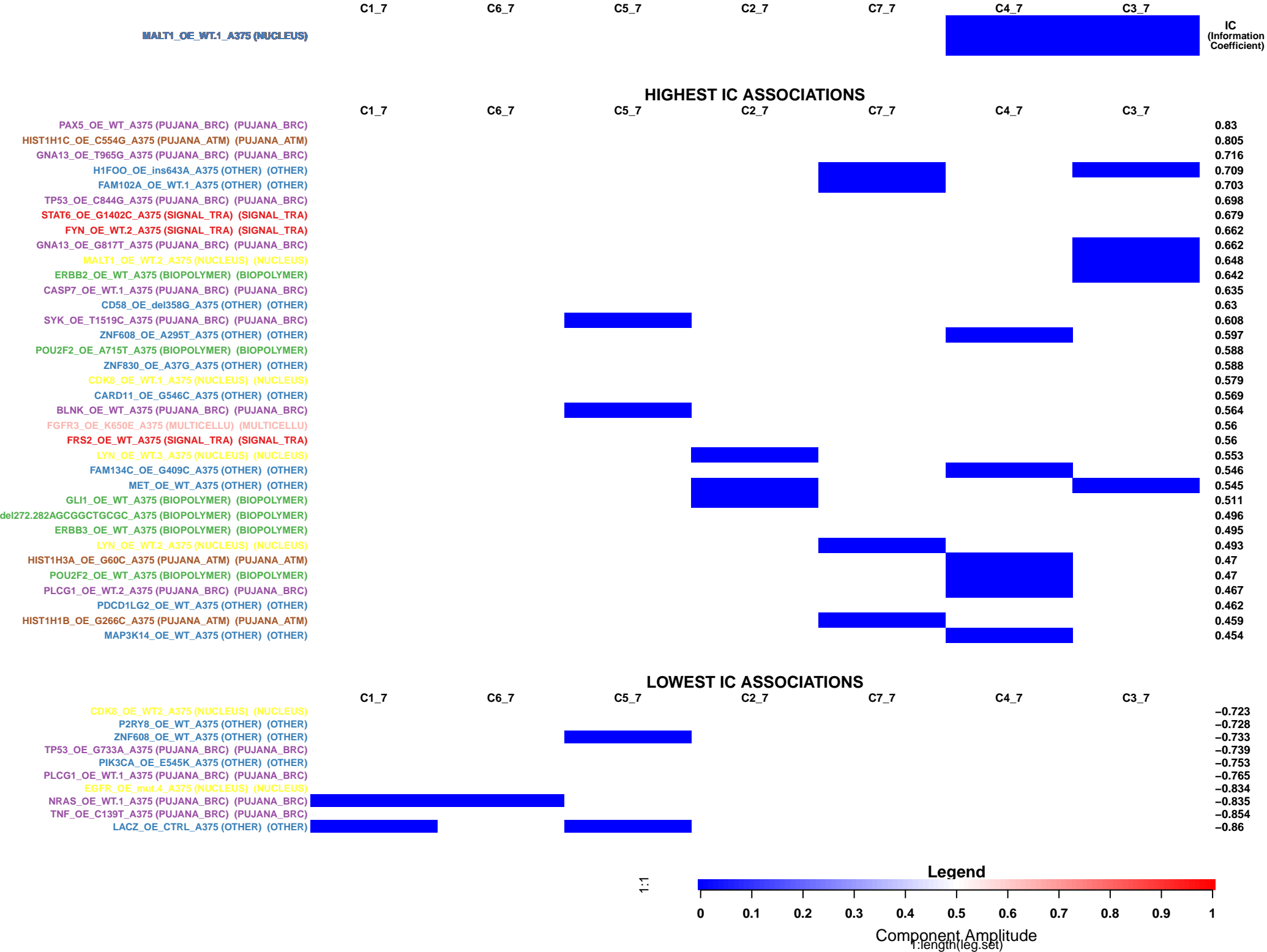
GLI1\_OE\_WT\_A375 (BIOPOLYMER)

GLI1\_OE\_WT\_A375 (BIOPOLYMER)

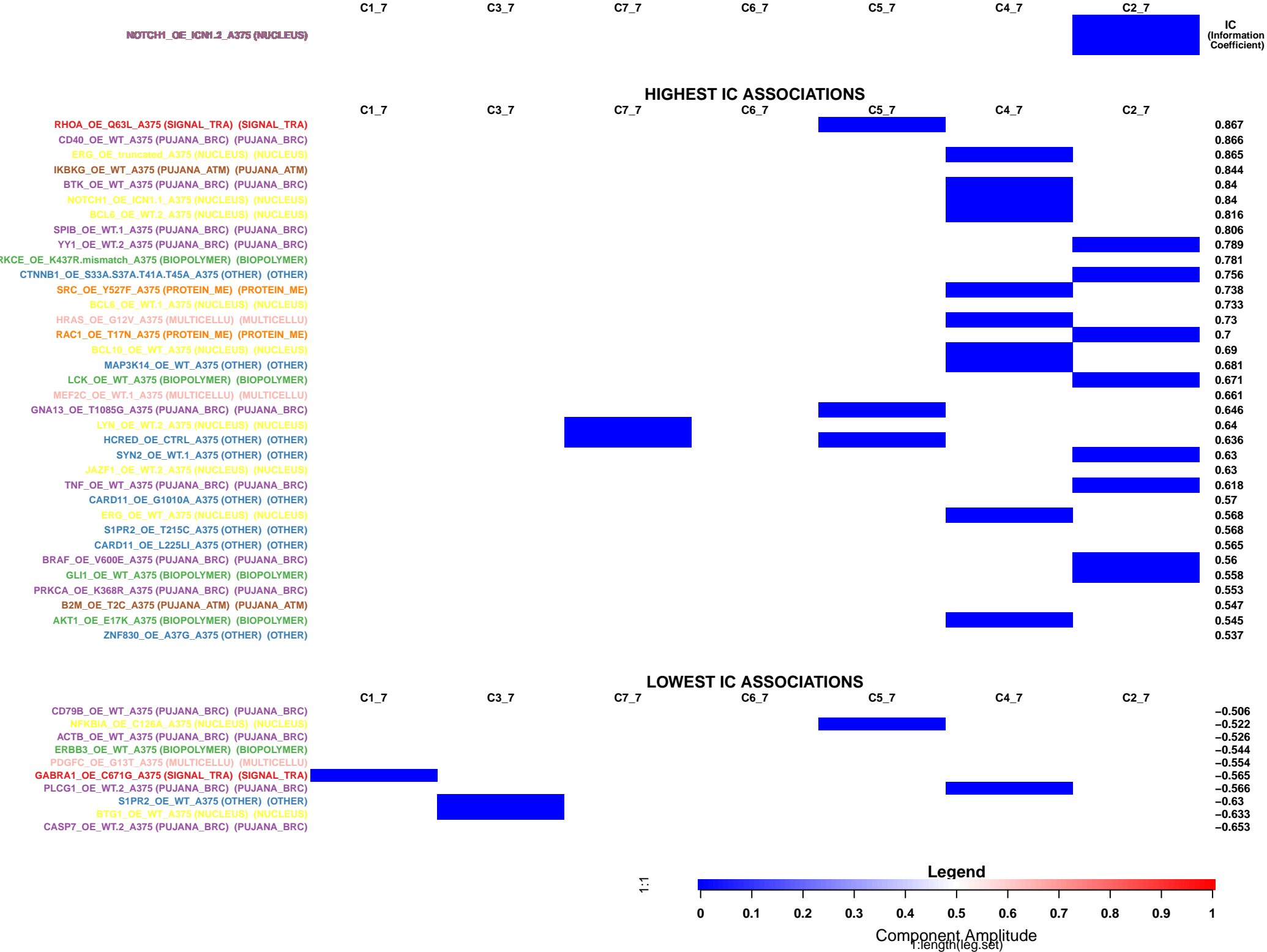
IC  
(Information  
Coefficient)



MALT1\_OE\_WT.1\_A375 (NUCLEUS)



NOTCH1\_OE\_ICN1.2\_A375 (NUCLEUS)

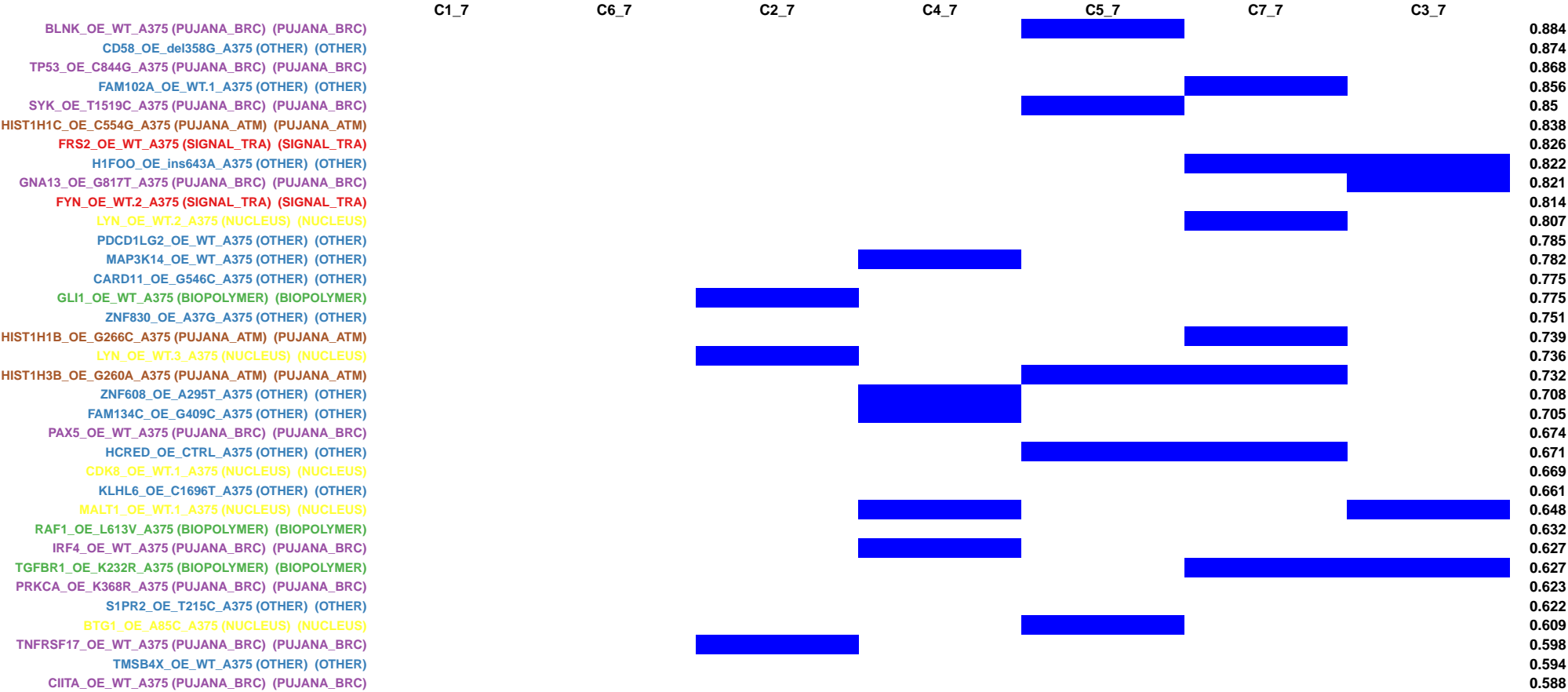


MALT1\_OE\_WT.2\_A375 (NUCLEUS)

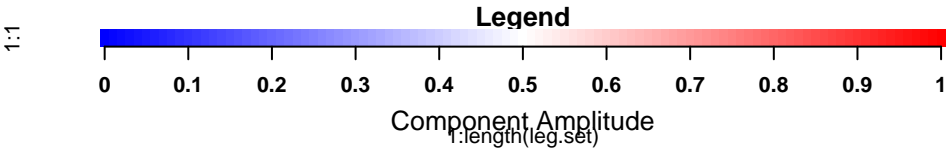
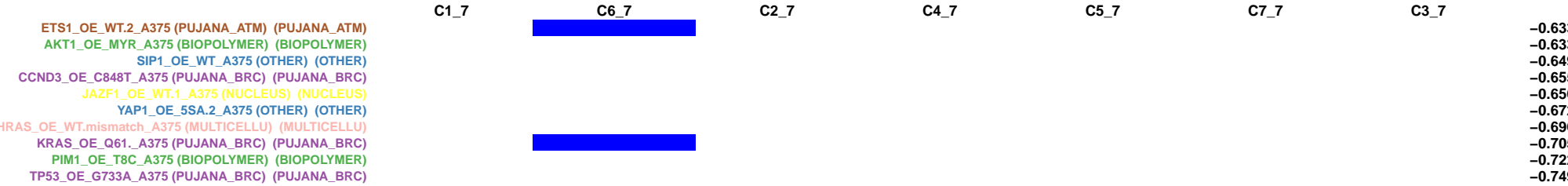
MALT1\_OE\_WT.2\_A375 (NUCLEUS)

IC  
(Information  
Coefficient)

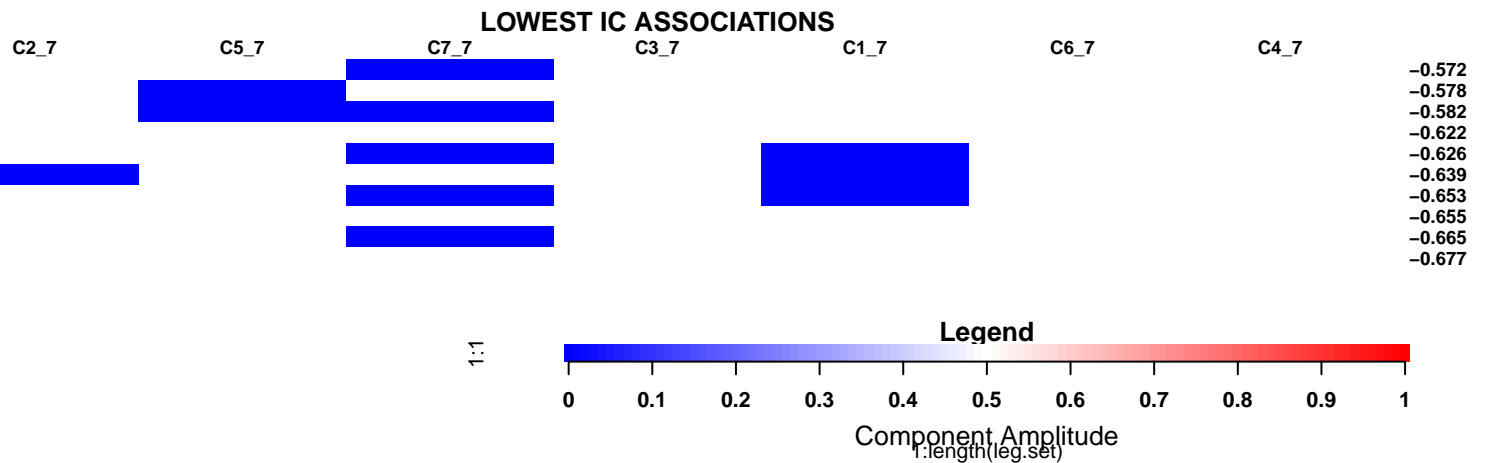
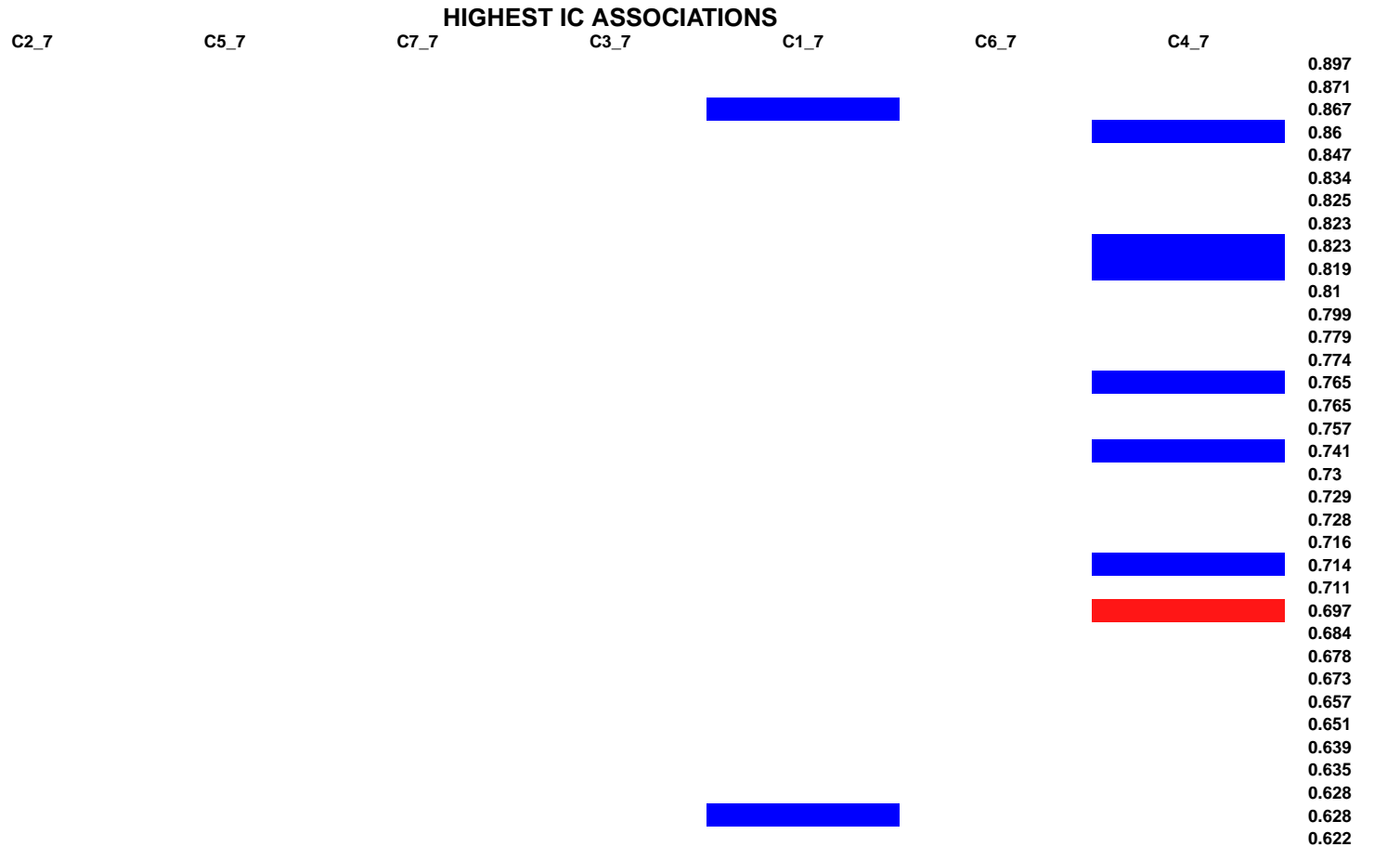
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS

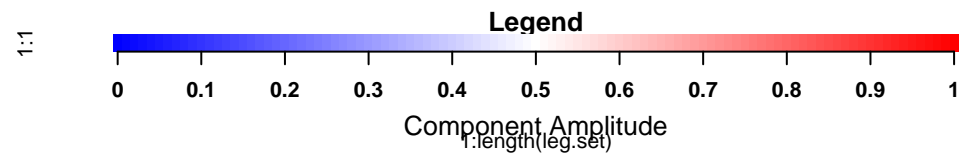
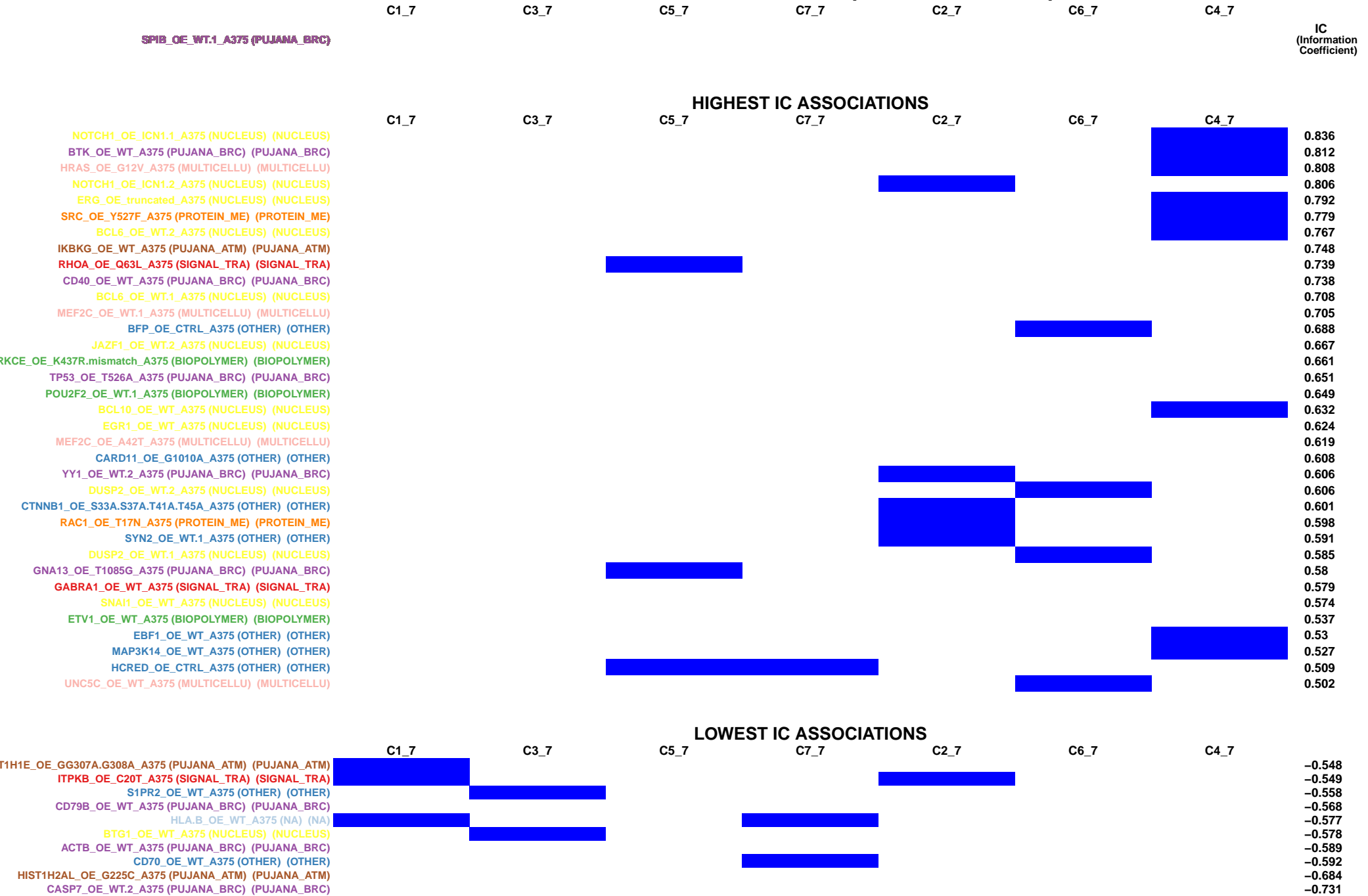


C2_7	C5_7	C7_7	C3_7	C1_7	C6_7	C4_7	IC (Information Coefficient)
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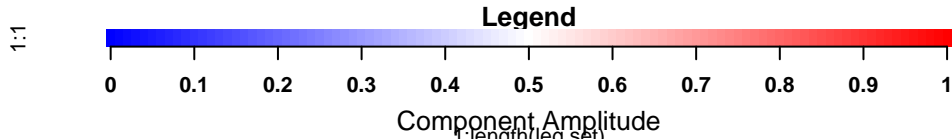
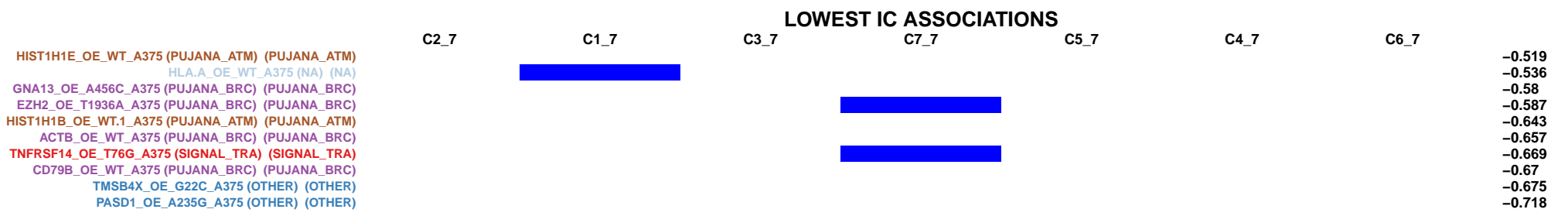
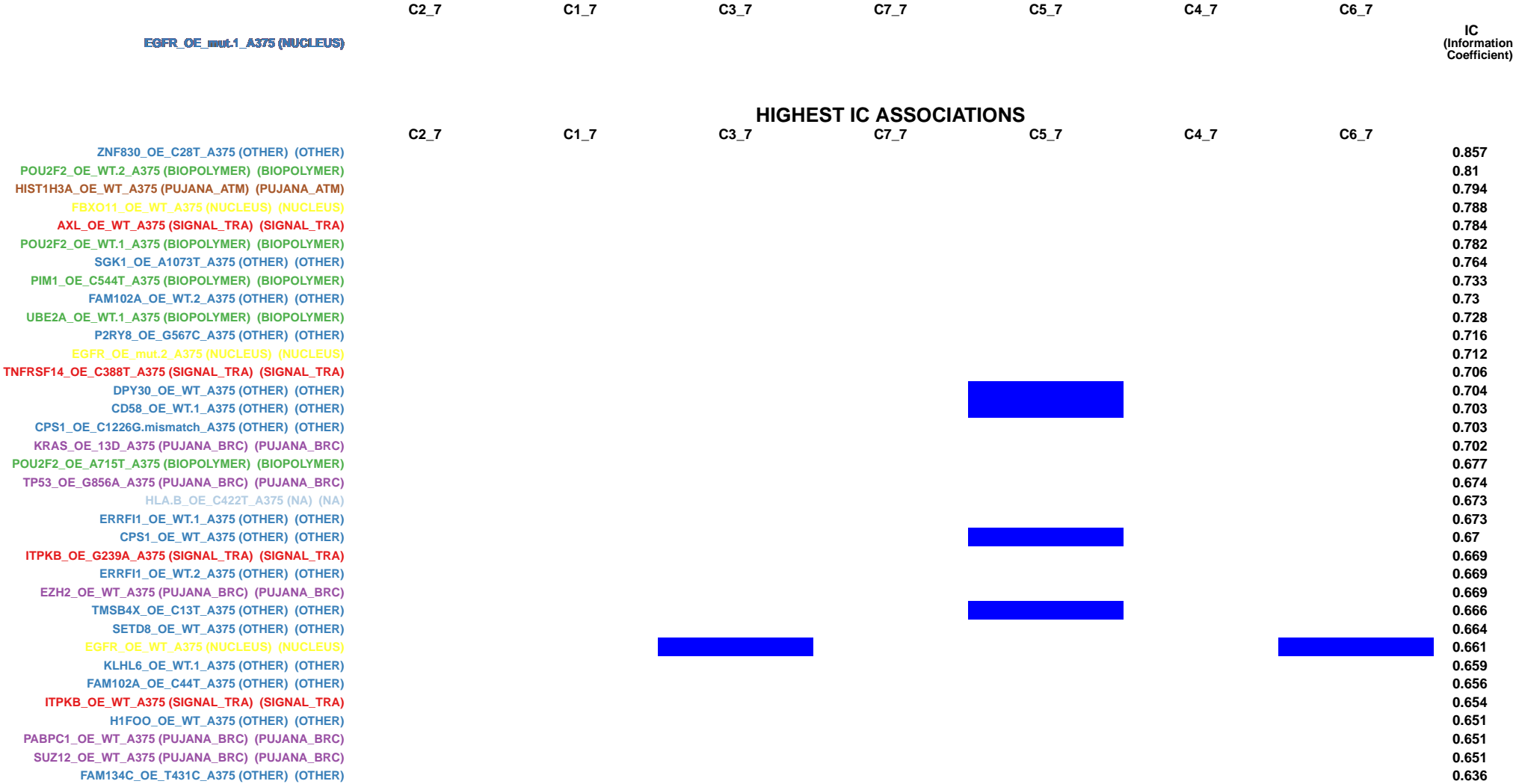




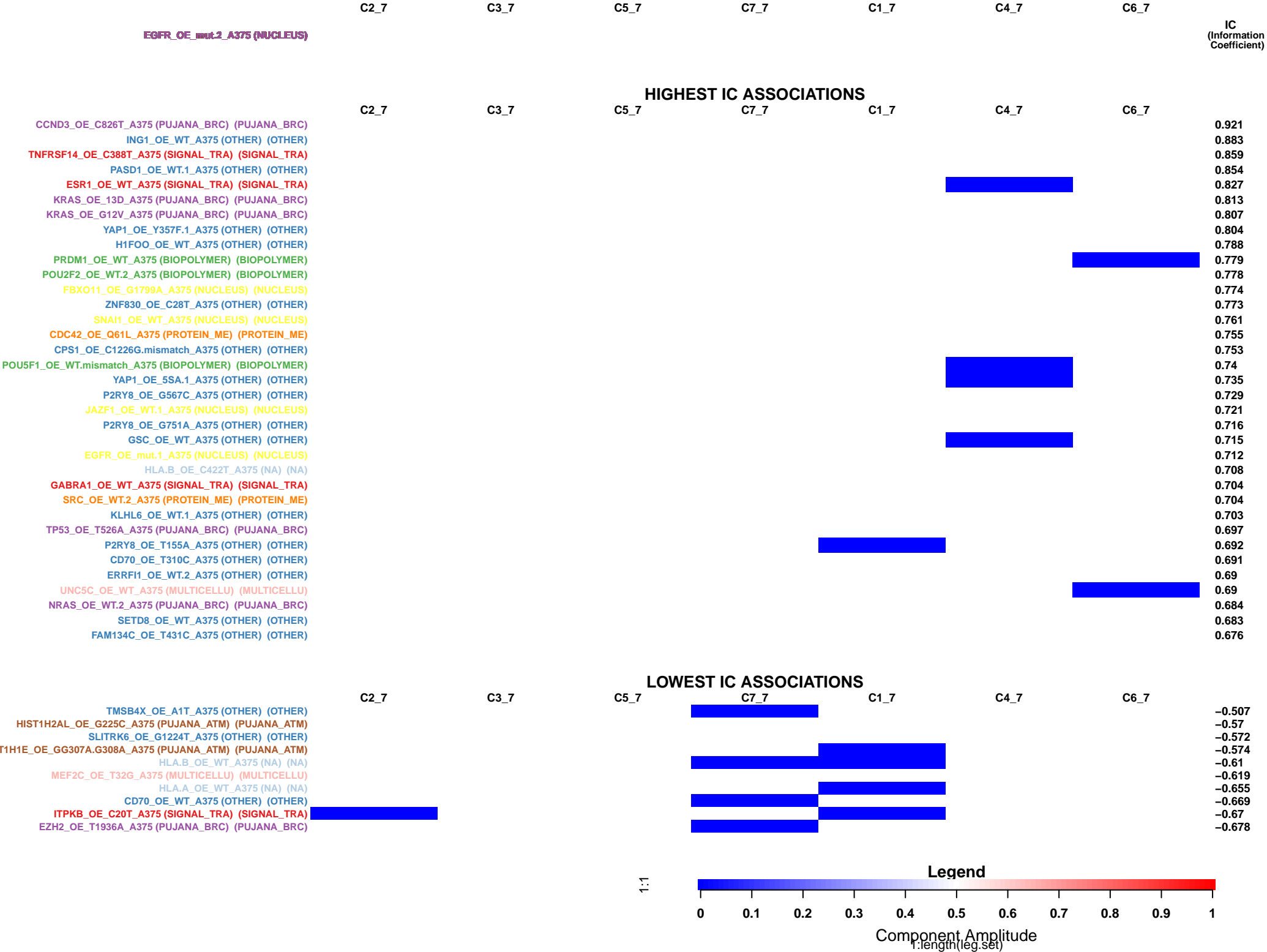
# SPIB\_OE\_WT.1\_A375 (PUJANA\_BRC)



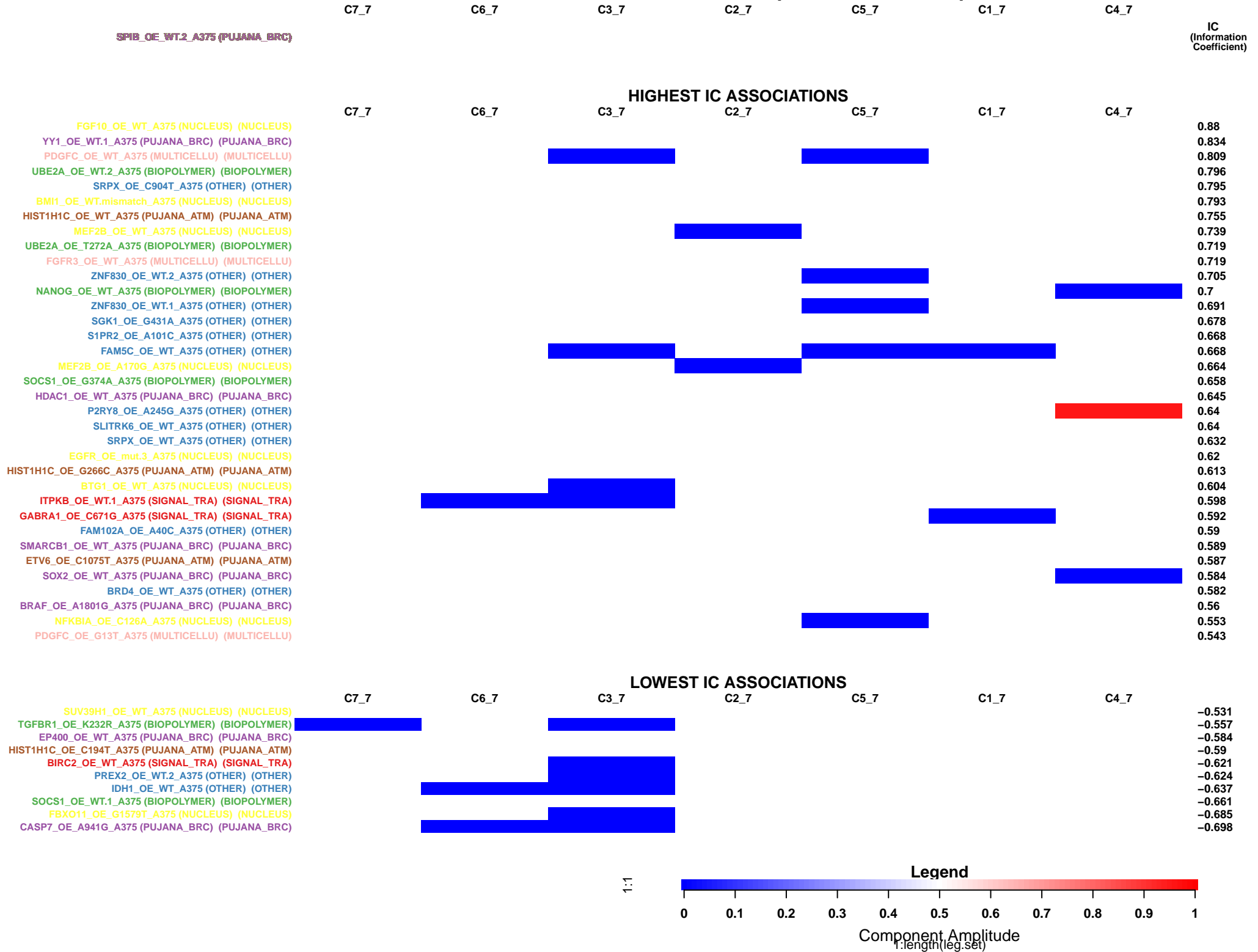
EGFR\_OE\_mut.1\_A375 (NUCLEUS)



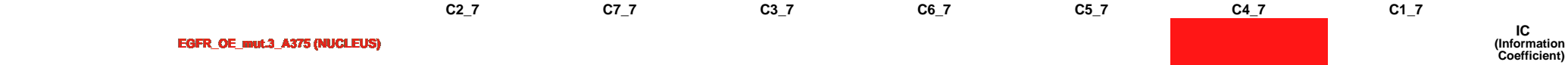
EGFR\_OE\_mut.2\_A375 (NUCLEUS)



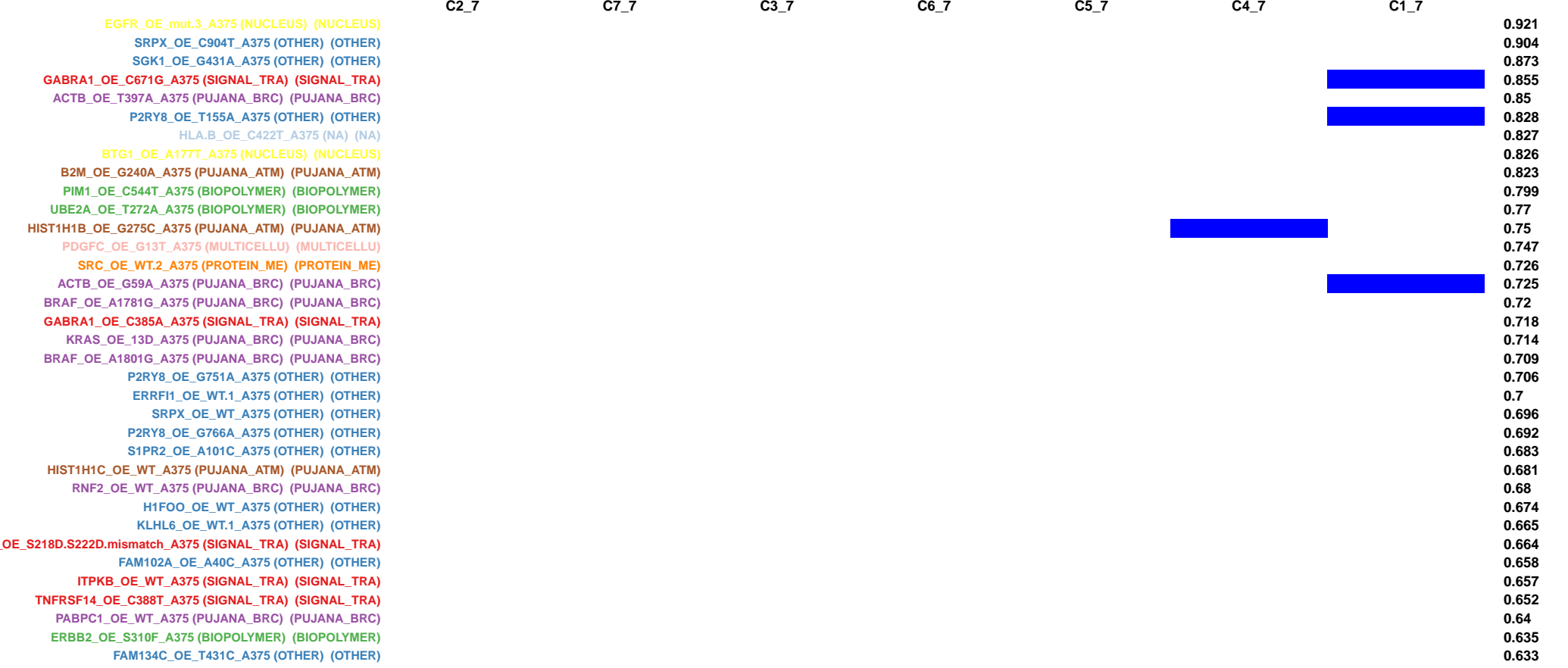
# SPIB\_OE\_WT.2\_A375 (PUJANA\_BRC)



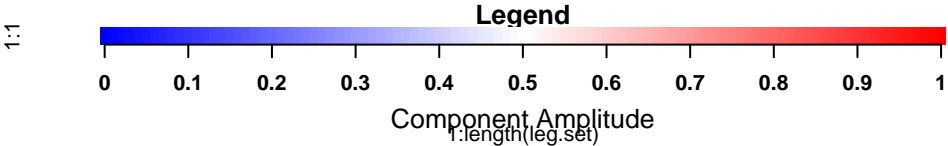
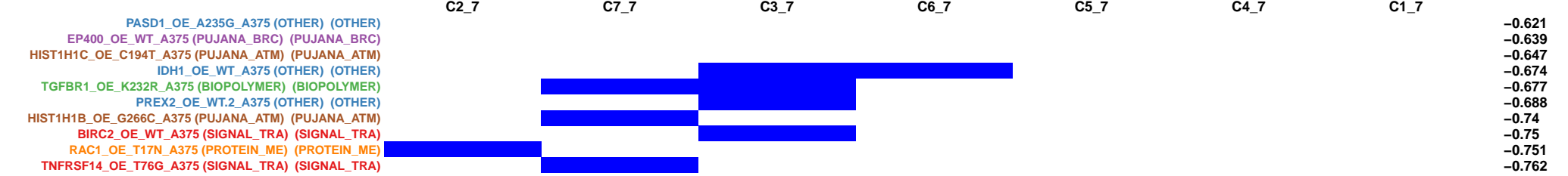
EGFR\_OE\_mut.3\_A375 (NUCLEUS)



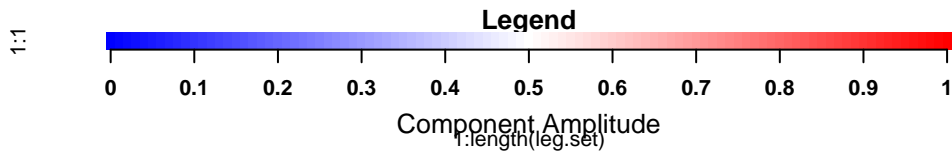
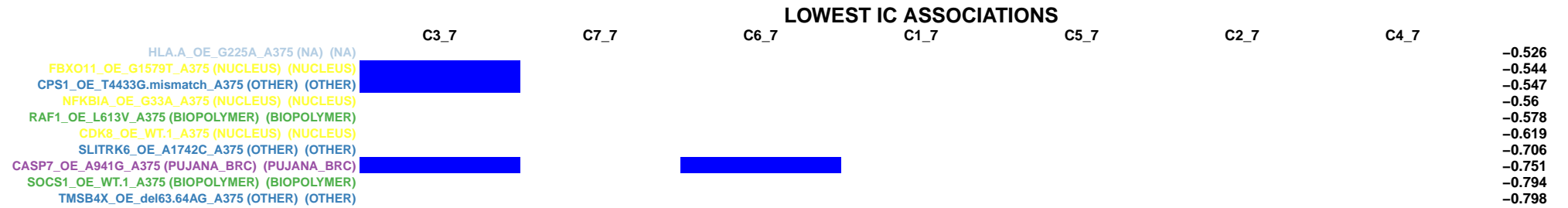
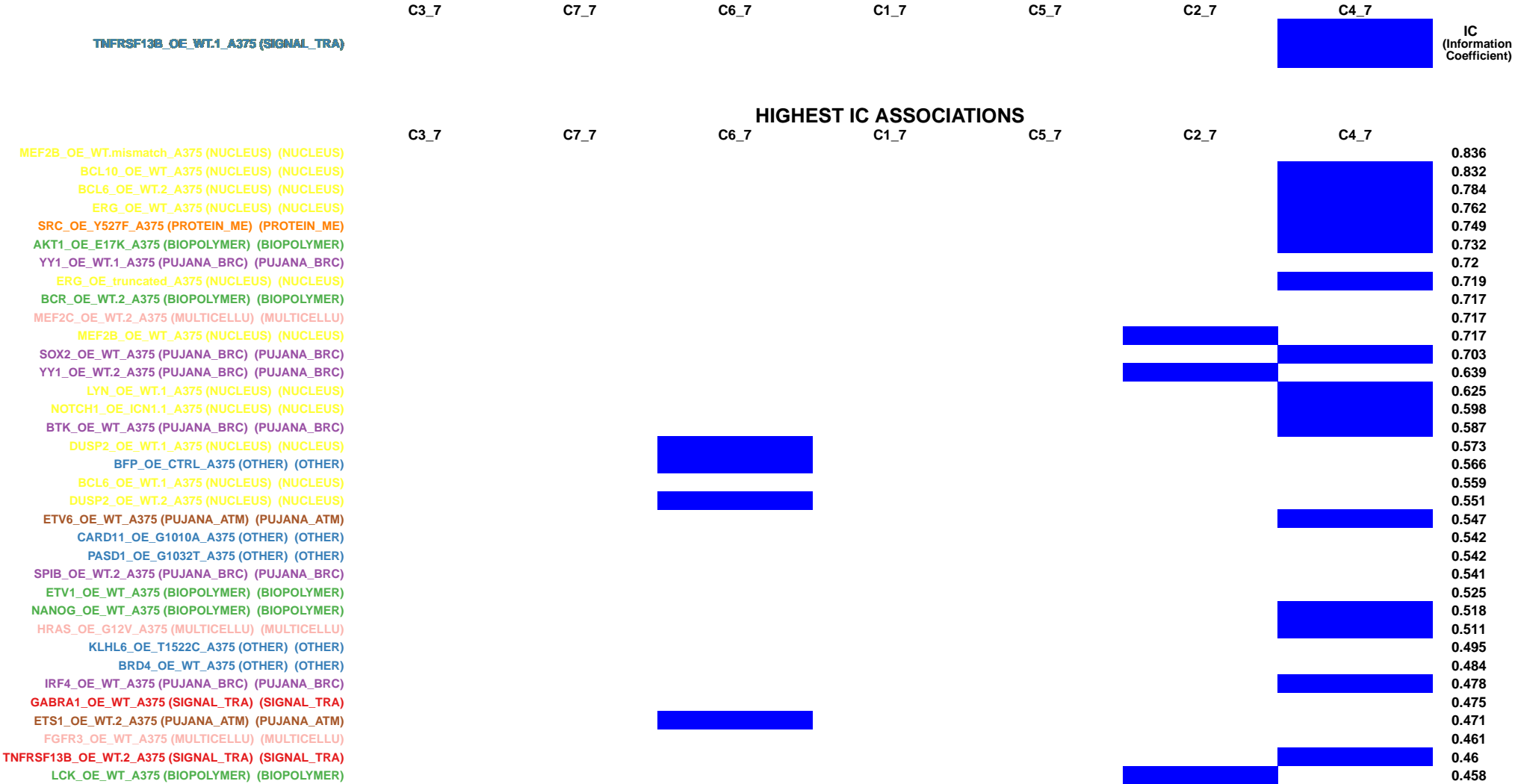
HIGHEST IC ASSOCIATIONS



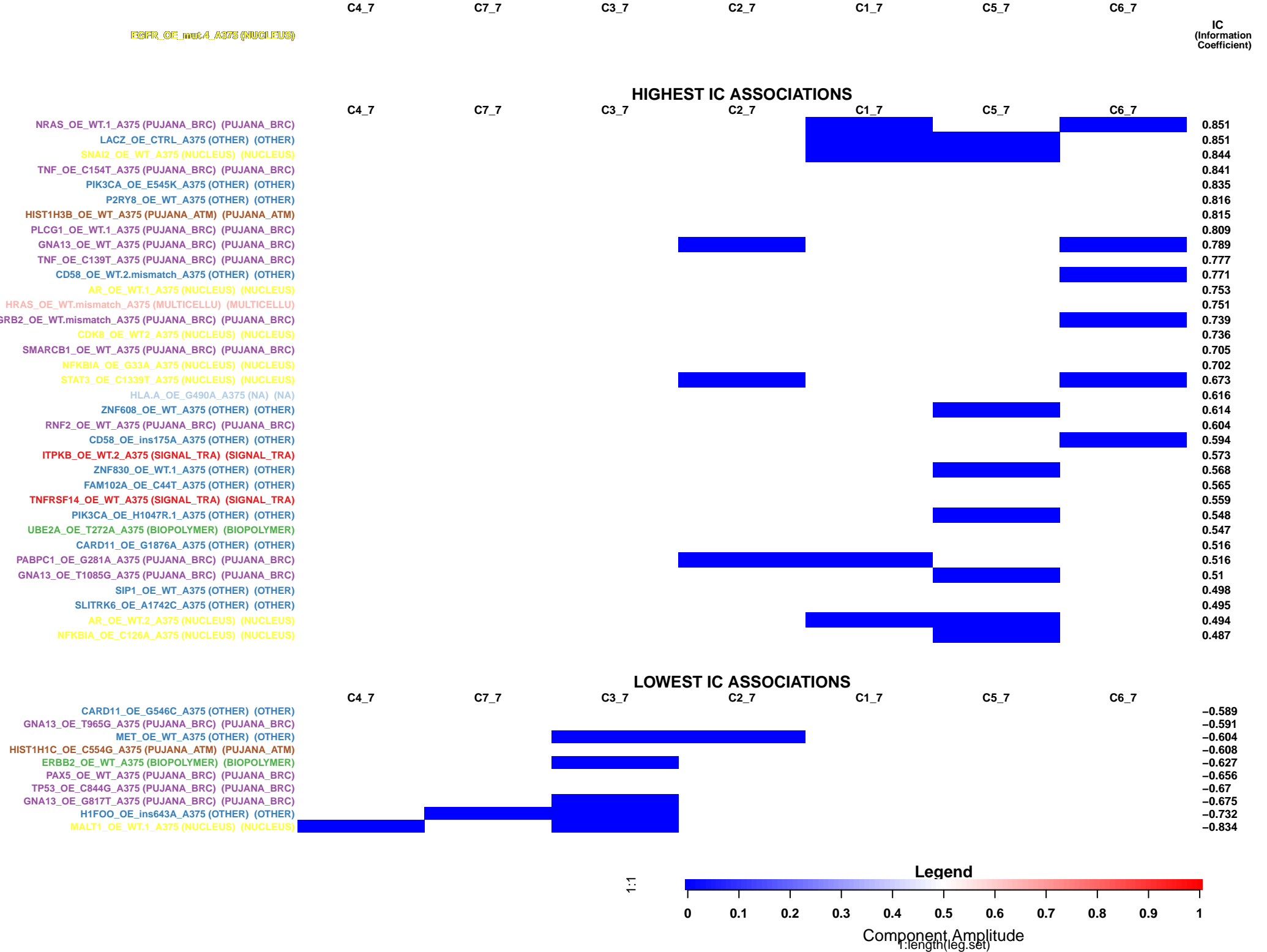
LOWEST IC ASSOCIATIONS



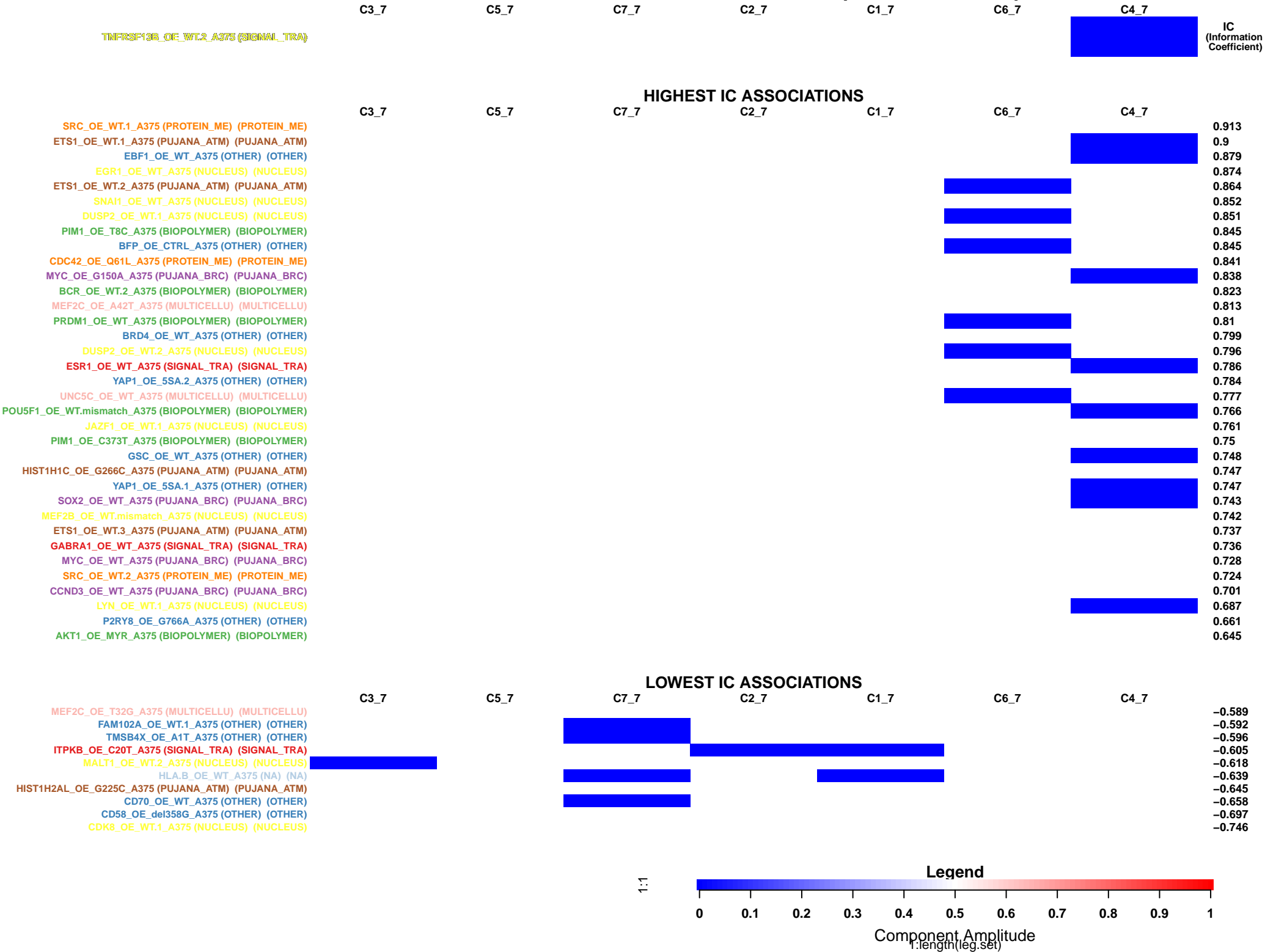
TNFRSF13B\_OE\_WT.1\_A375 (SIGNAL\_TRA)



EGFR\_OE\_mut.4\_A375 (NUCLEUS)

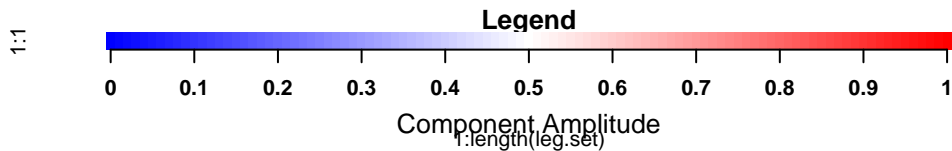
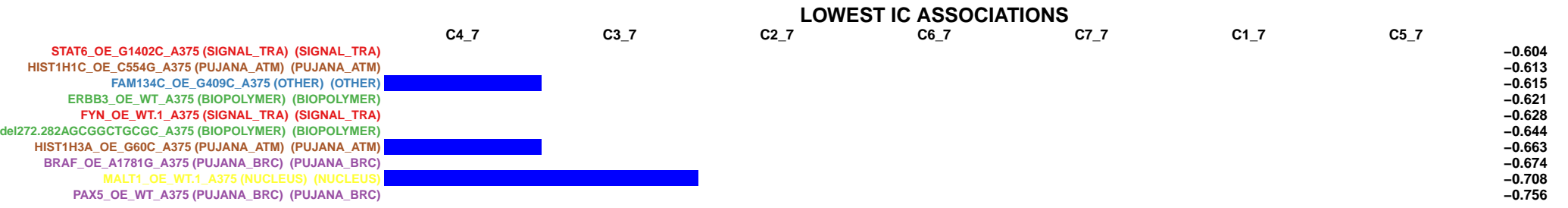
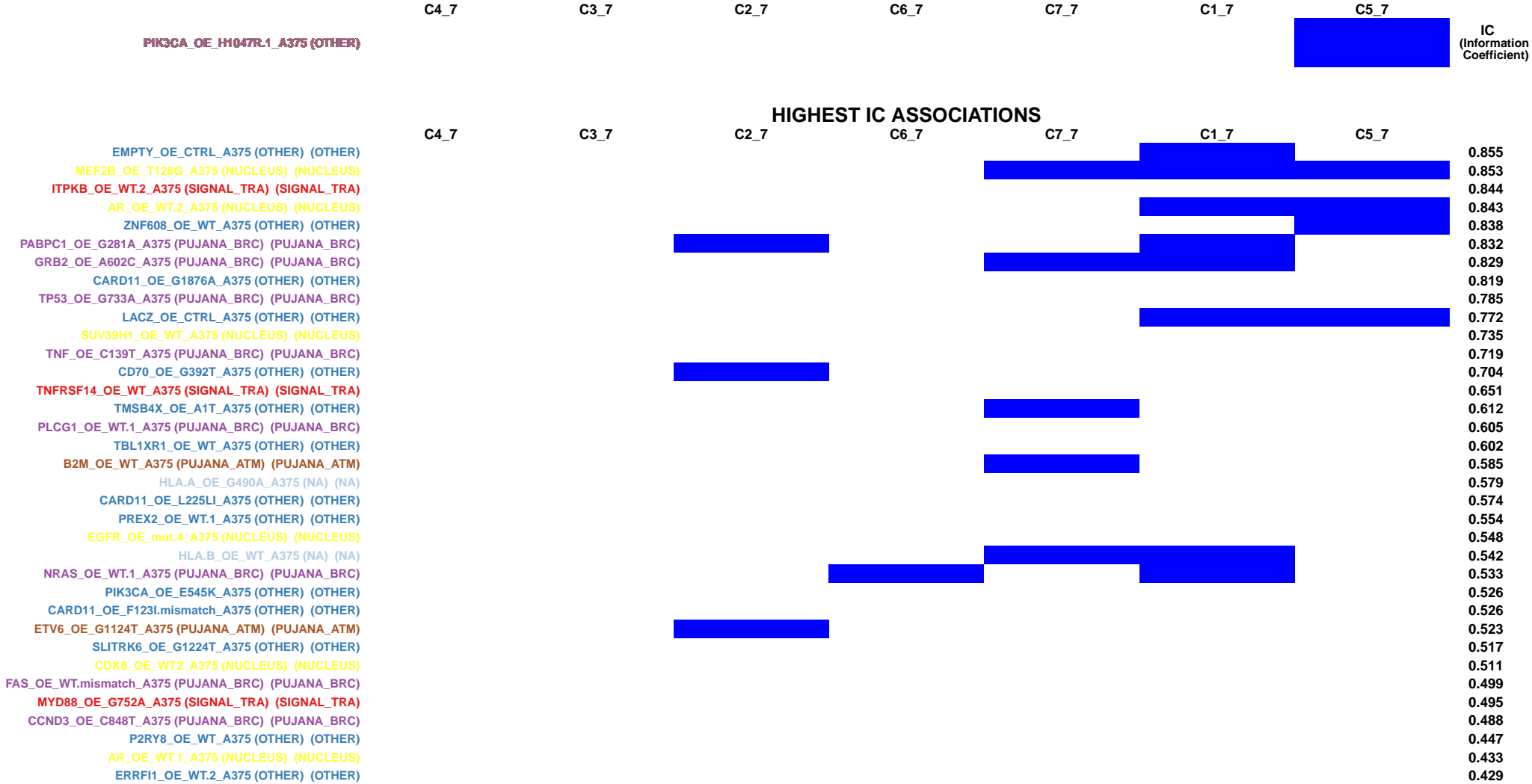


TNFRSF13B\_OE\_WT.2\_A375 (SIGNAL\_TRA)

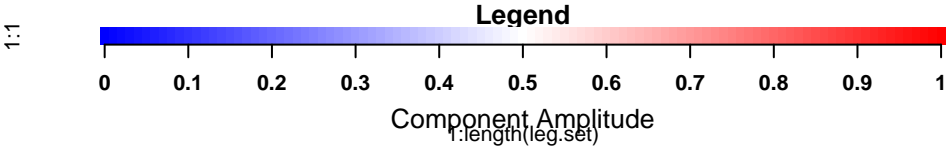
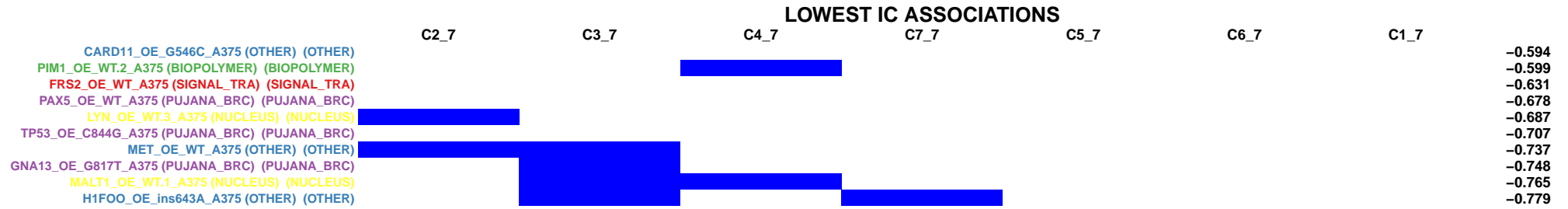
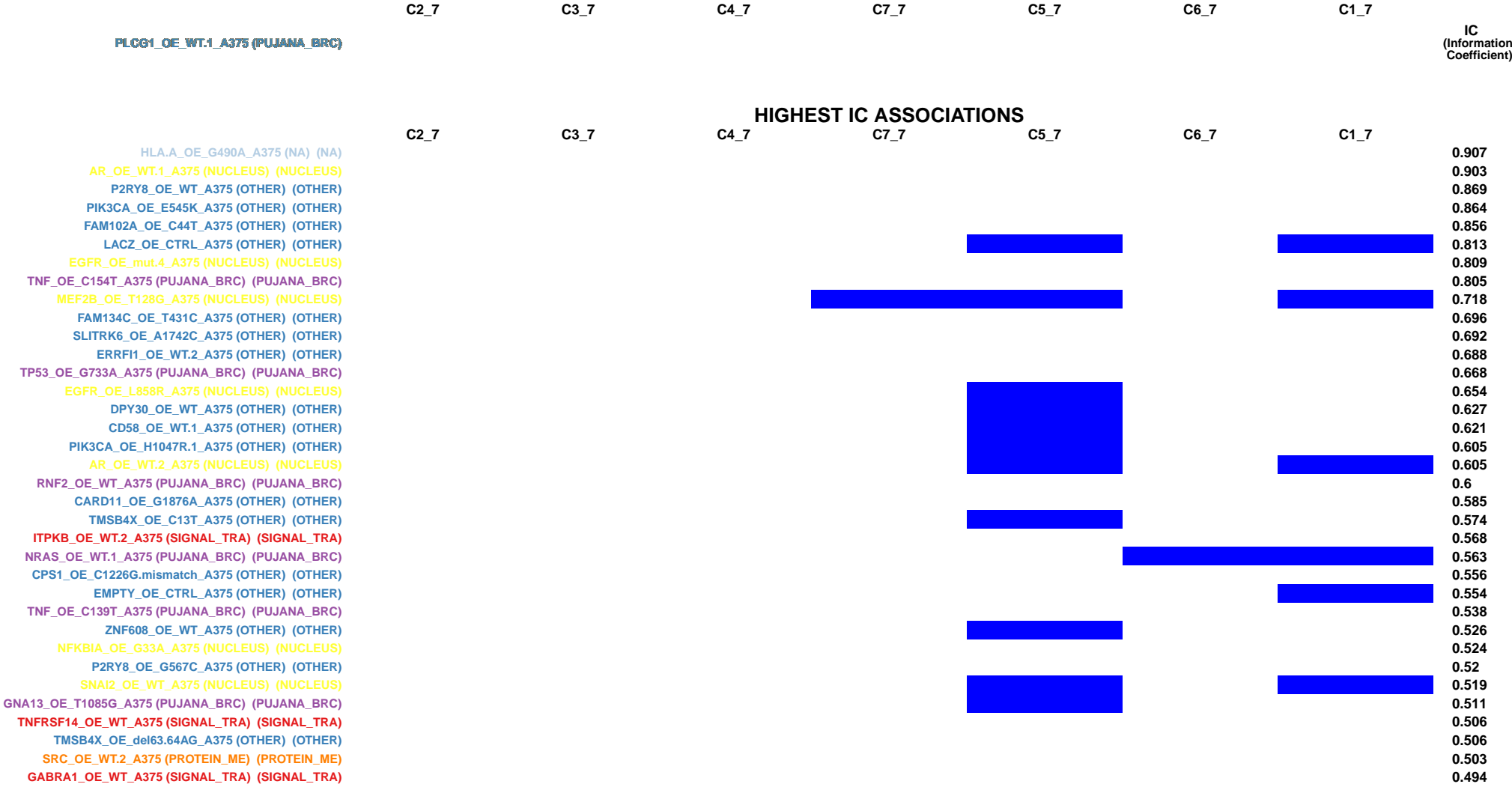




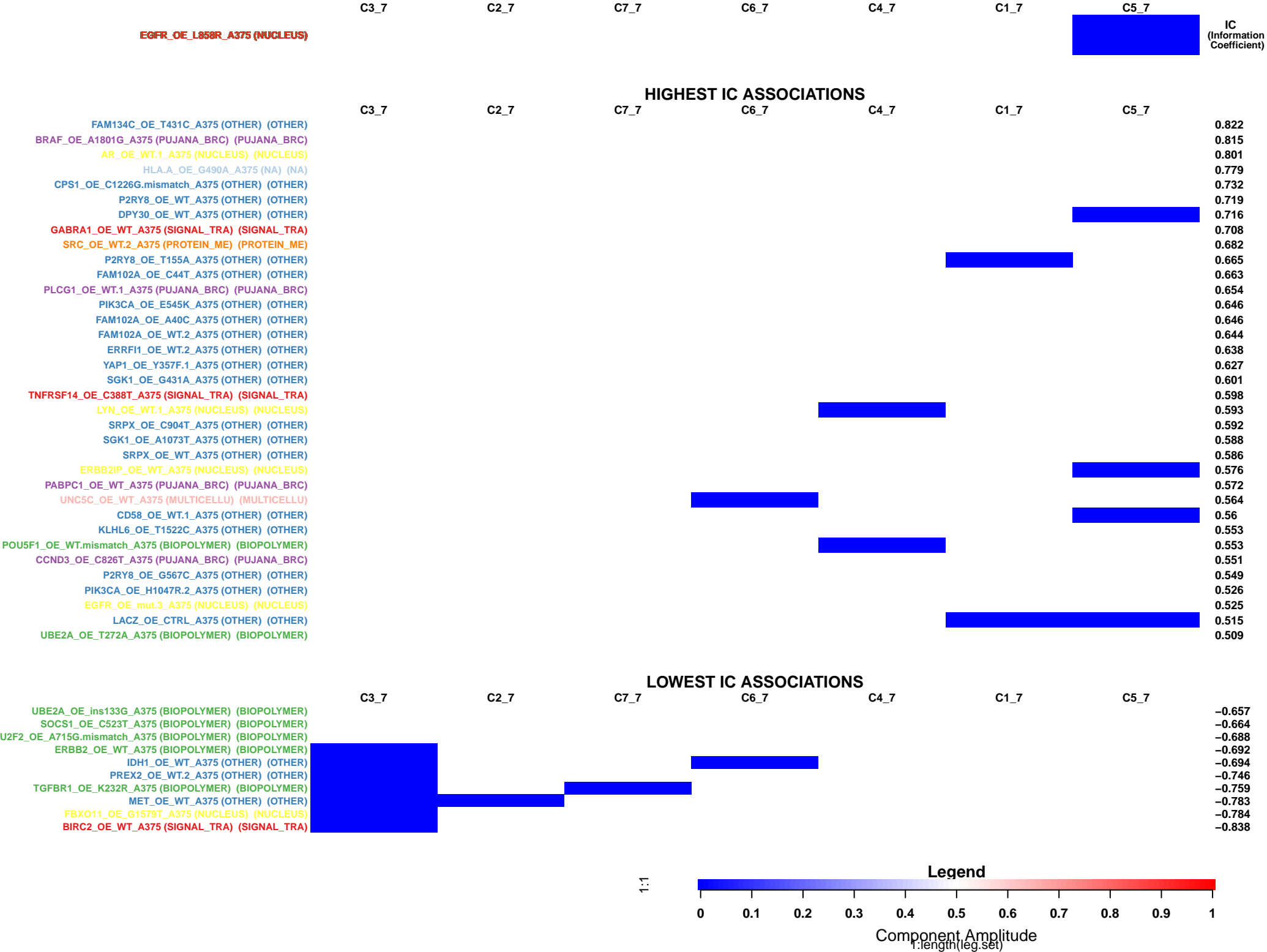
PIK3CA\_OE\_H1047R.1\_A375 (OTHER)



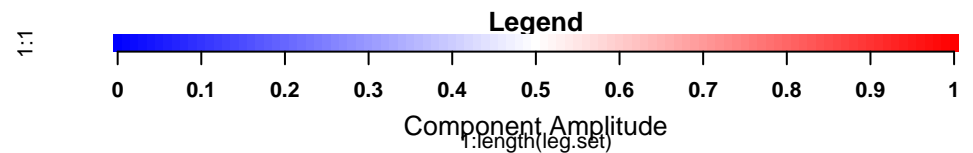
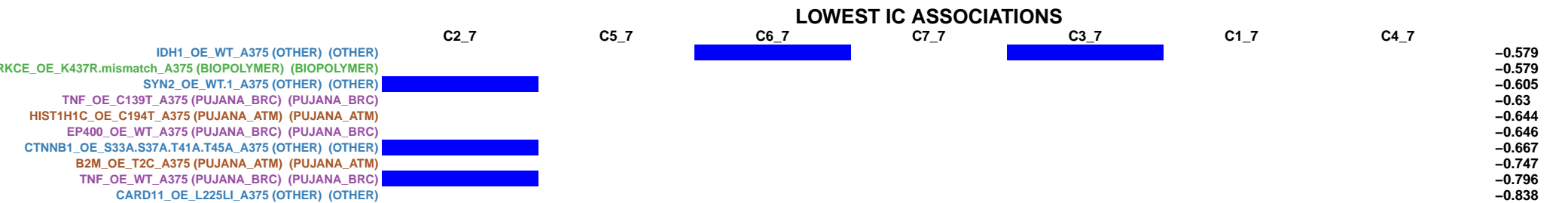
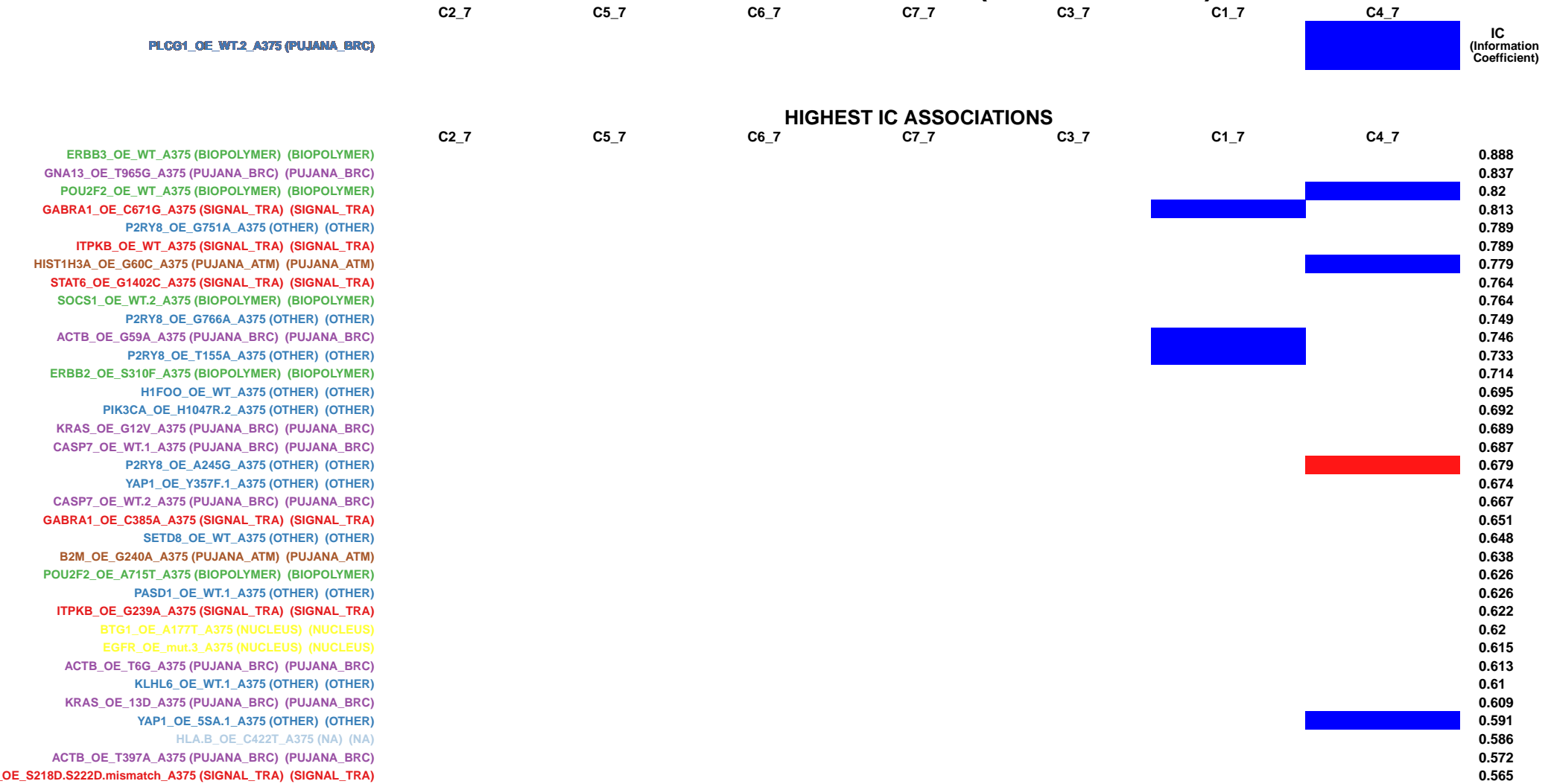
PLCG1\_OE\_WT.1\_A375 (PUJANA\_BRC)



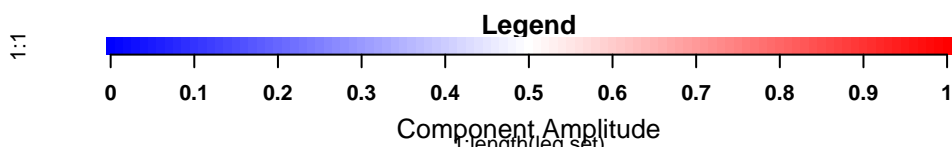
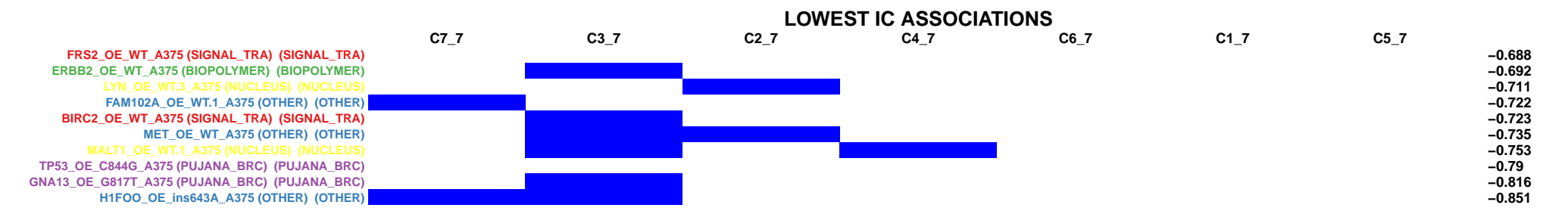
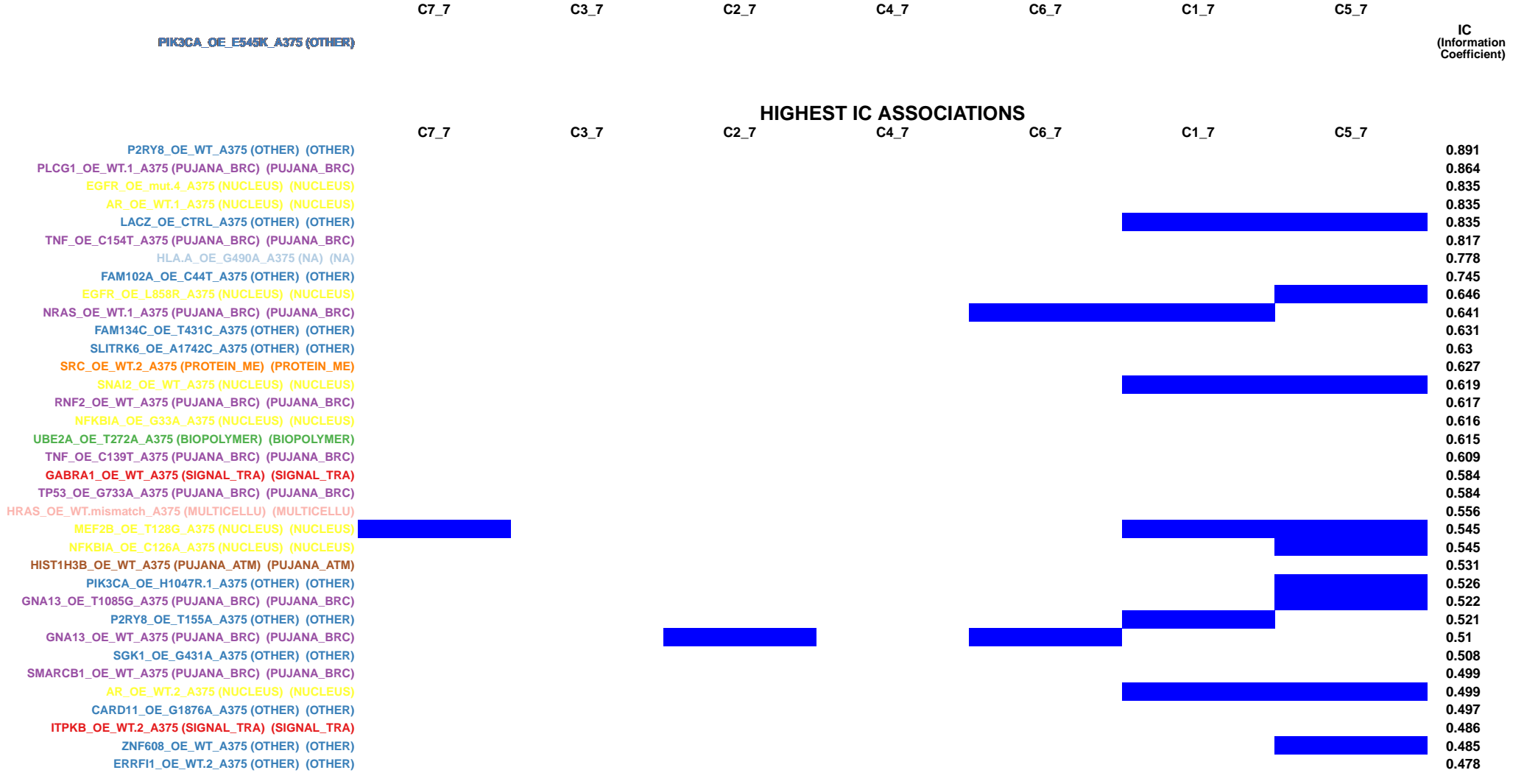
EGFR\_OE\_L858R\_A375 (NUCLEUS)



# PLCG1\_OE\_WT.2\_A375 (PUJANA\_BRC)



PIK3CA\_OE\_E545K\_A375 (OTHER)

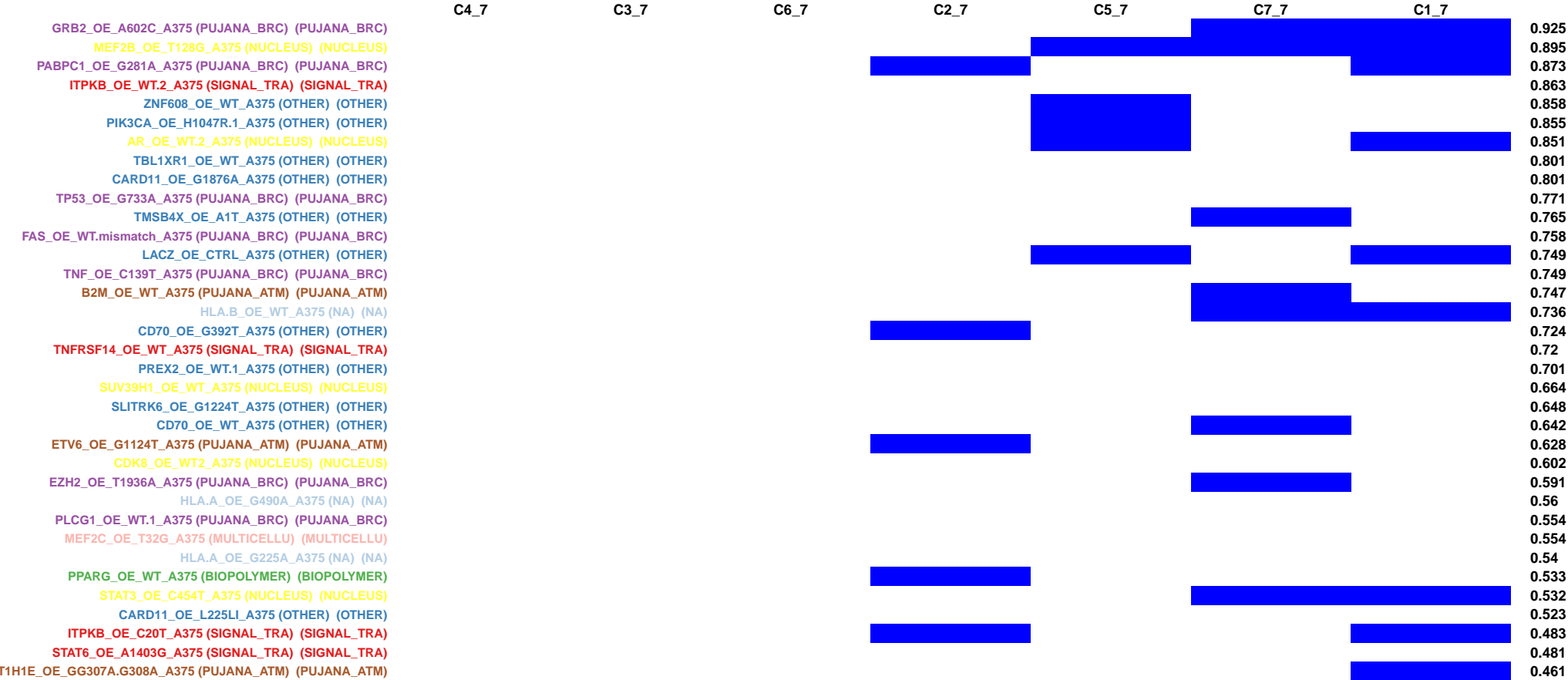


EMPTY\_OE\_CTRL\_A375 (OTHER)

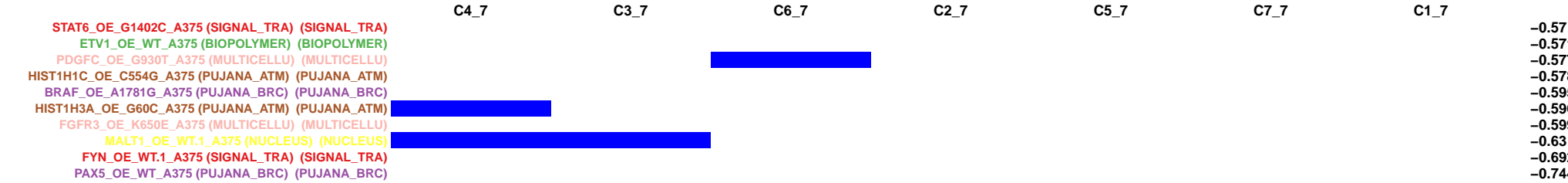
EMPTY\_OE\_CTRL\_A375 (OTHER)

IC  
(Information  
Coefficient)

HIGHEST IC ASSOCIATIONS

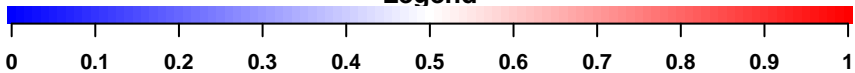


LOWEST IC ASSOCIATIONS



1:1

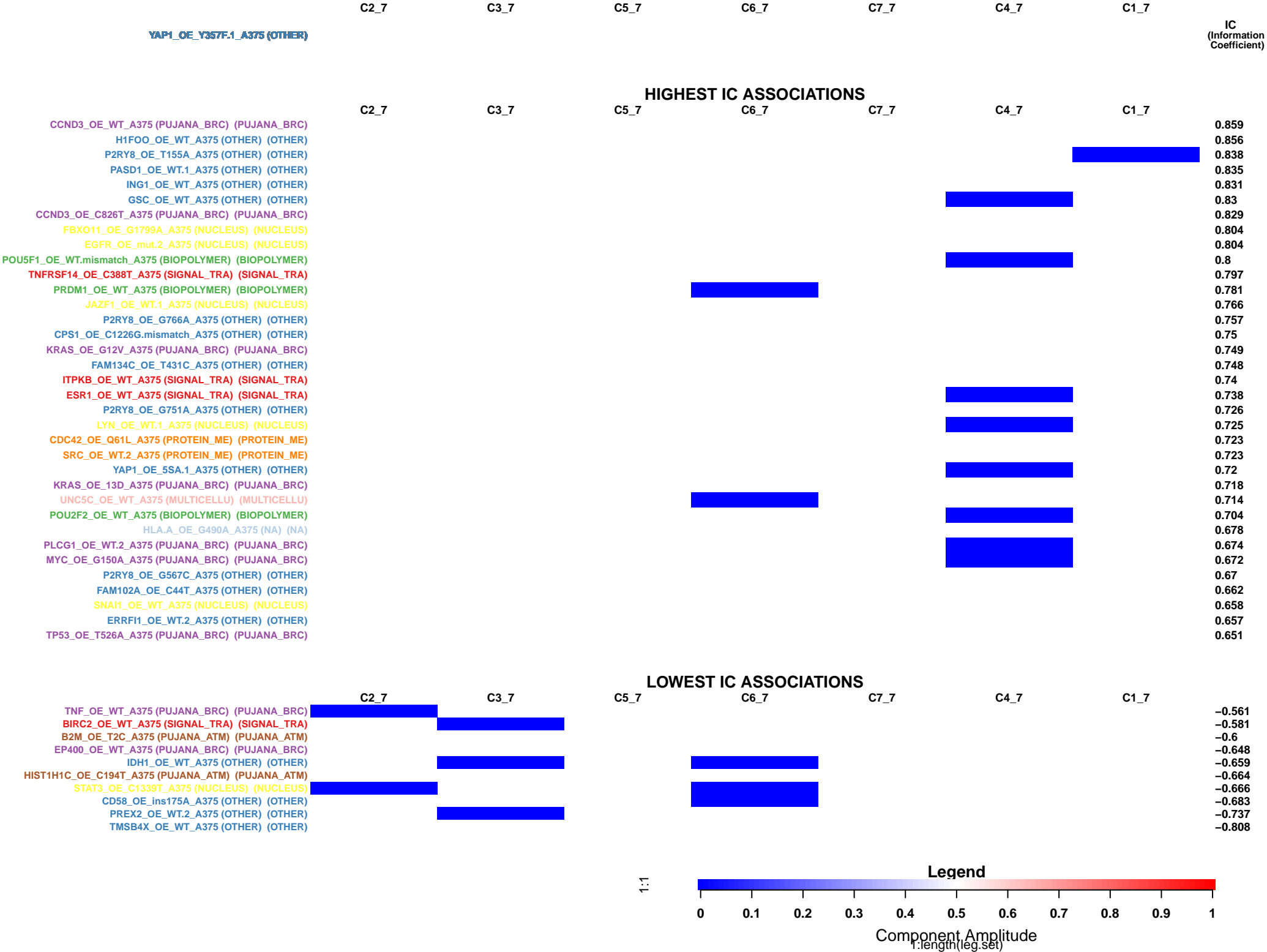
Legend



Component Amplitude

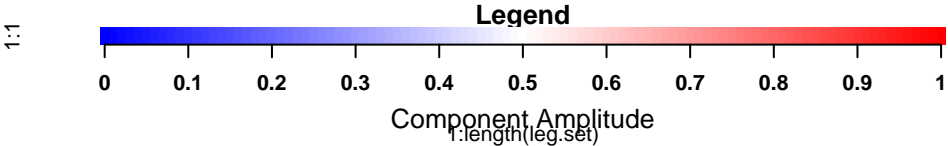
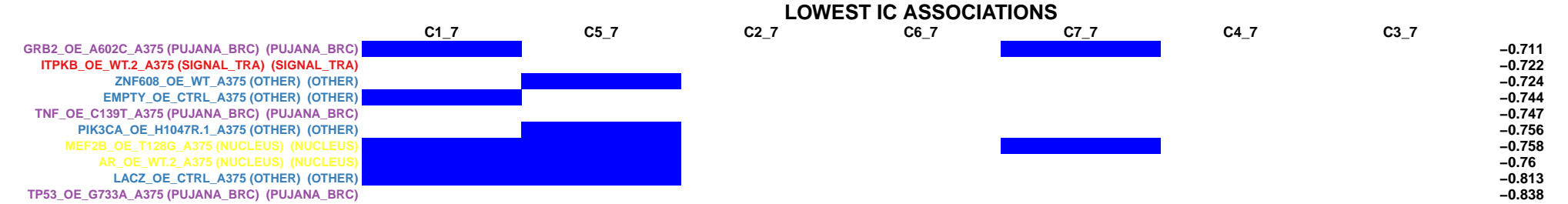
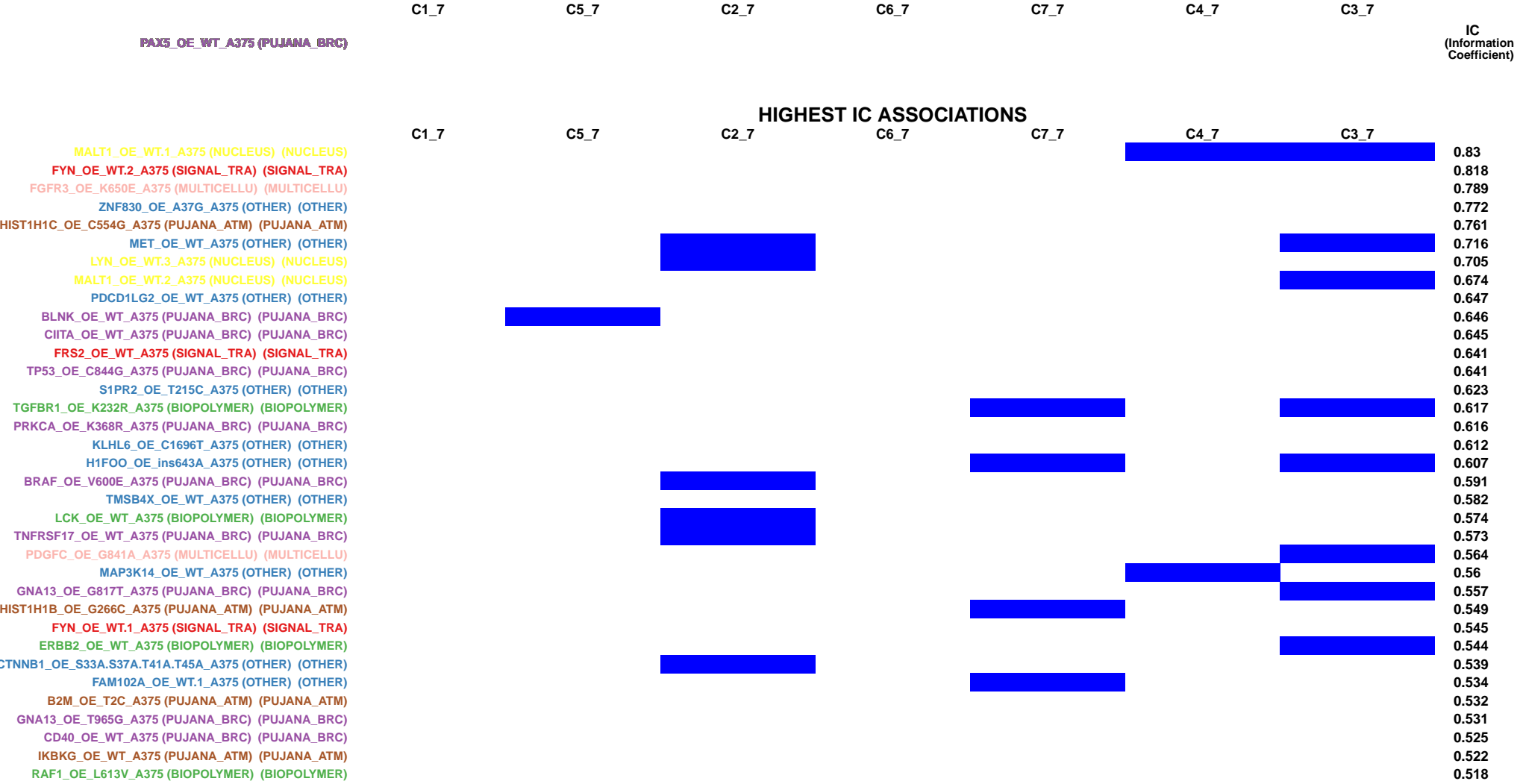
1:length(leg.set)

YAP1\_OE\_Y357F.1\_A375 (OTHER)



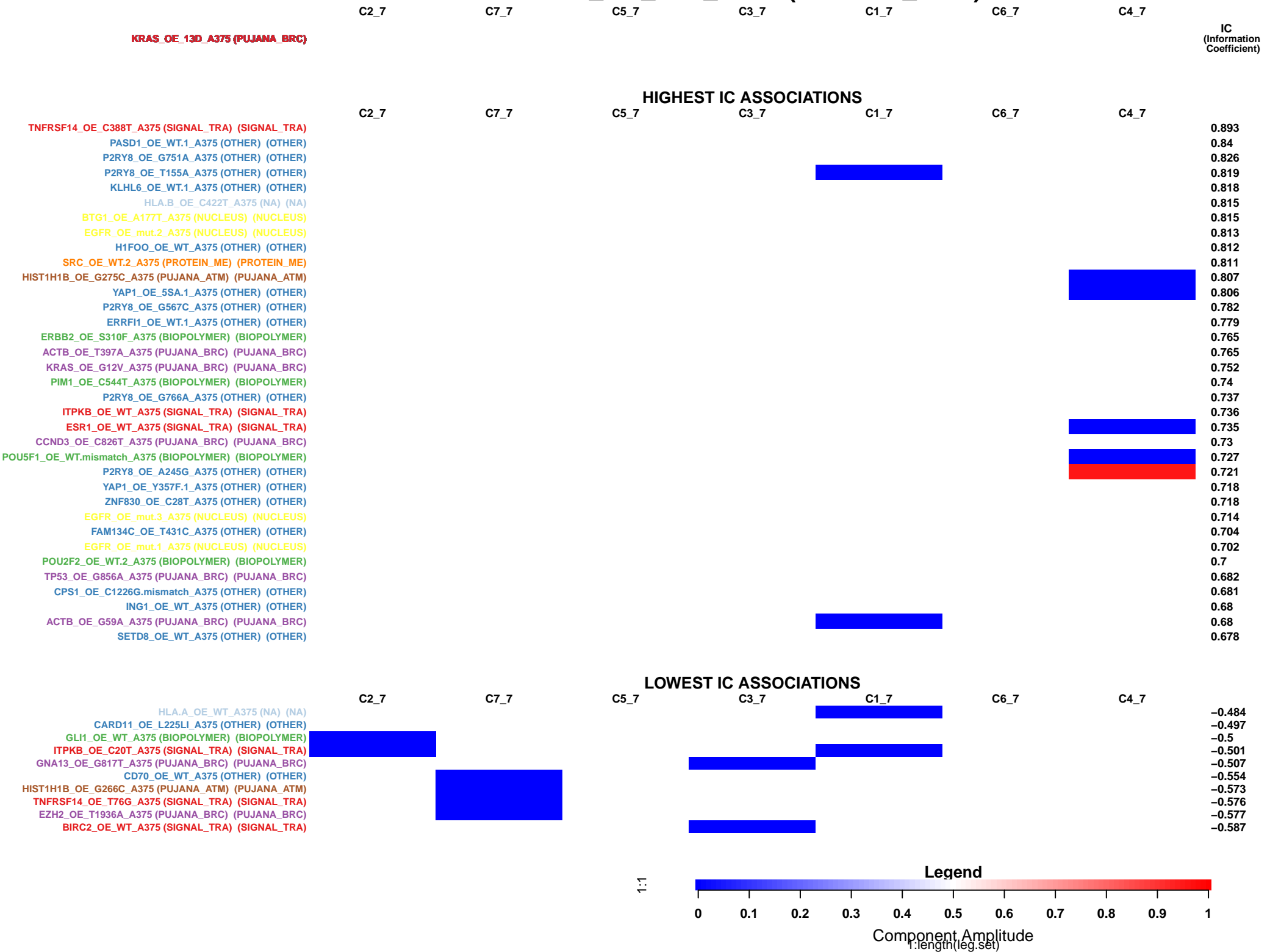
PAX5\_OE\_WT\_A375 (PUJANA\_BRC)

PAX5\_OE\_WT\_A375 (PUJANA\_BRC)

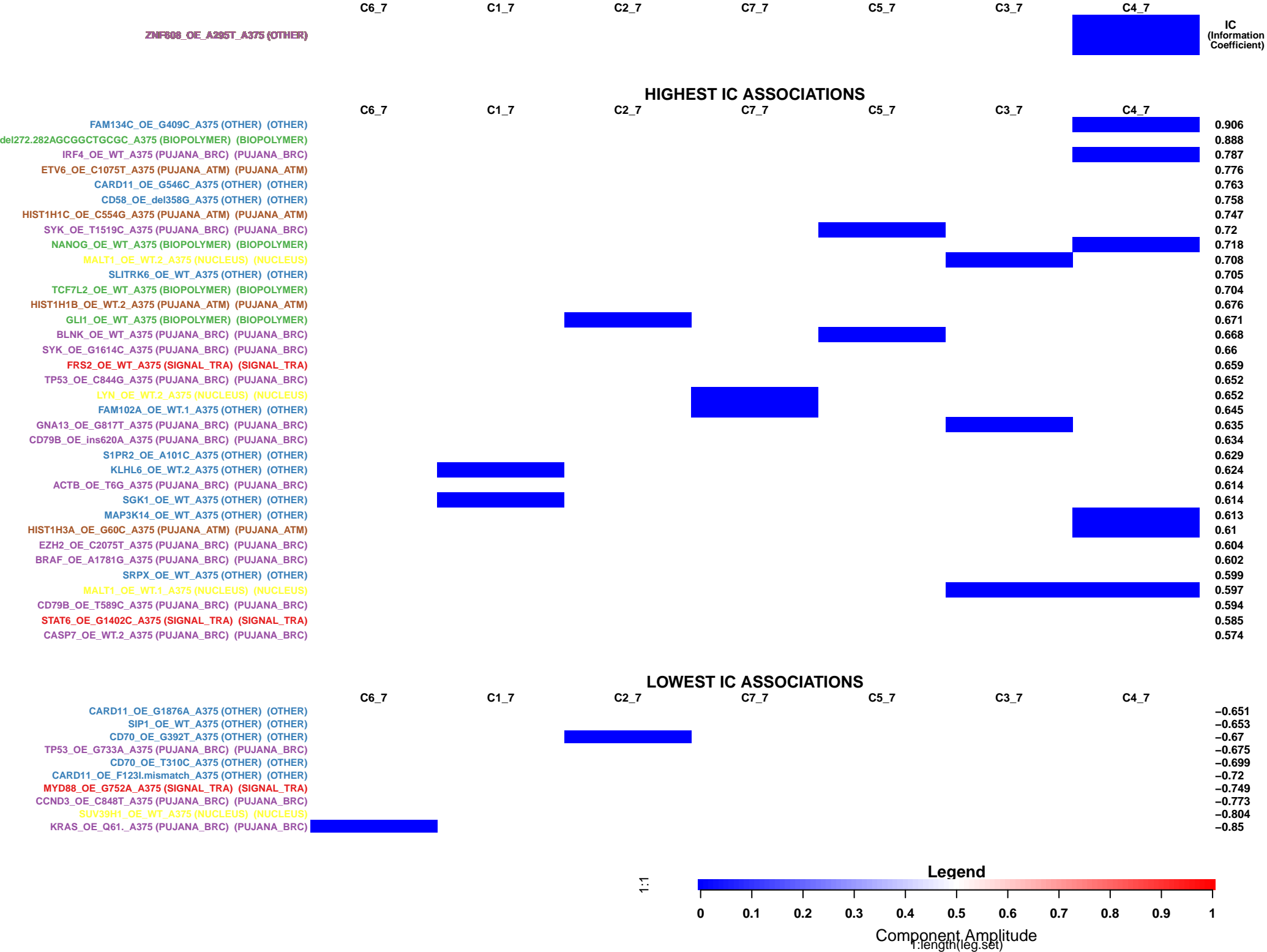




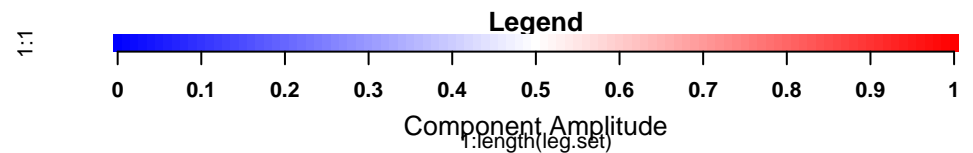
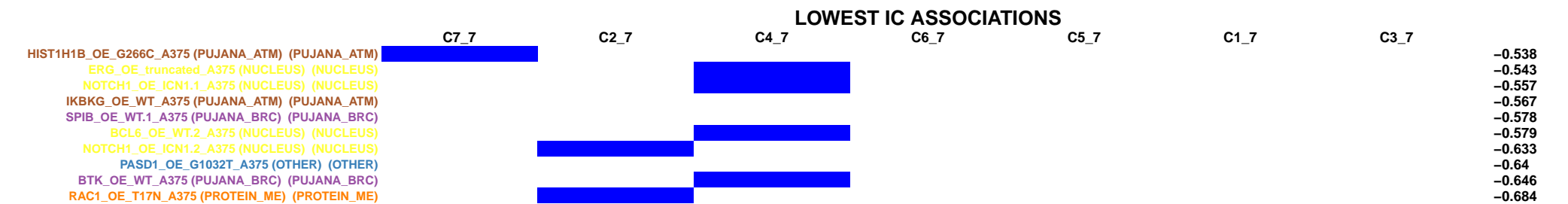
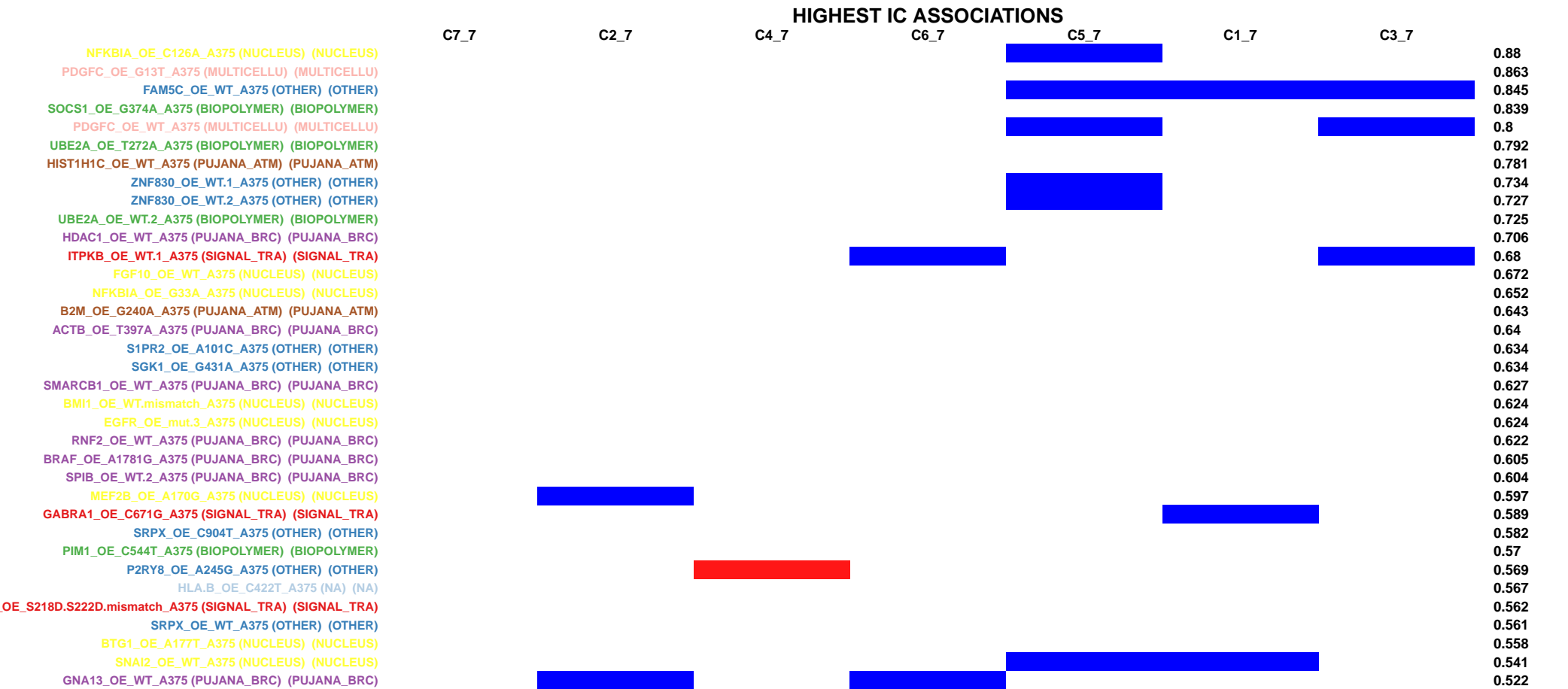
# KRAS\_OE\_13D\_A375 (PUJANA\_BRC)



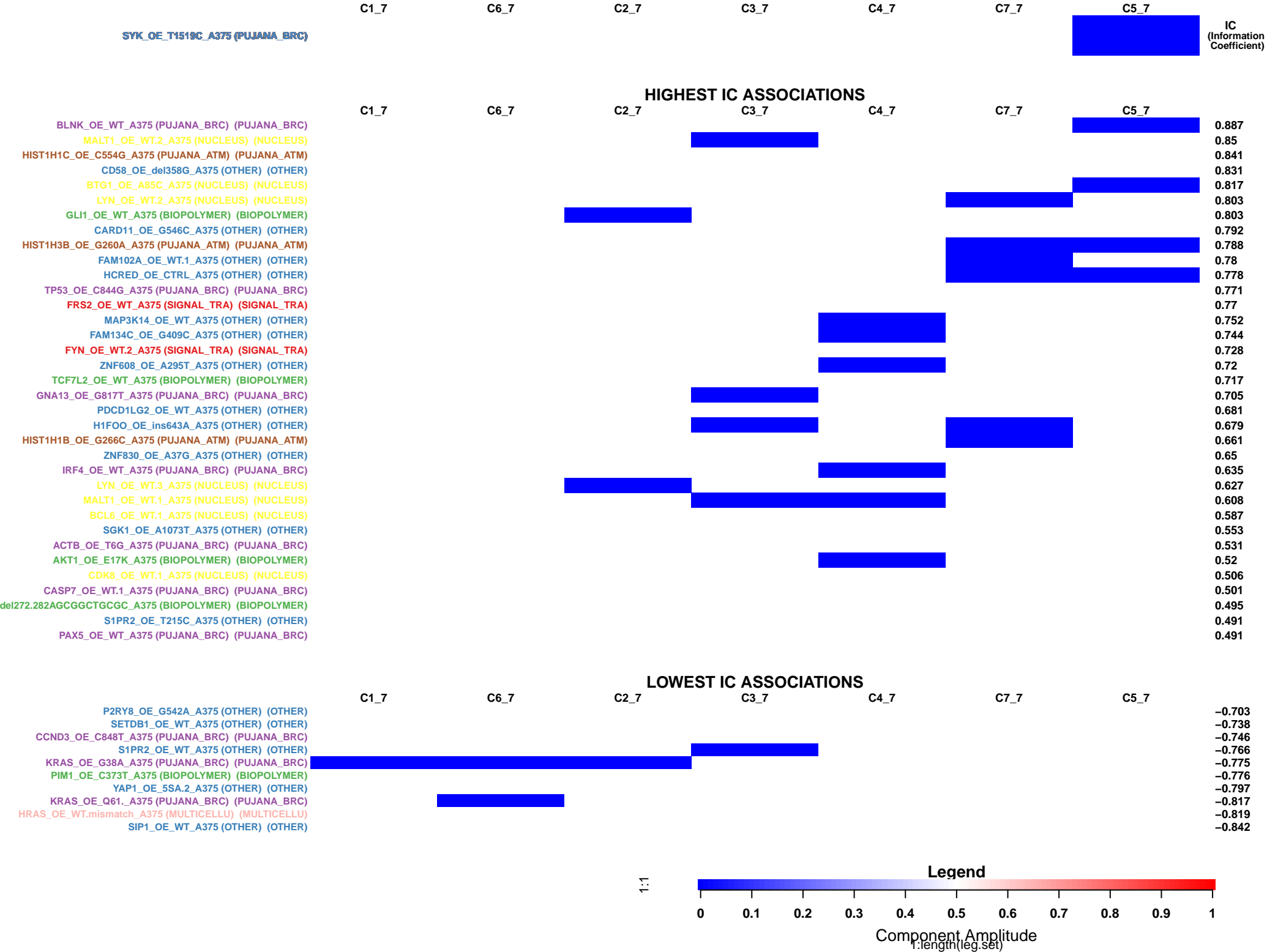
ZNF608\_OE\_A295T\_A375 (OTHER)



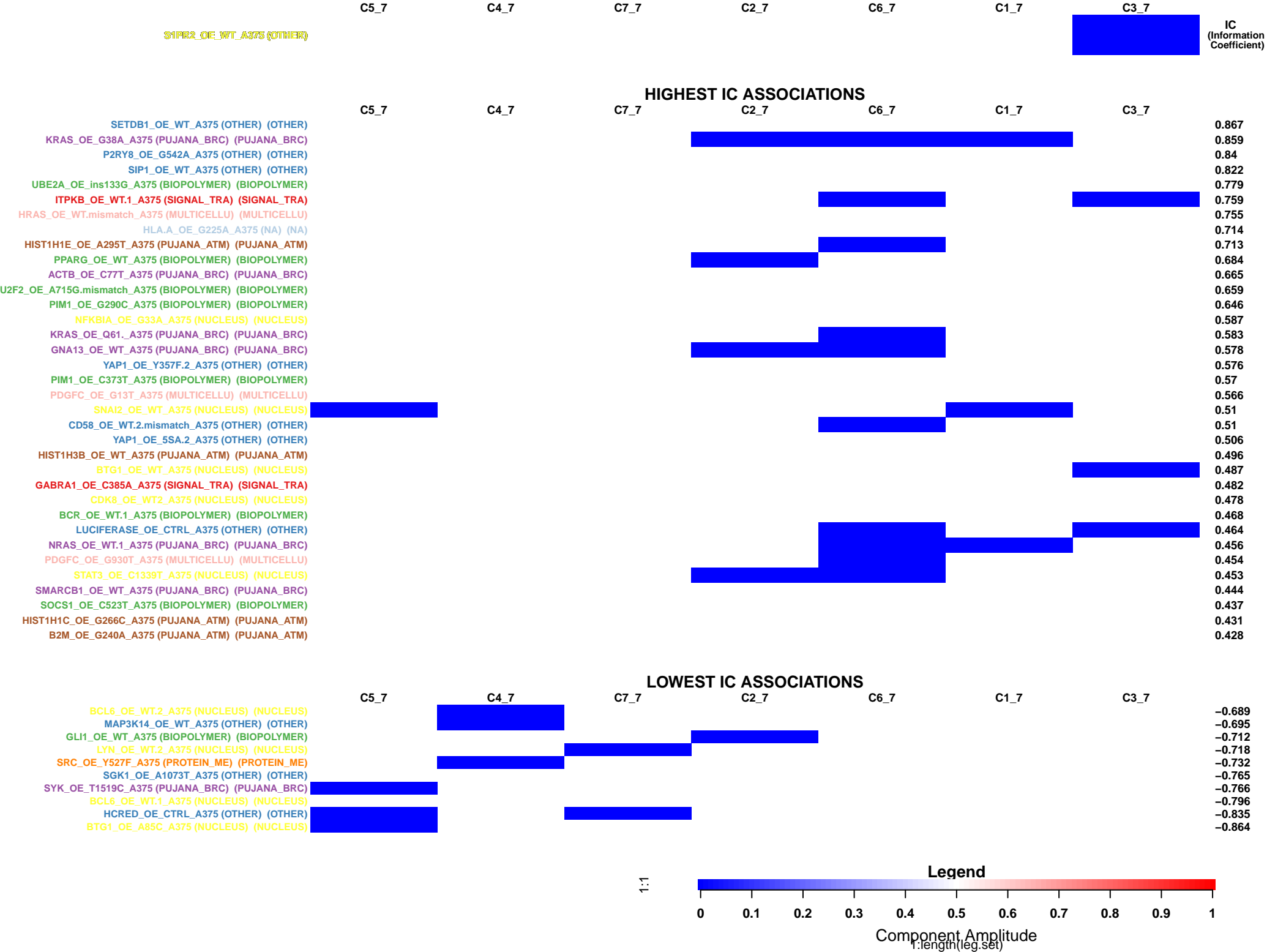
C7_7	C2_7	C4_7	C6_7	C5_7	C1_7	C3_7	IC (Information Coefficient)
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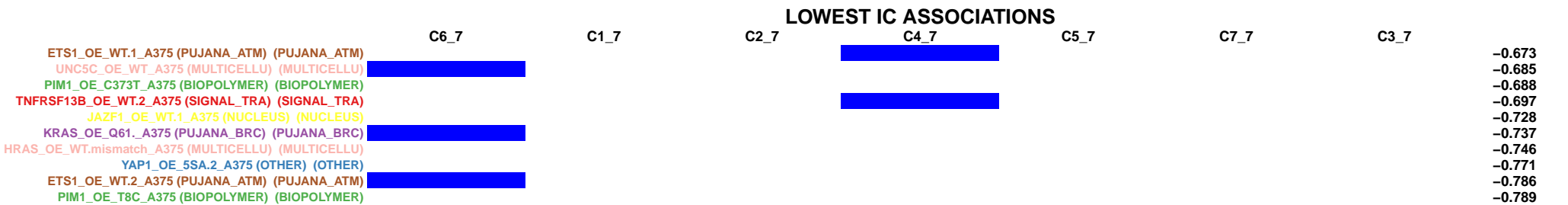
SYK\_OE\_T1519C\_A375 (PUJANA\_BRC)



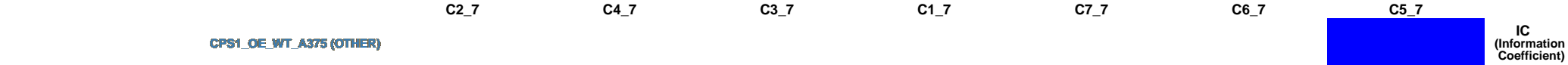
S1PR2\_OE\_WT\_A375 (OTHER)



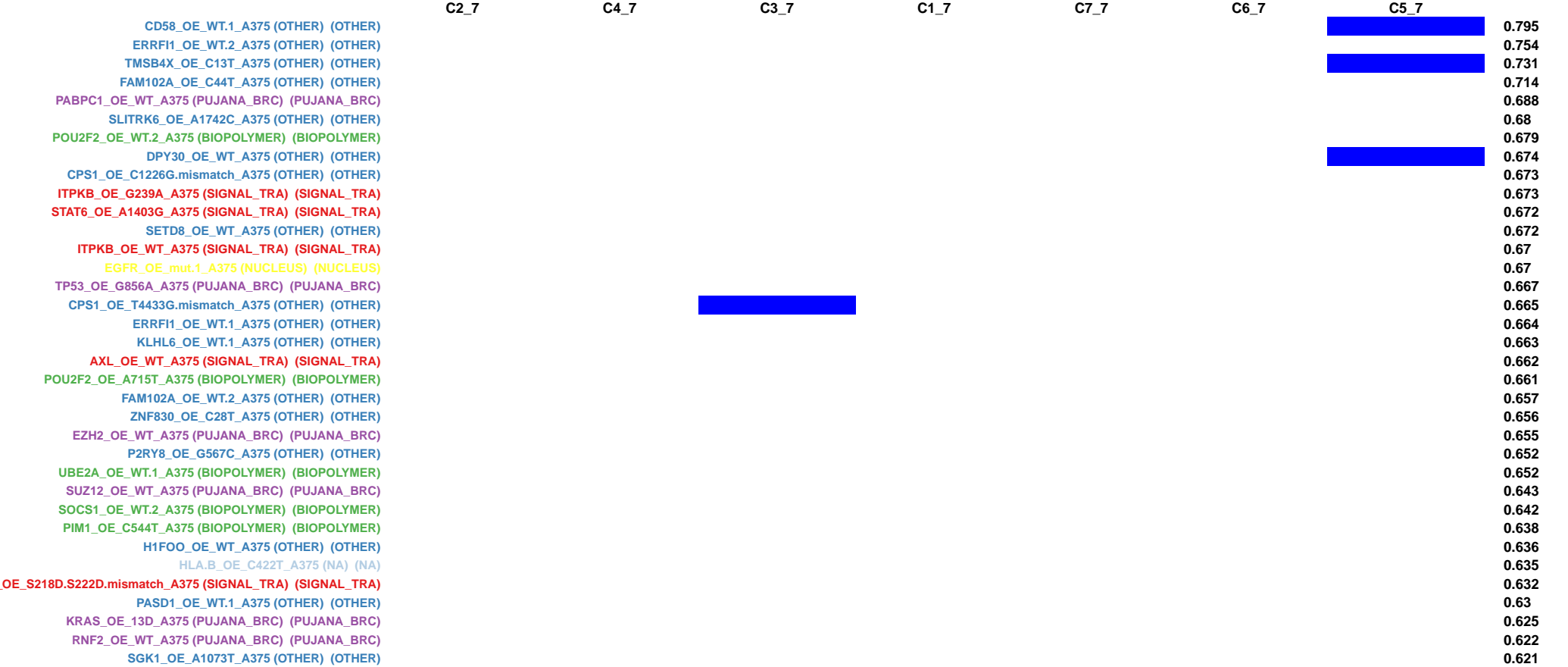
C6_7	C1_7	C2_7	C4_7	C5_7	C7_7	C3_7	IC (Information Coefficient)
------	------	------	------	------	------	------	------------------------------------



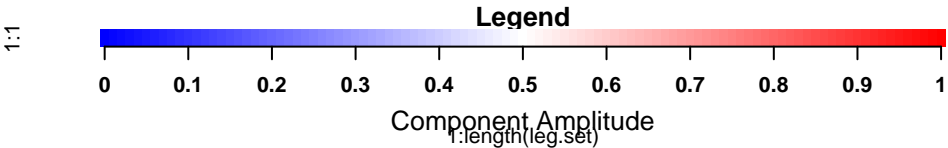
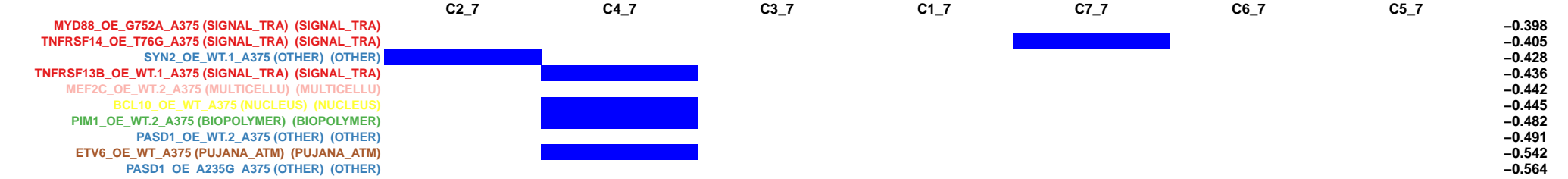
CPS1\_OE\_WT\_A375 (OTHER)



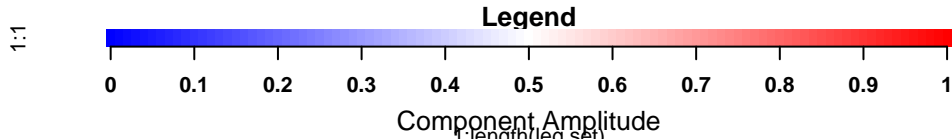
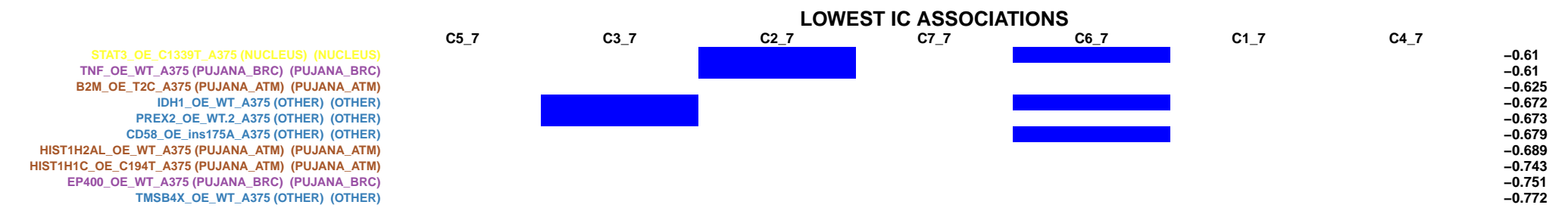
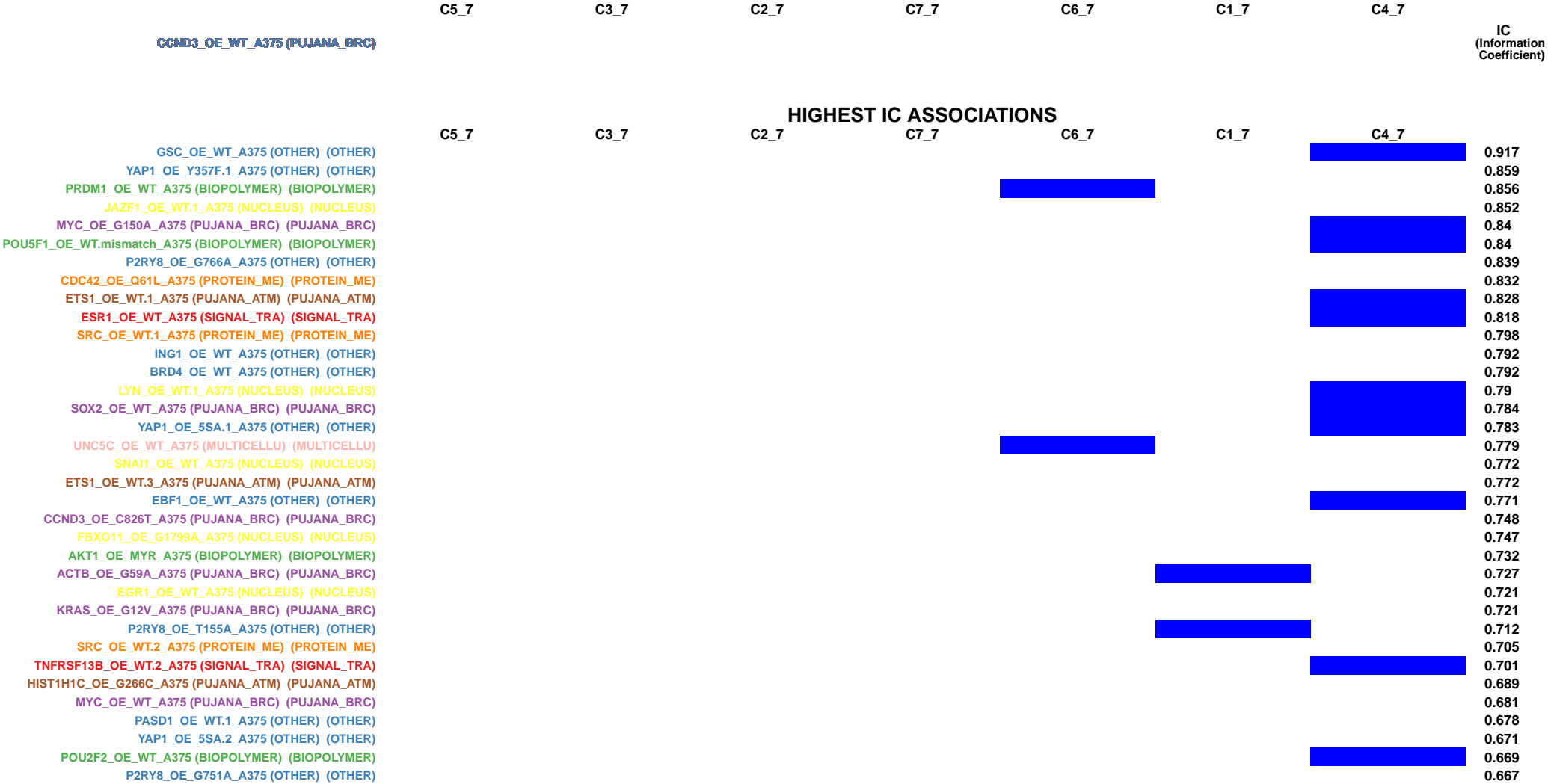
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS

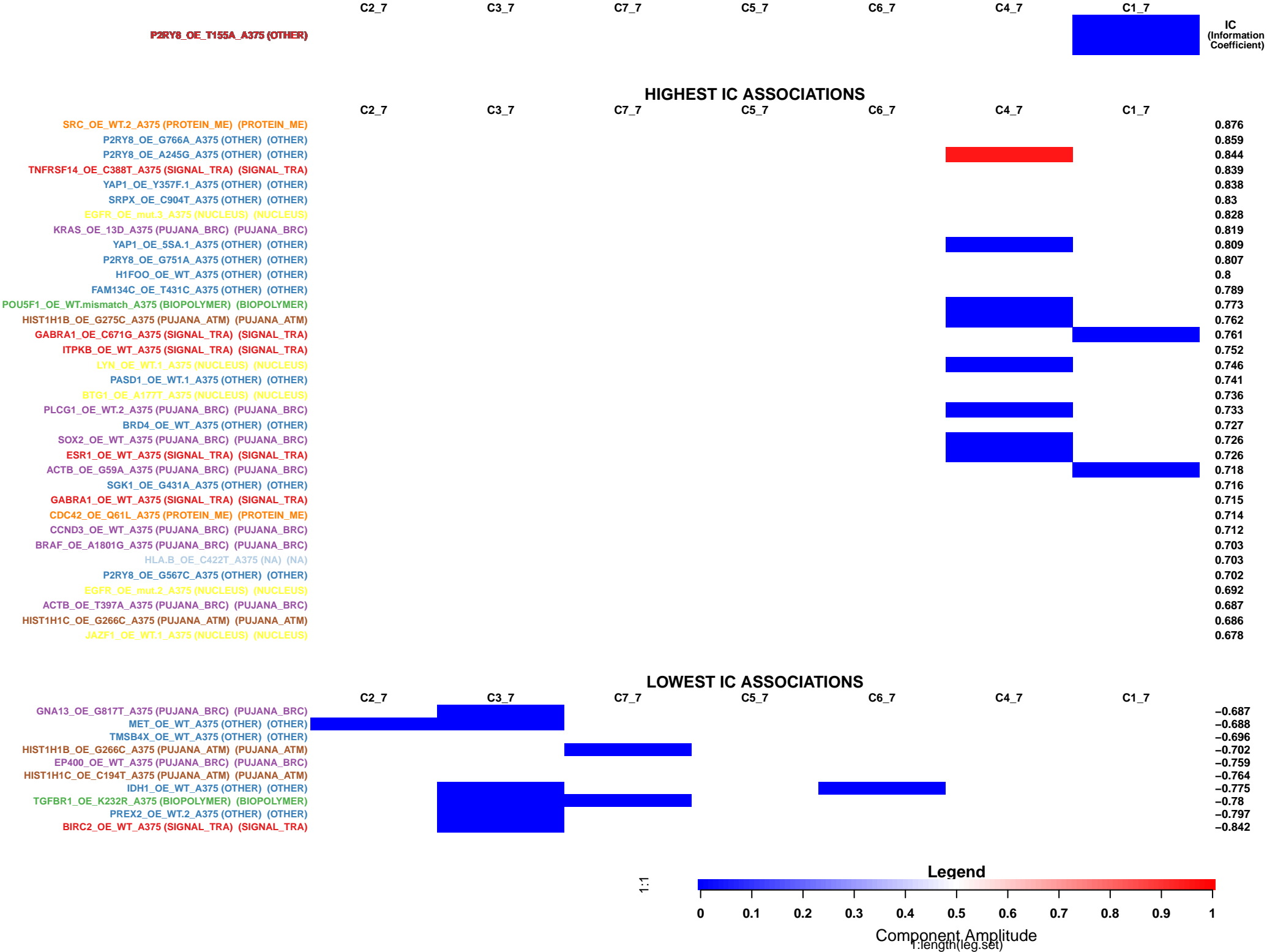


CCND3\_OE\_WT\_A375 (PUJANA\_BRC)

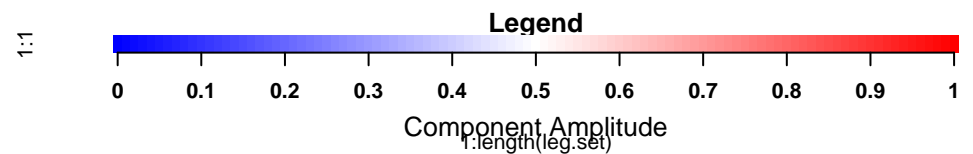
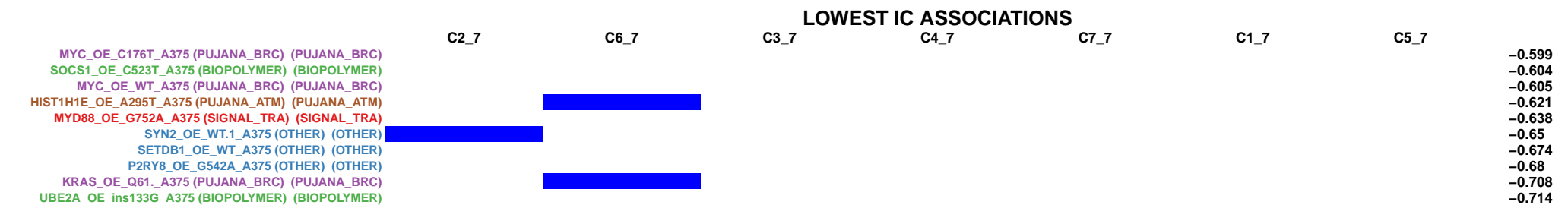
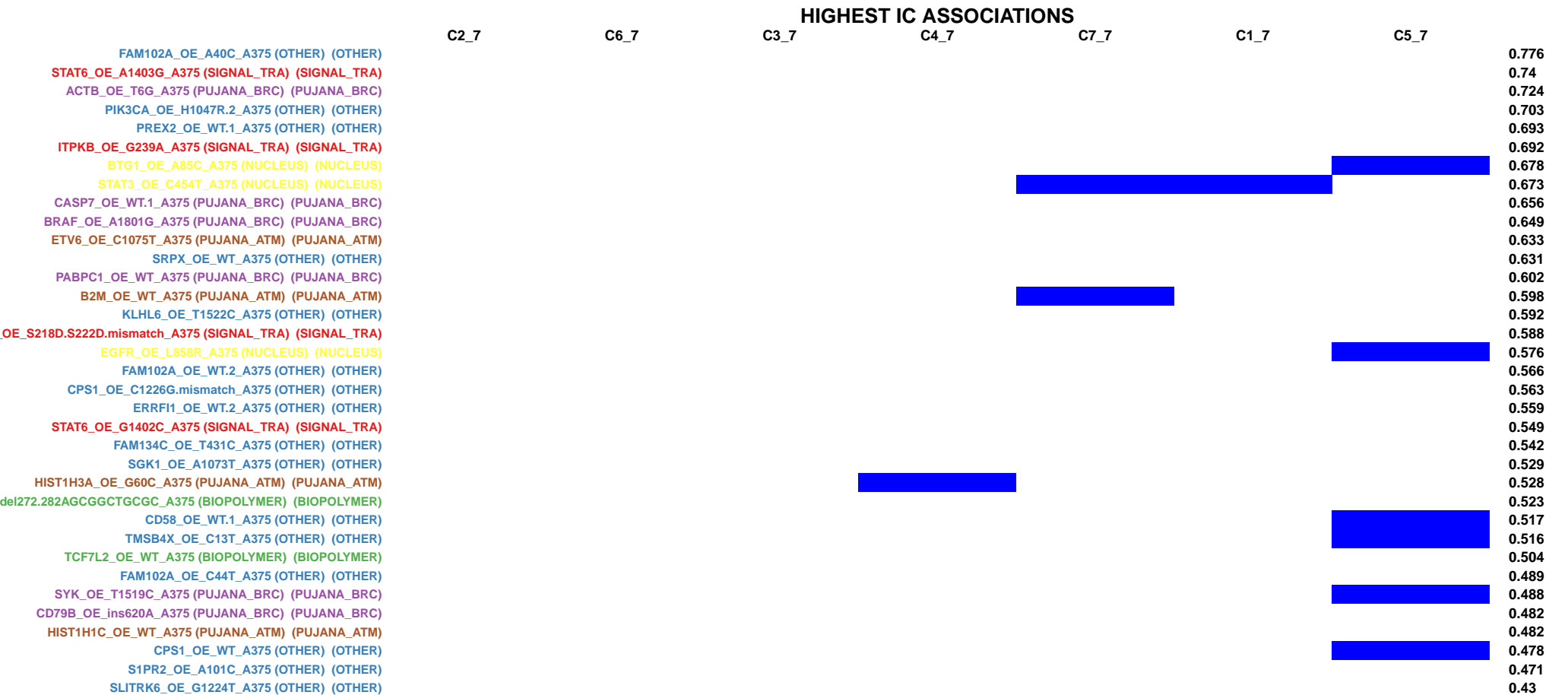




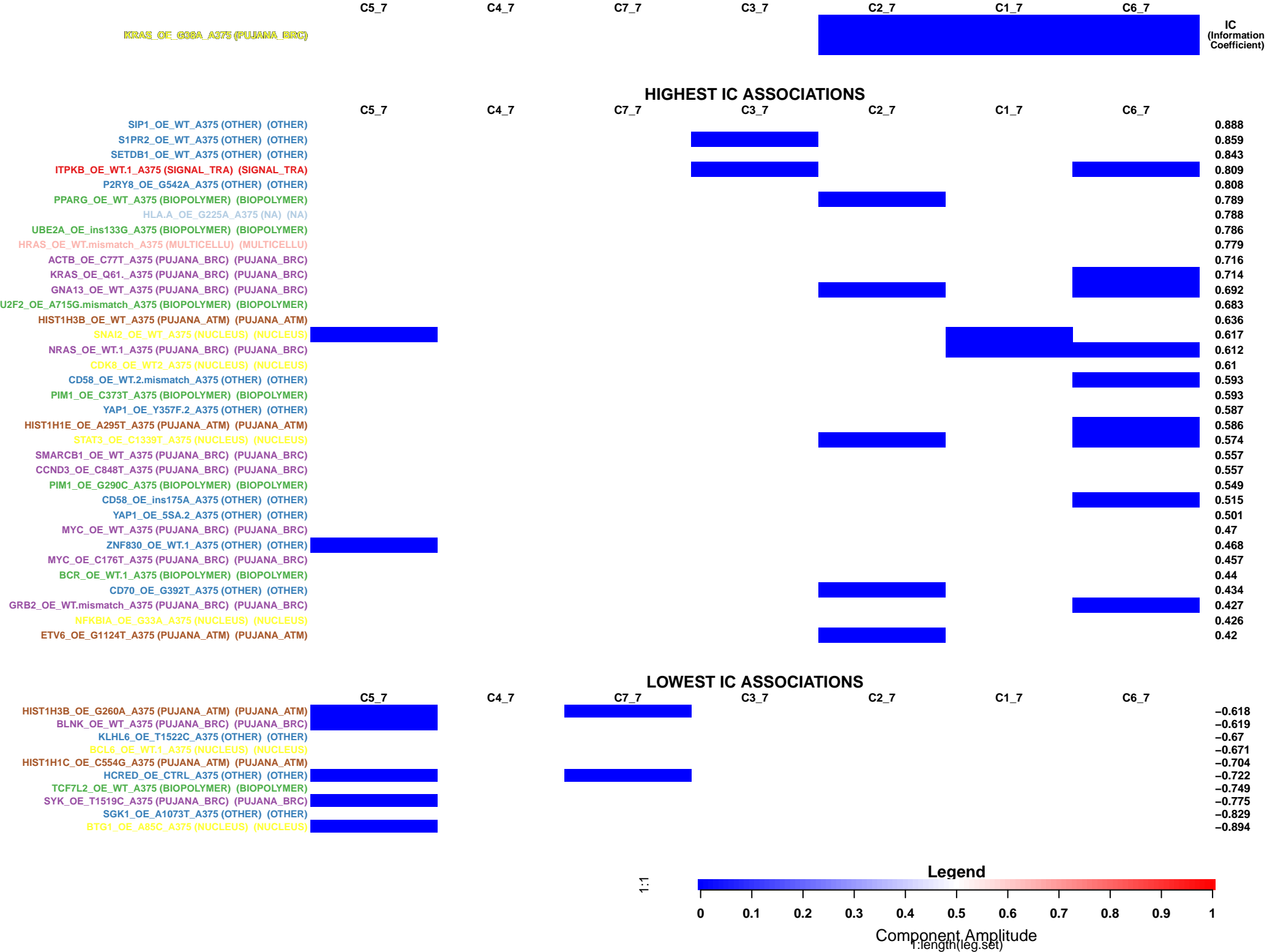
P2RY8\_OE\_T155A\_A375 (OTHER)



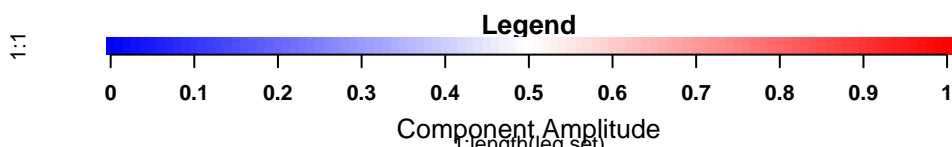
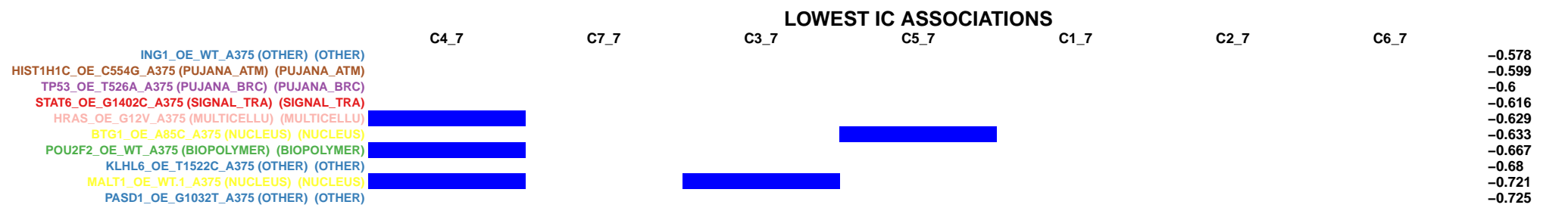
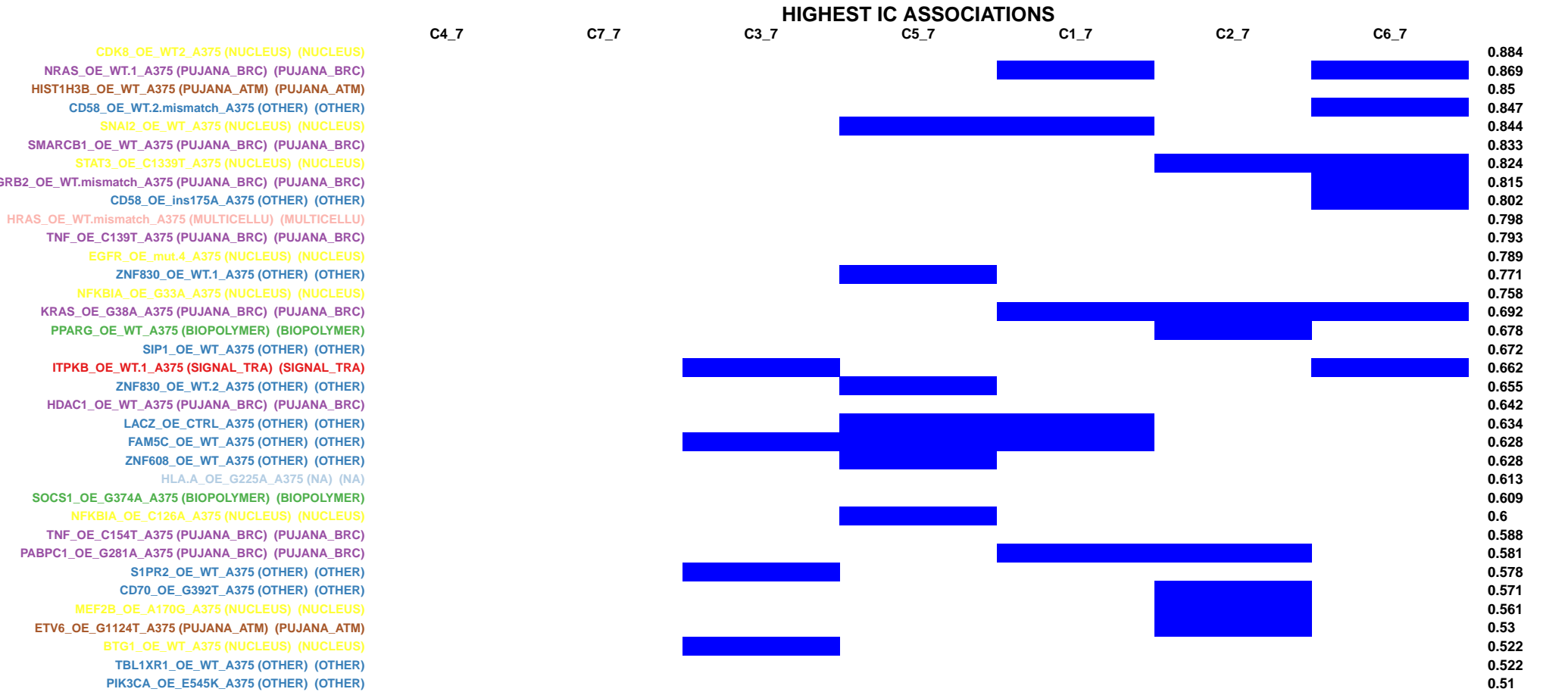
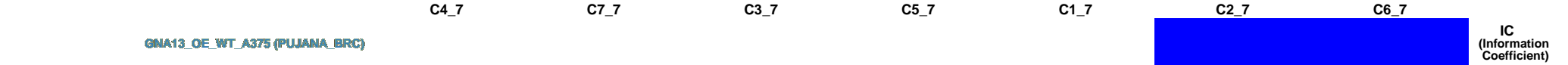
C2_7	C6_7	C3_7	C4_7	C7_7	C1_7	C5_7	IC (Information Coefficient)



KRAS\_OE\_G38A\_A375 (PUJANA\_BRC)



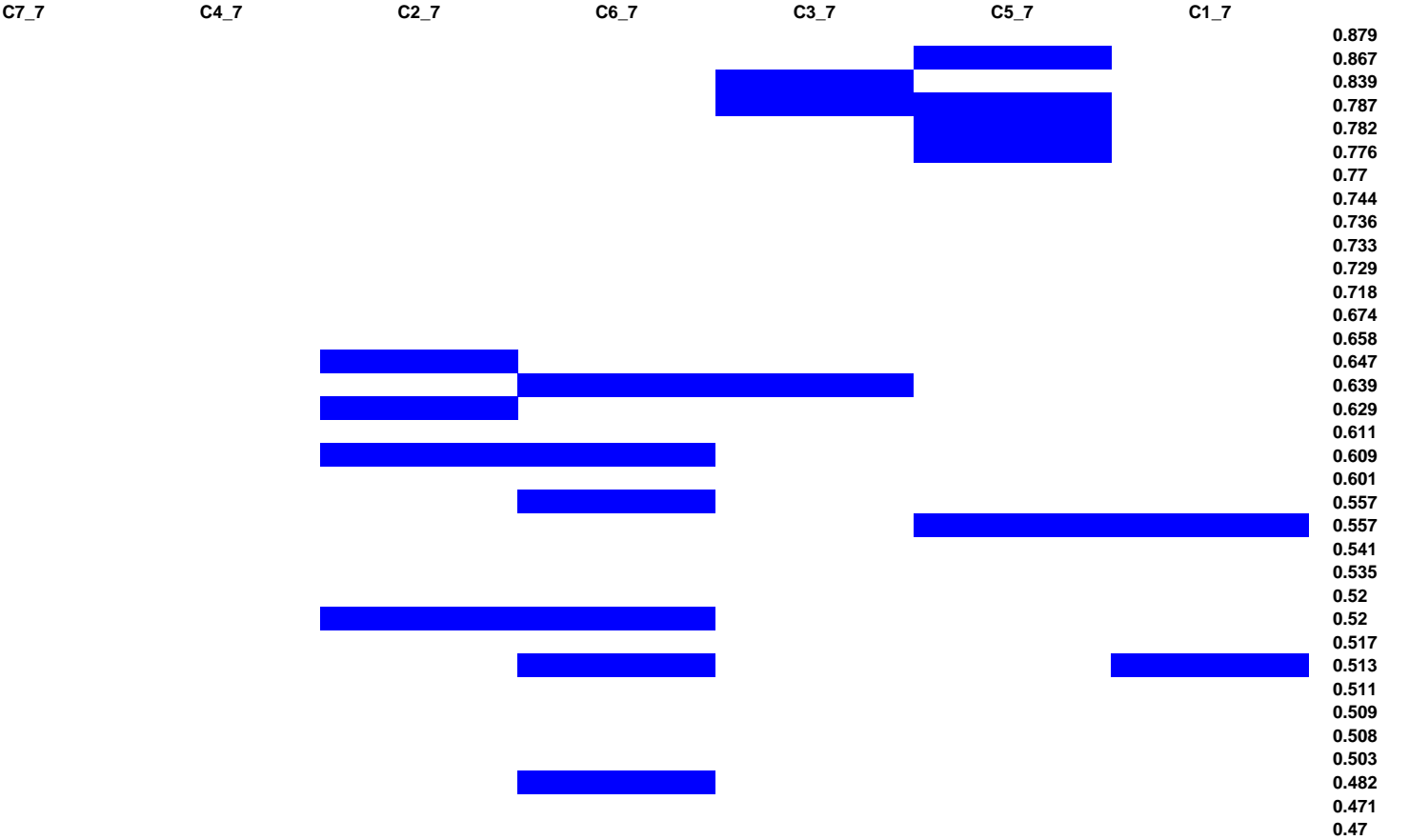
GNA13\_OE\_WT\_A375 (PUJANA\_BRC)



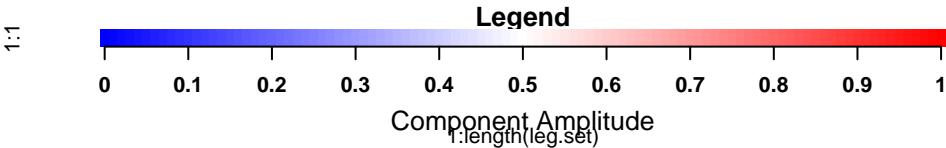
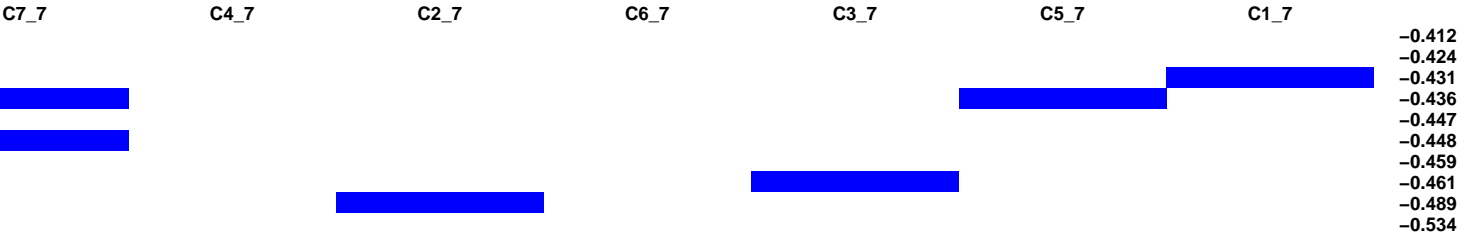
SOCS1\_OE\_G374A\_A375 (BIOPOLYMER)



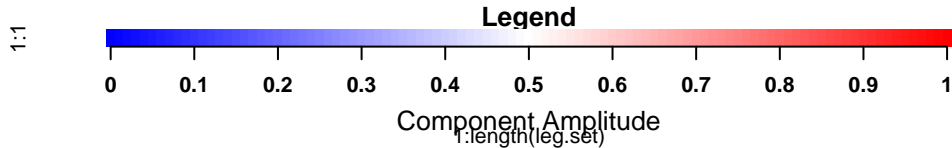
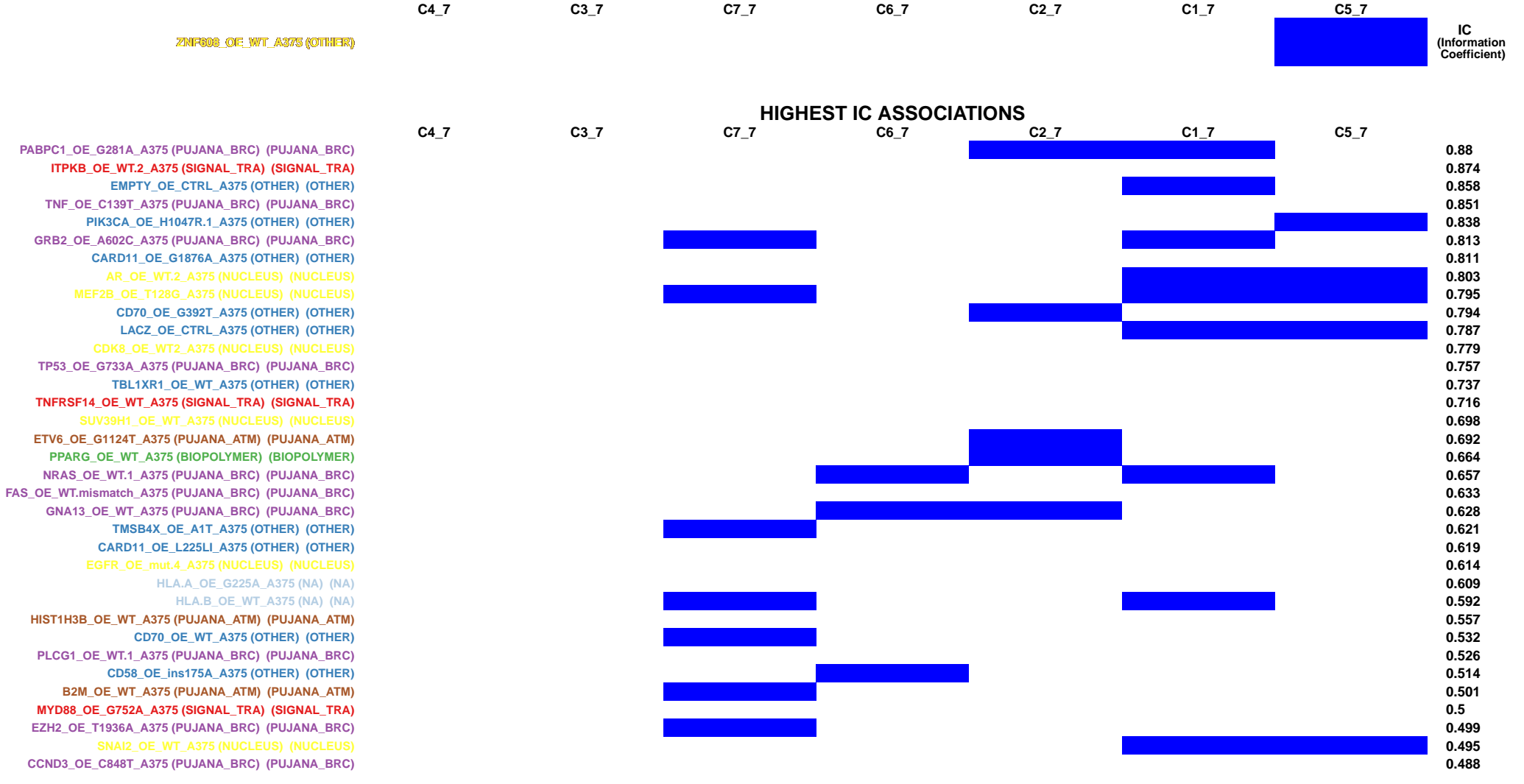
HIGHEST IC ASSOCIATIONS



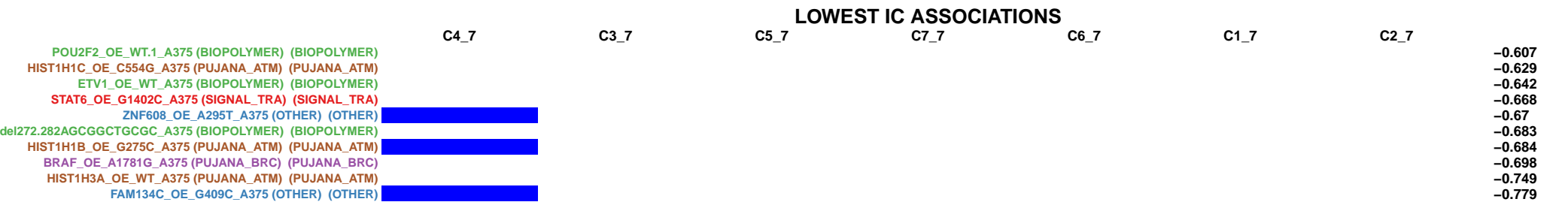
LOWEST IC ASSOCIATIONS



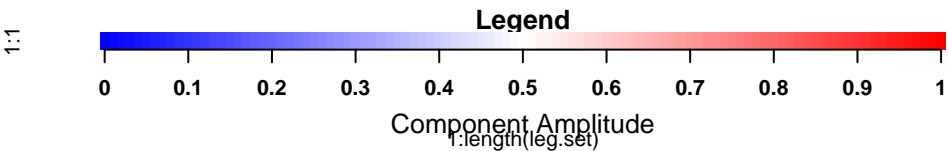
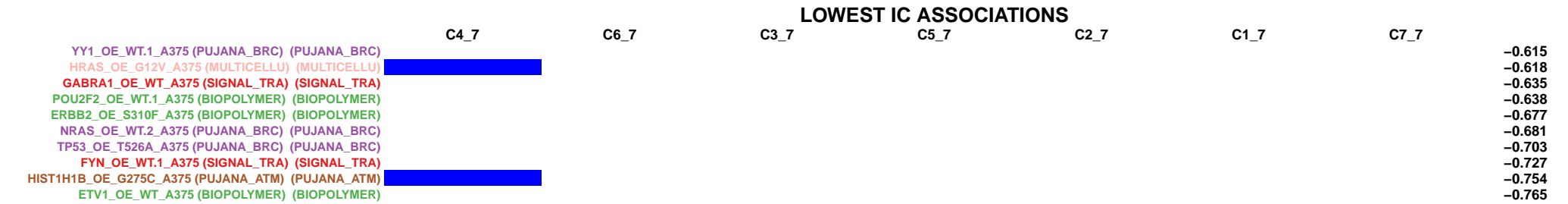
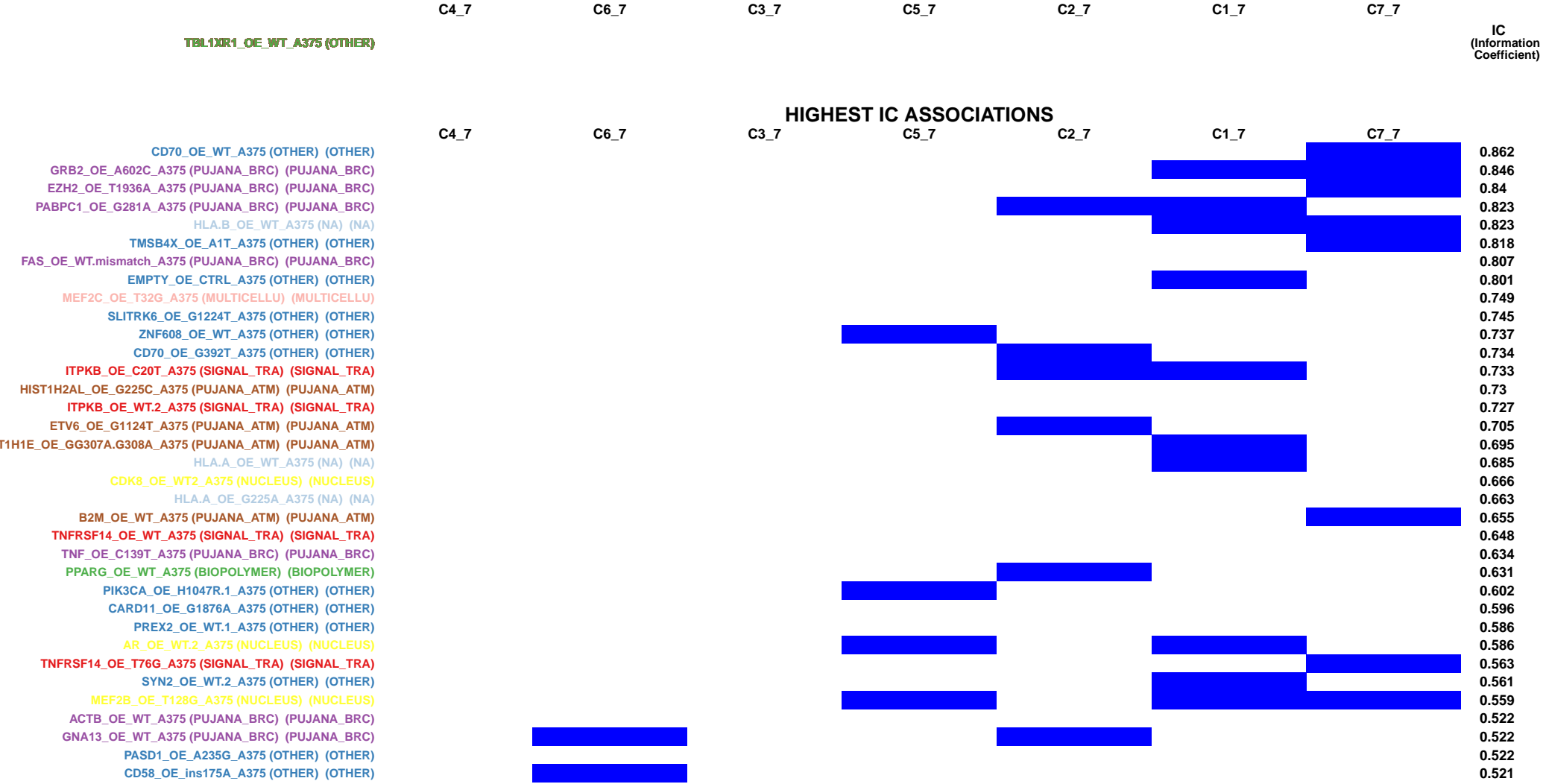
ZNF608\_OE\_WT\_A375 (OTHER)



C4_7	C3_7	C5_7	C7_7	C6_7	C1_7	C2_7	IC (Information Coefficient)
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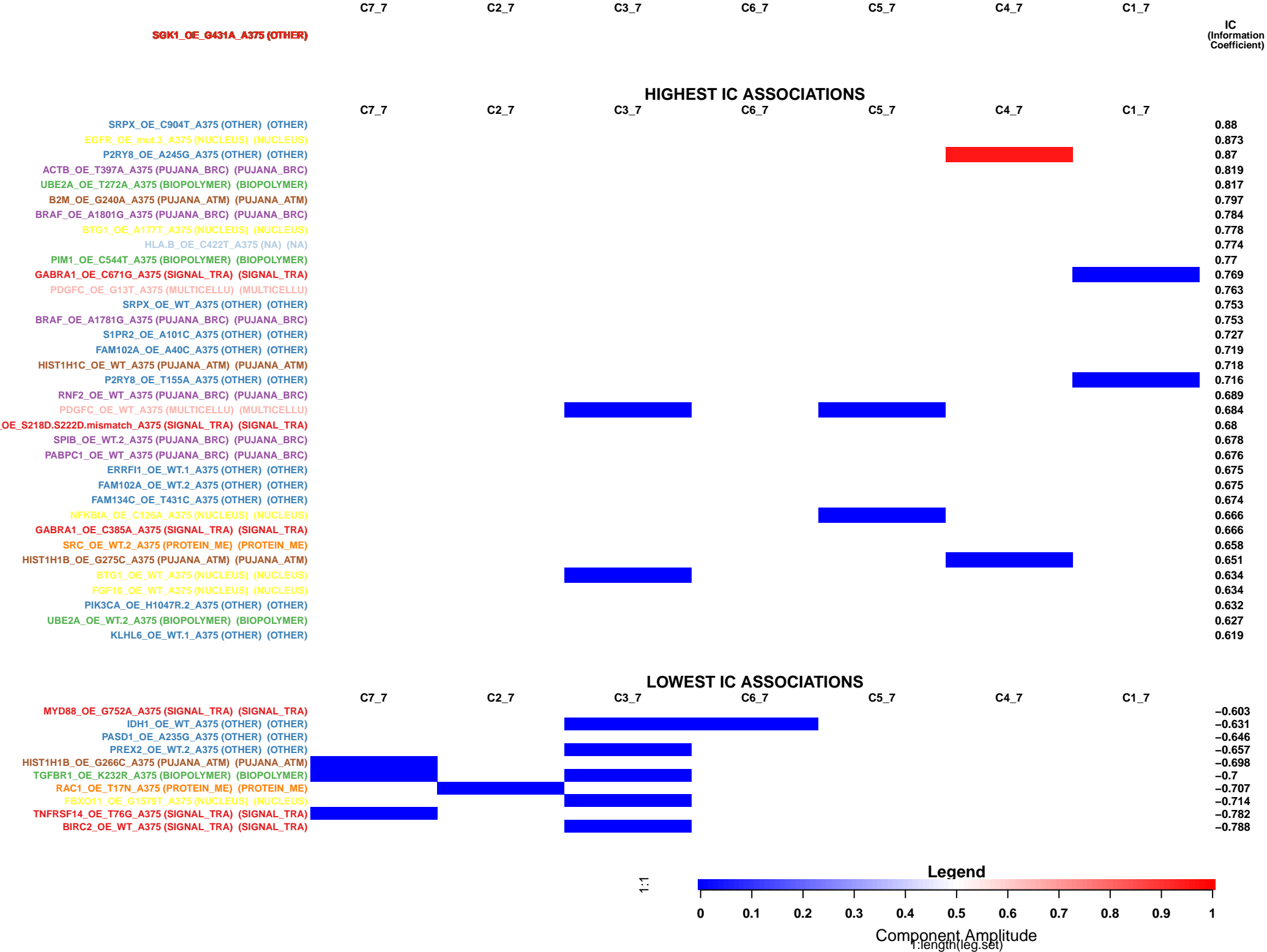


TBL1XR1\_OE\_WT\_A375 (OTHER)

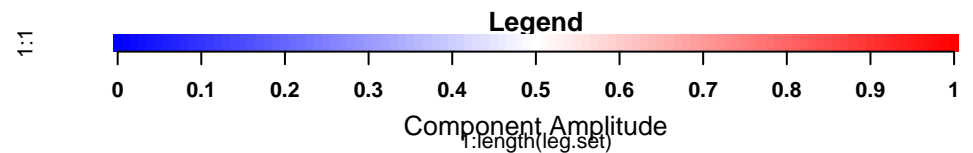
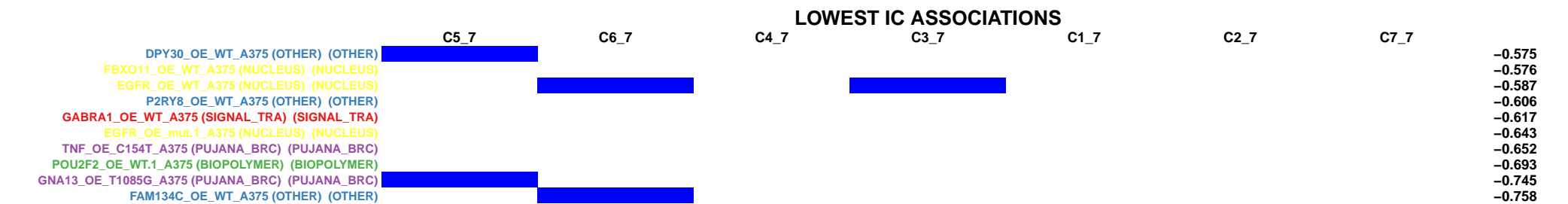
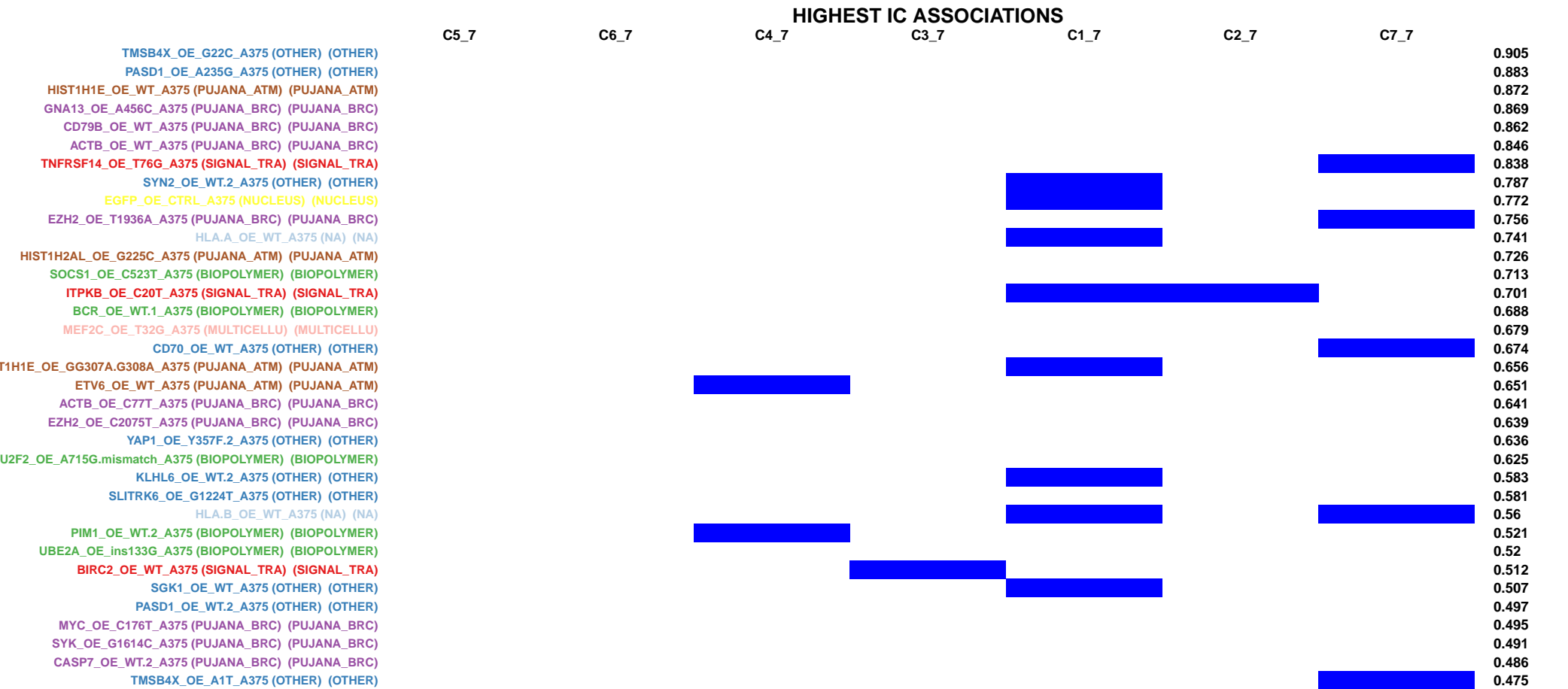




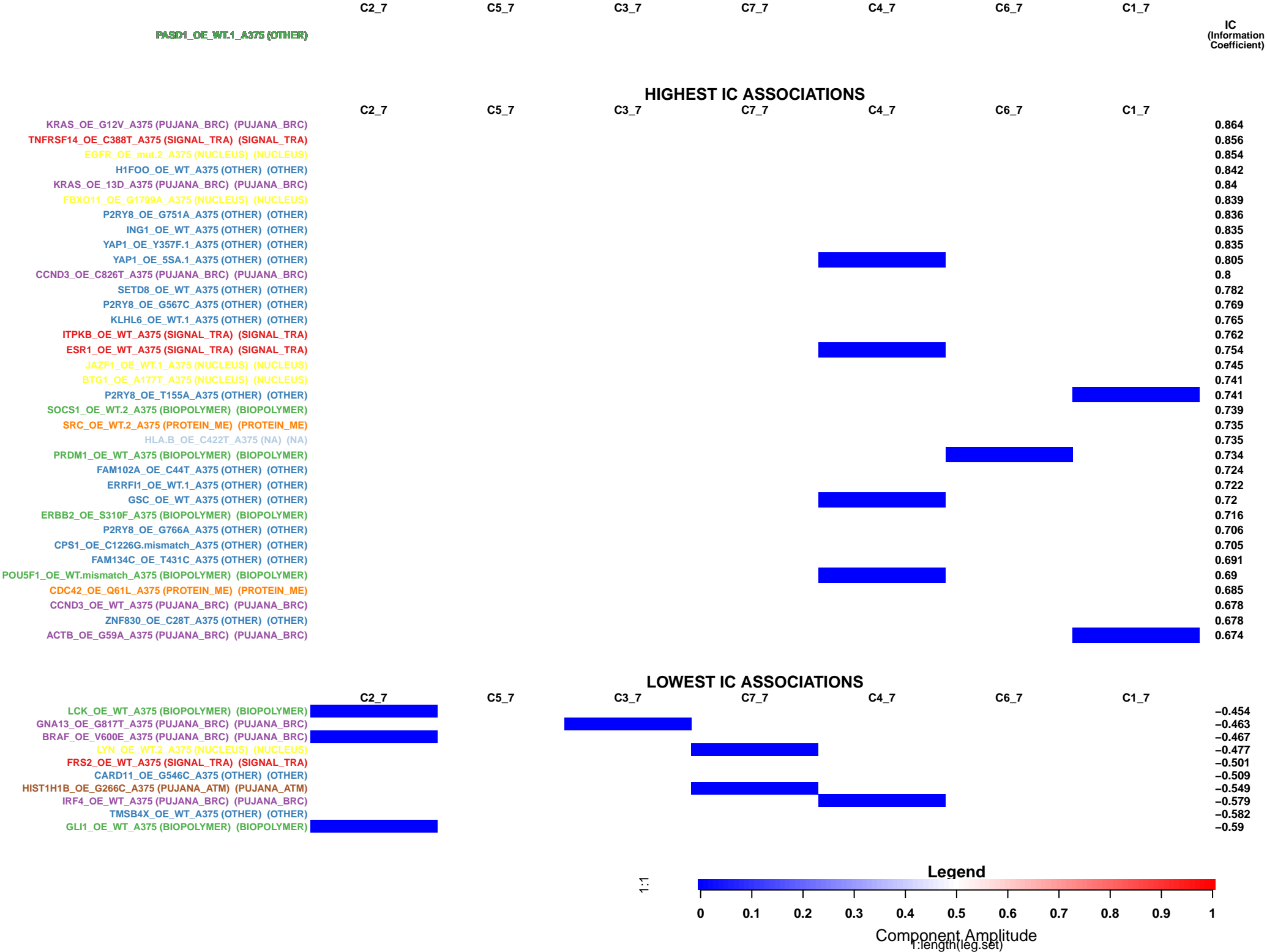
SGK1\_OE\_G431A\_A375 (OTHER)



C5_7	C6_7	C4_7	C3_7	C1_7	C2_7	C7_7	IC (Information Coefficient)
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PASD1\_OE\_WT.1\_A375 (OTHER)



C2\_7                      C6\_7                      C1\_7                      C5\_7                      C7\_7                      C4\_7                      C3\_7

IC  
Information  
Coefficient)

C2\_7                      C6\_7                      C1\_7                      C5\_7                      C7\_7                      C4\_7                      C3\_7



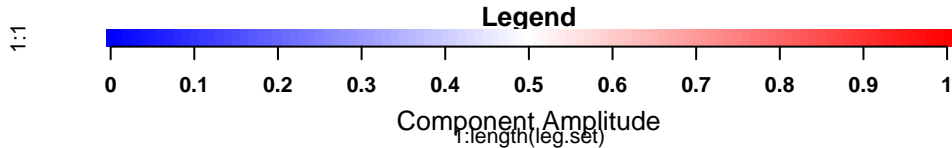
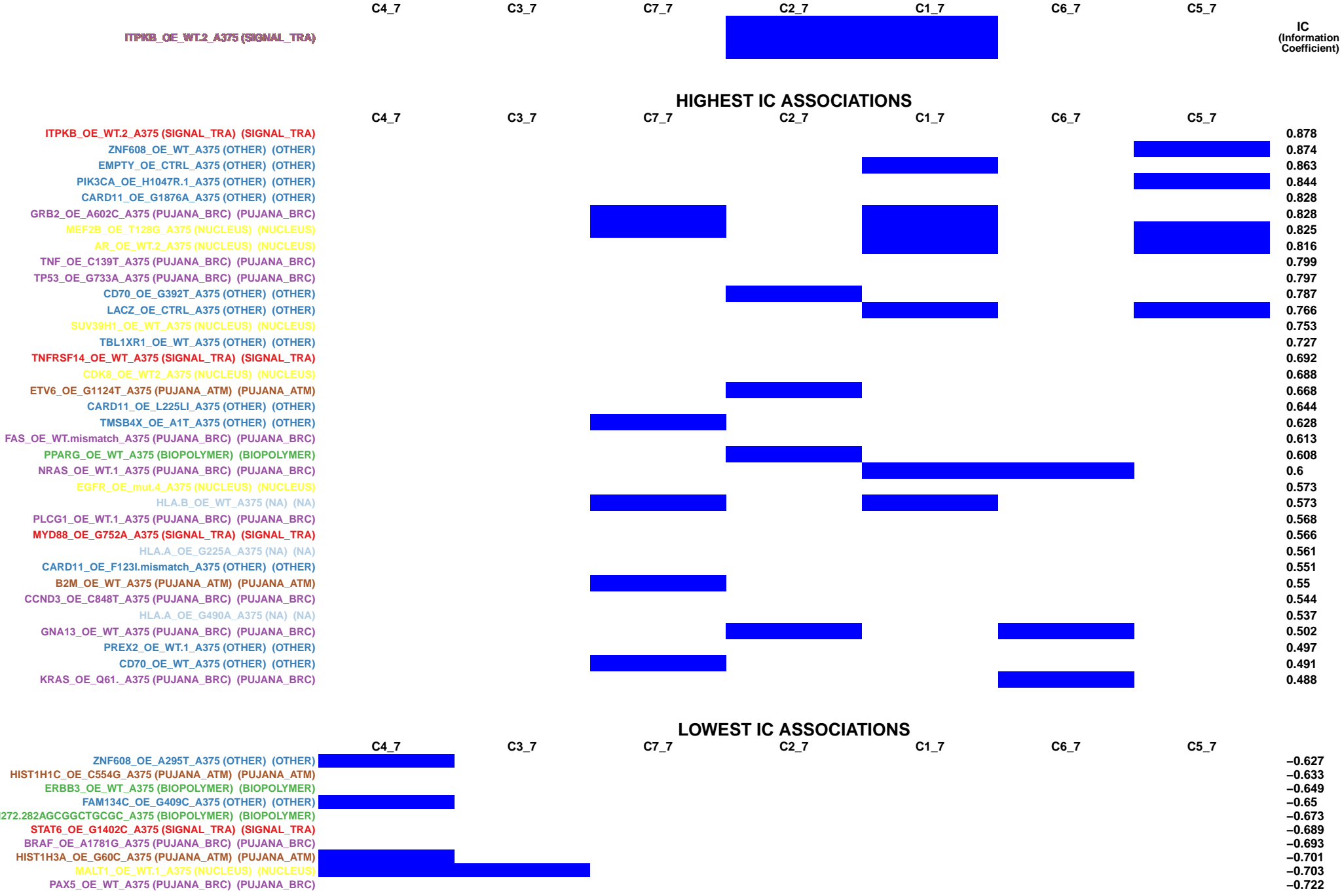
**C2\_7                      C6\_7                      C1\_7                      C5\_7                      C7\_7                      C4\_7                      C3\_7**



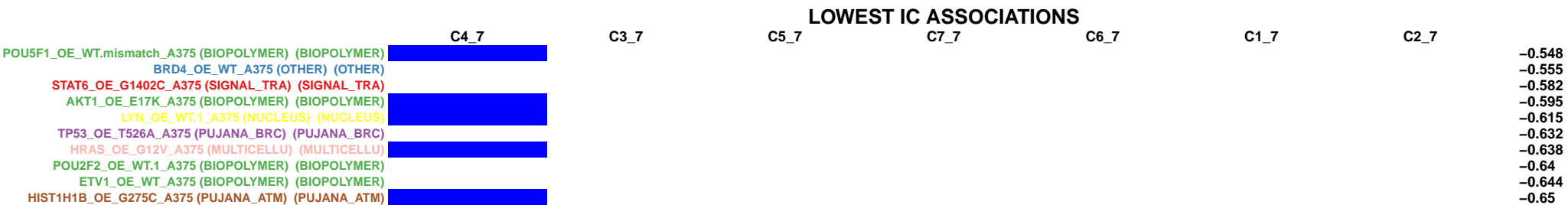
0      0.1      0.2      0.3      0.4      0.5      0.6      0.7      0.8      0.9      1

```
1:length(lég.set)
```

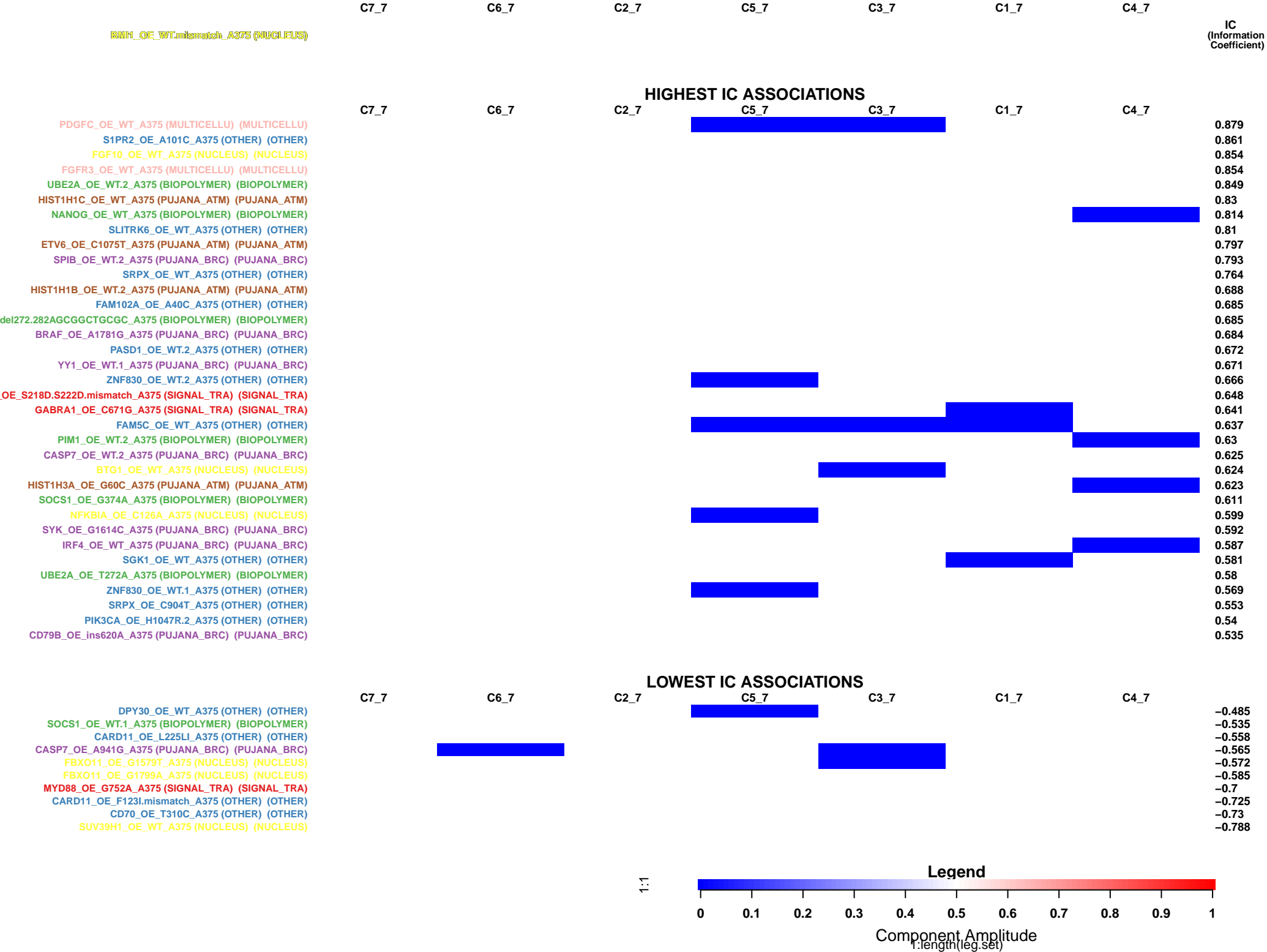
ITPKB\_OE\_WT.2\_A375 (SIGNAL\_TRA)



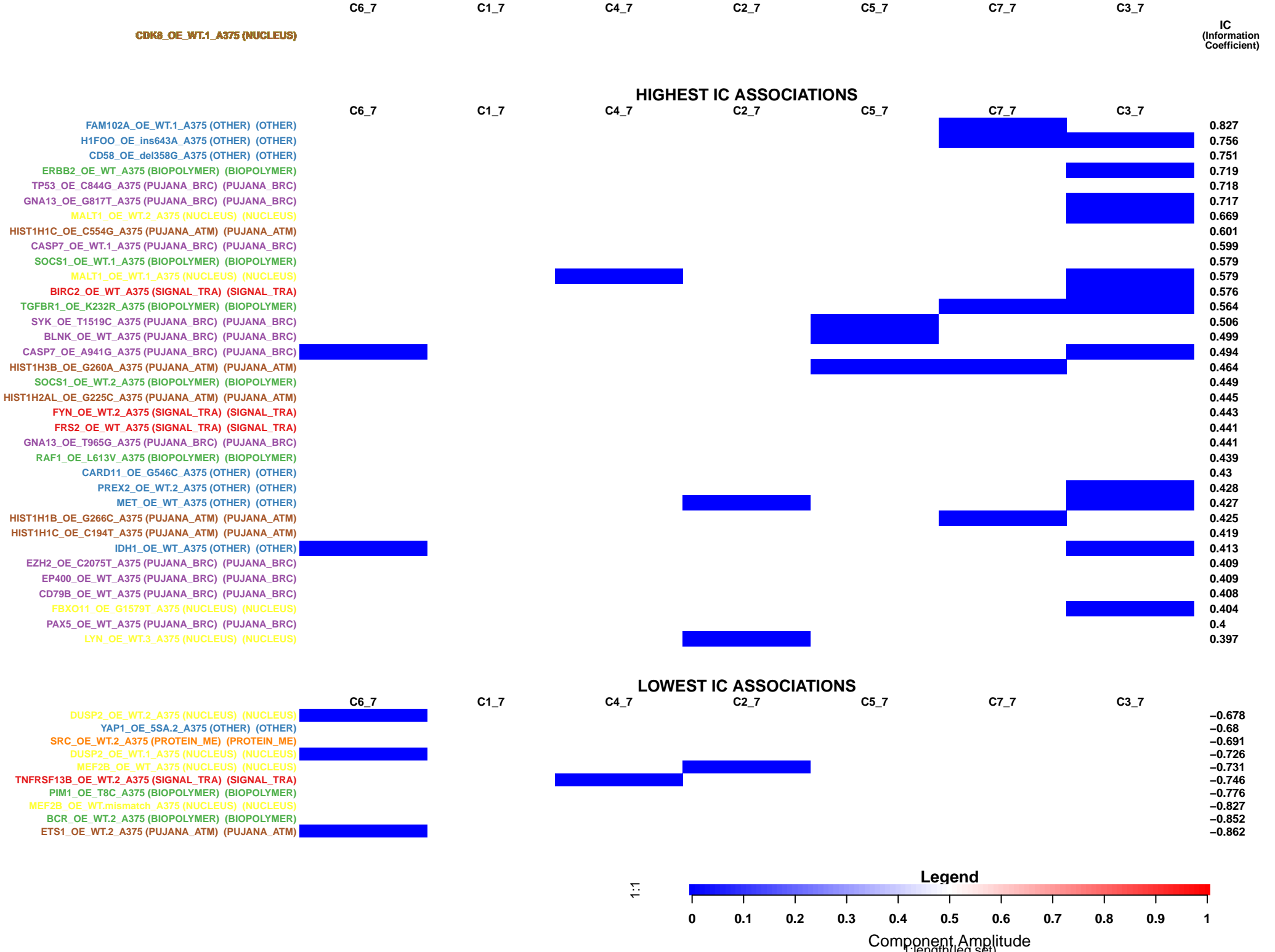
C4_7	C3_7	C5_7	C7_7	C6_7	C1_7	C2_7	IC (Information Coefficient)



BMI1\_OE\_WT.mismatch\_A375 (NUCLEUS)

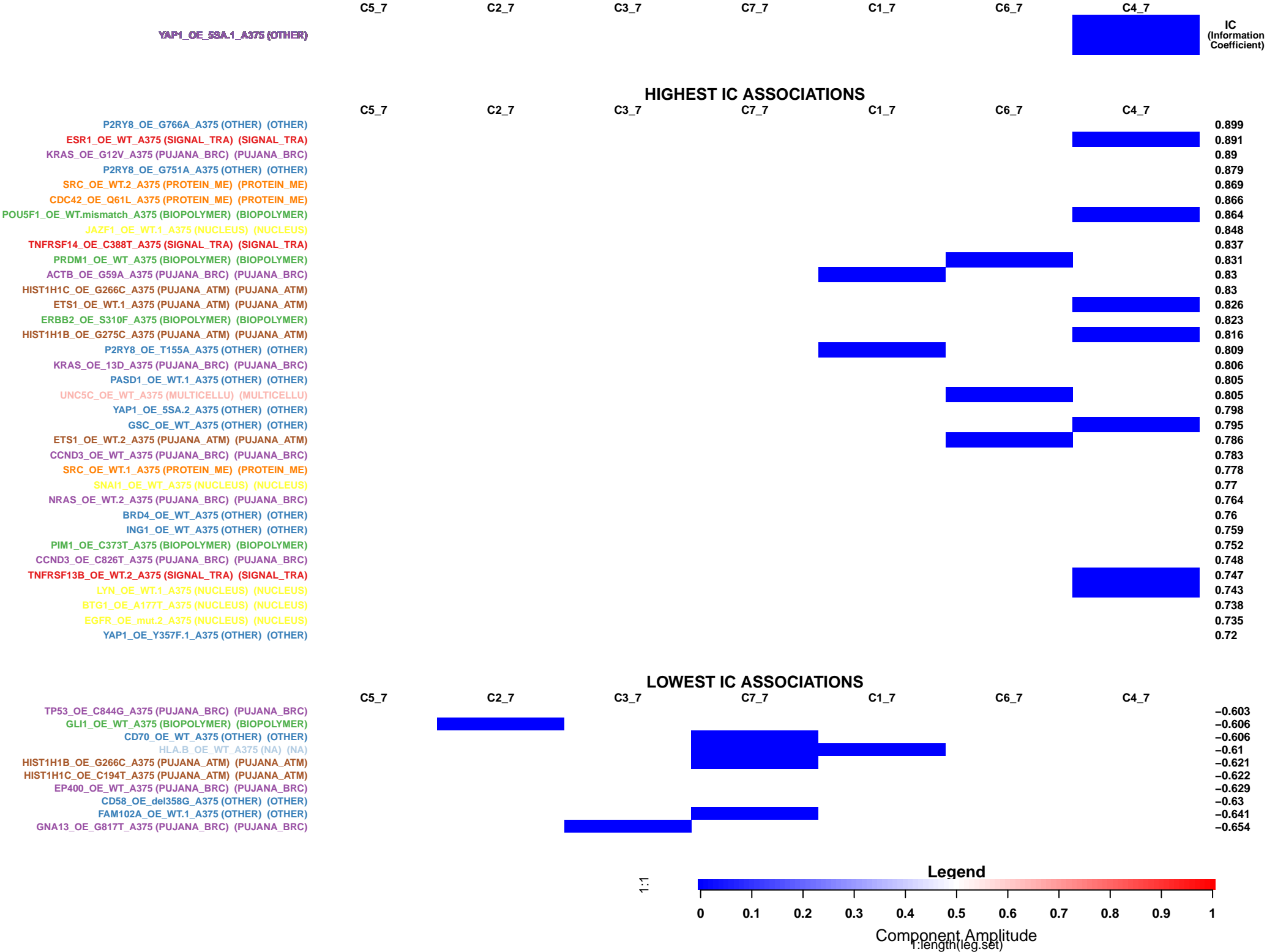


CDK8\_OE\_WT.1\_A375 (NUCLEUS)

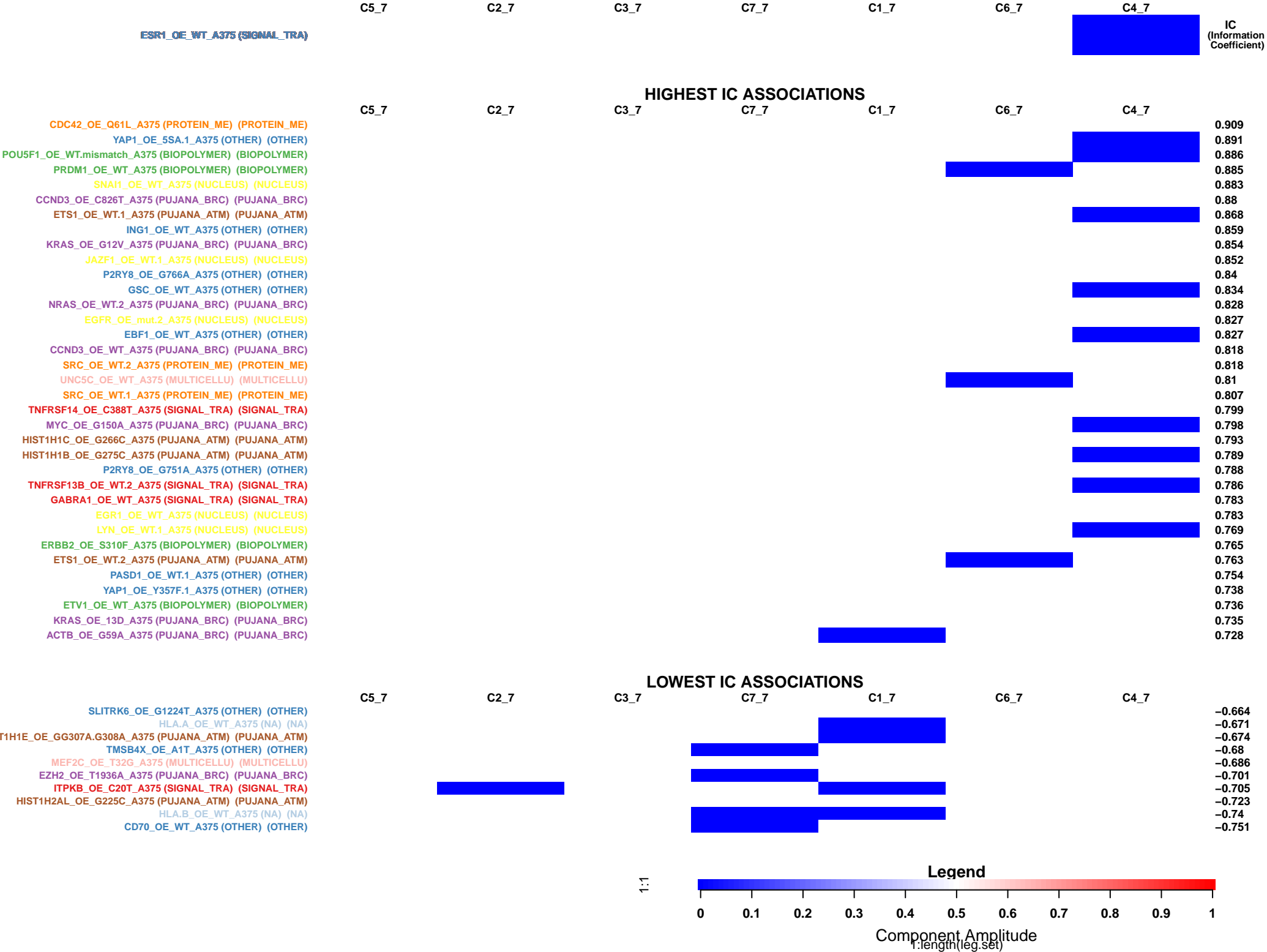




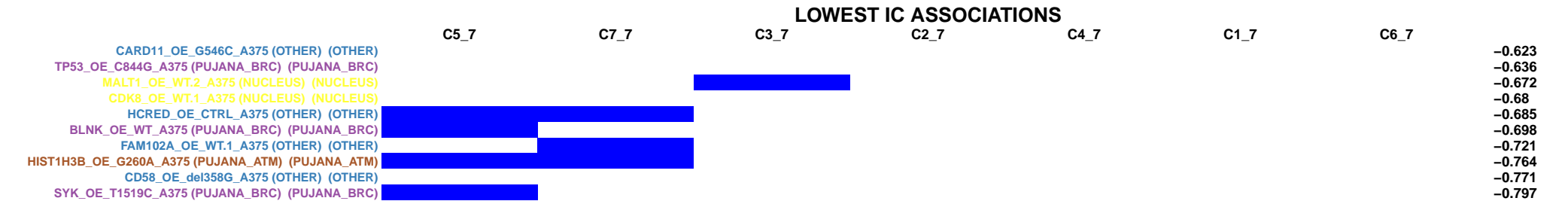
YAP1\_OE\_5SA.1\_A375 (OTHER)



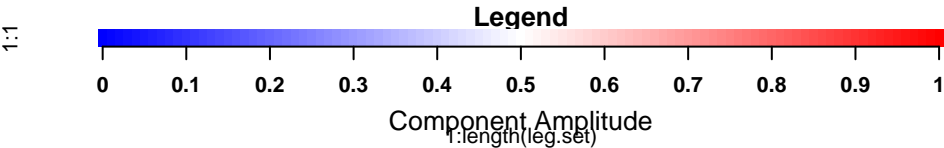
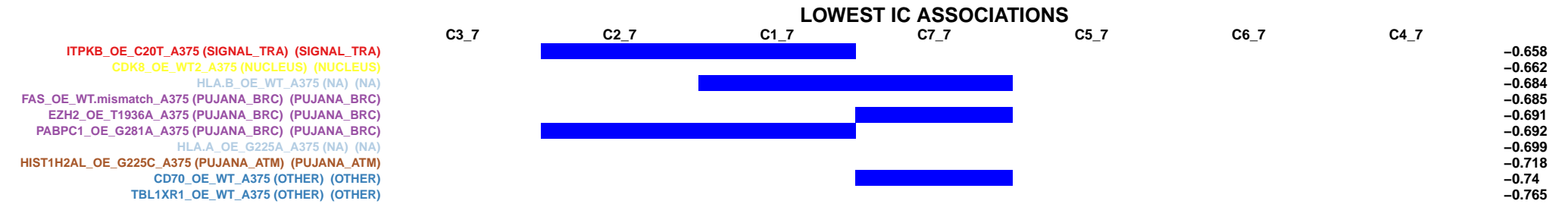
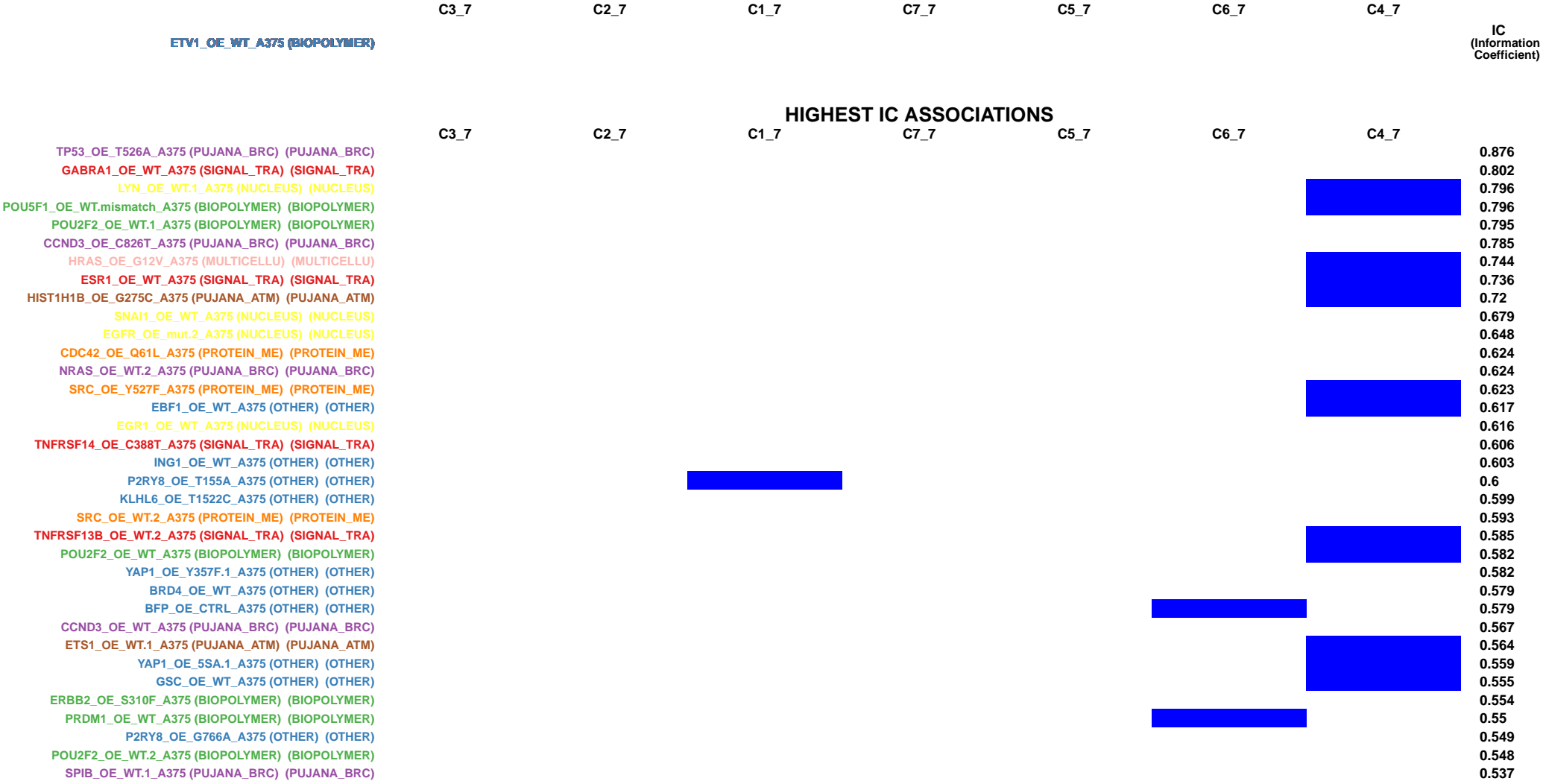
ESR1\_OE\_WT\_A375 (SIGNAL\_TRA)



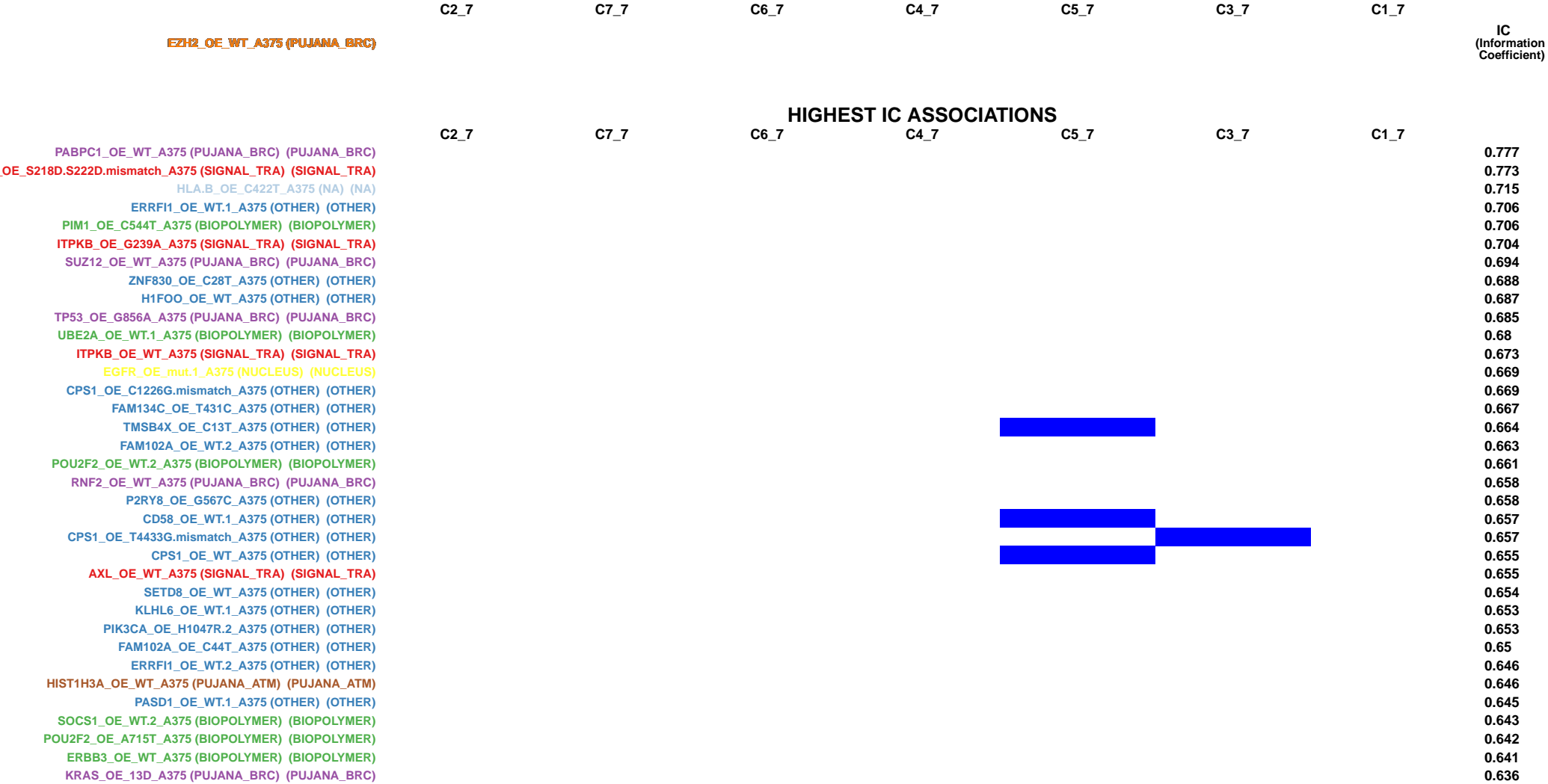
C5_7	C7_7	C3_7	C2_7	C4_7	C1_7	C6_7	IC (Information Coefficient)
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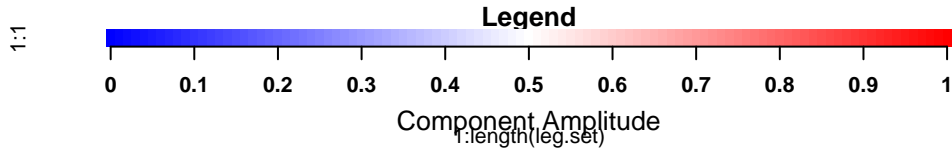
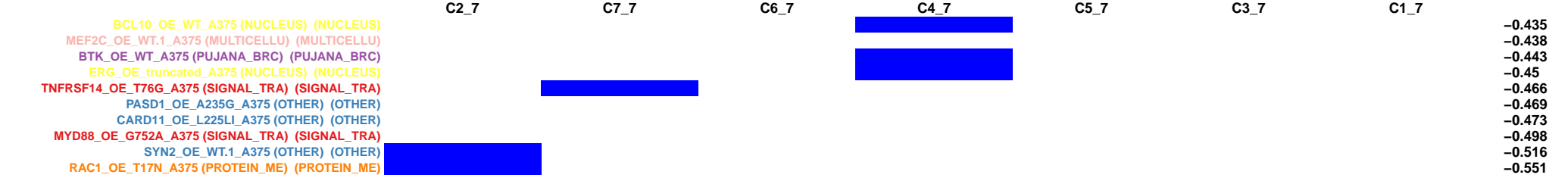
ETV1\_OE\_WT\_A375 (BIOPOLYMER)



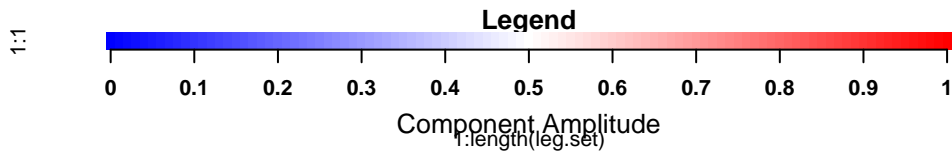
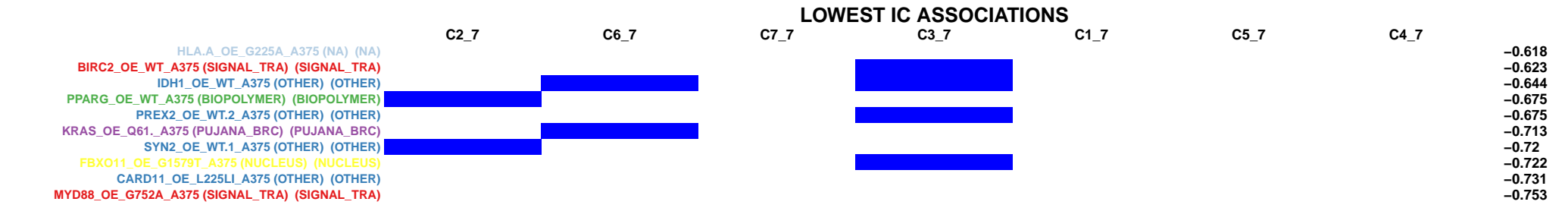
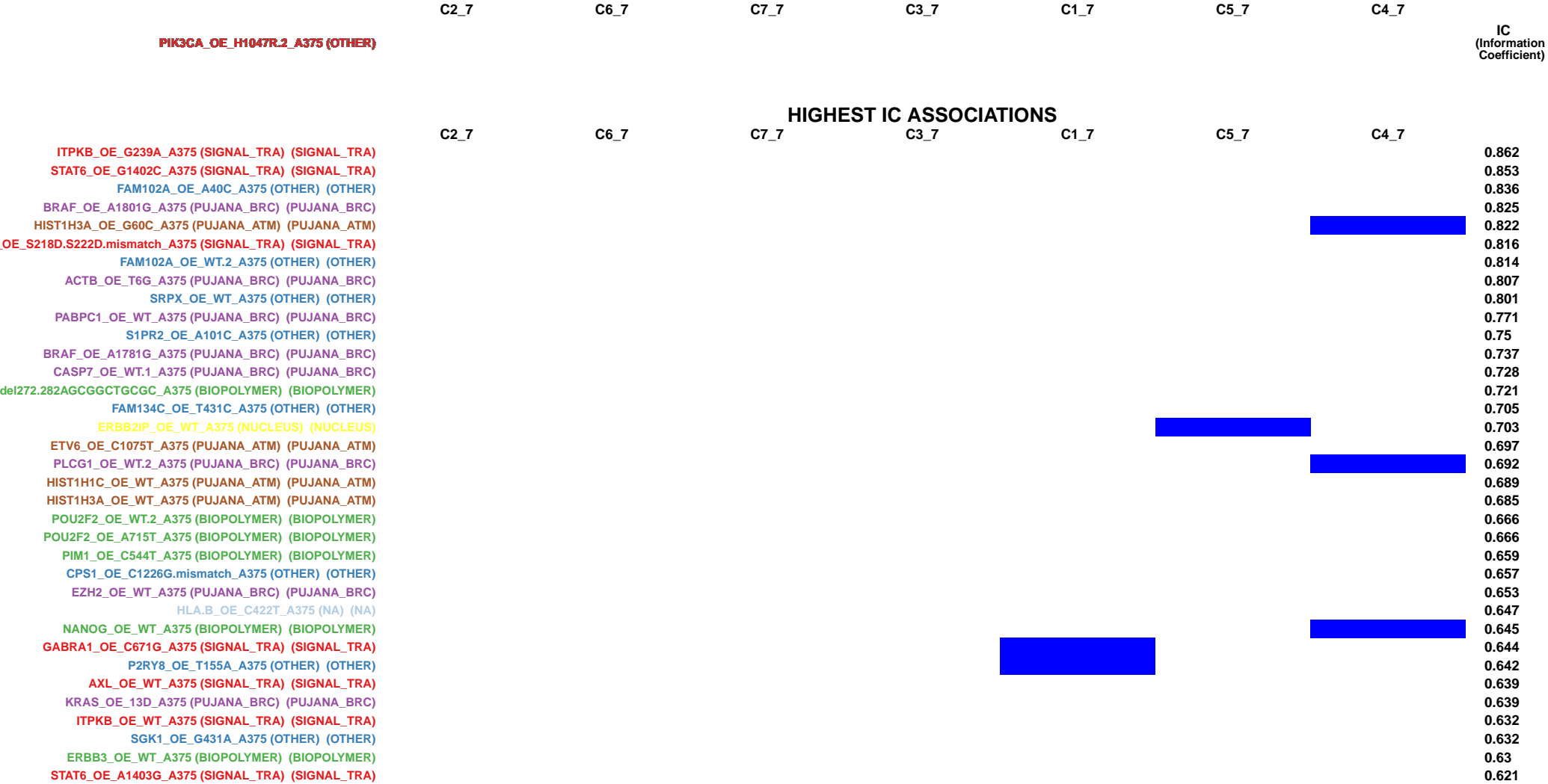
EZH2\_OE\_WT\_A375 (PUJANA\_BRC)



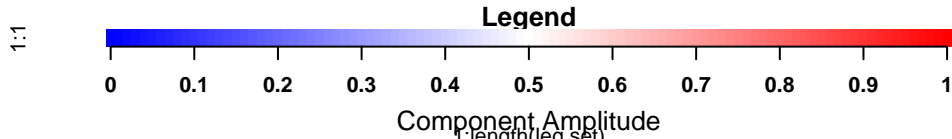
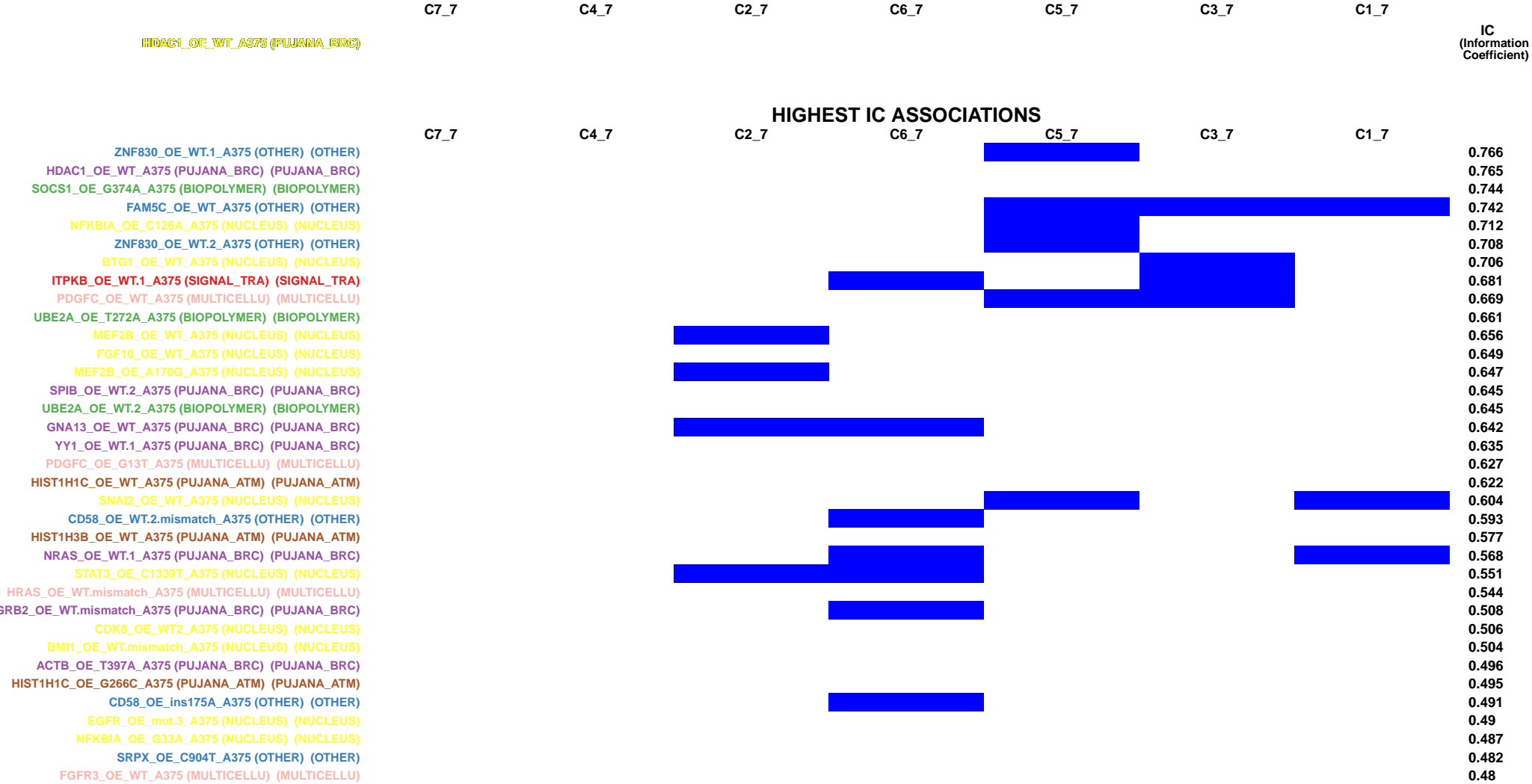
LOWEST IC ASSOCIATIONS



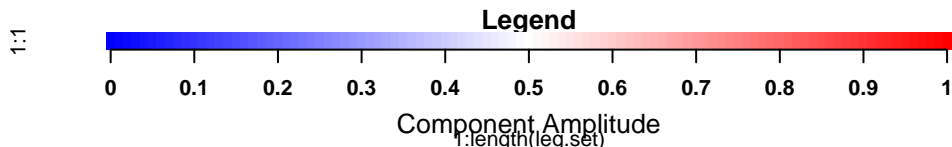
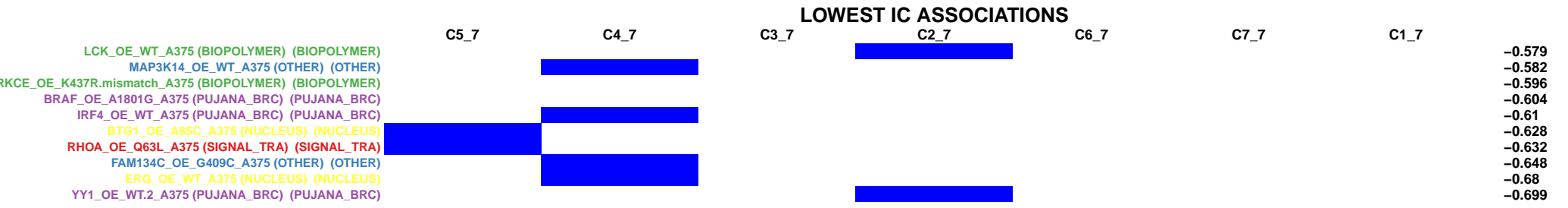
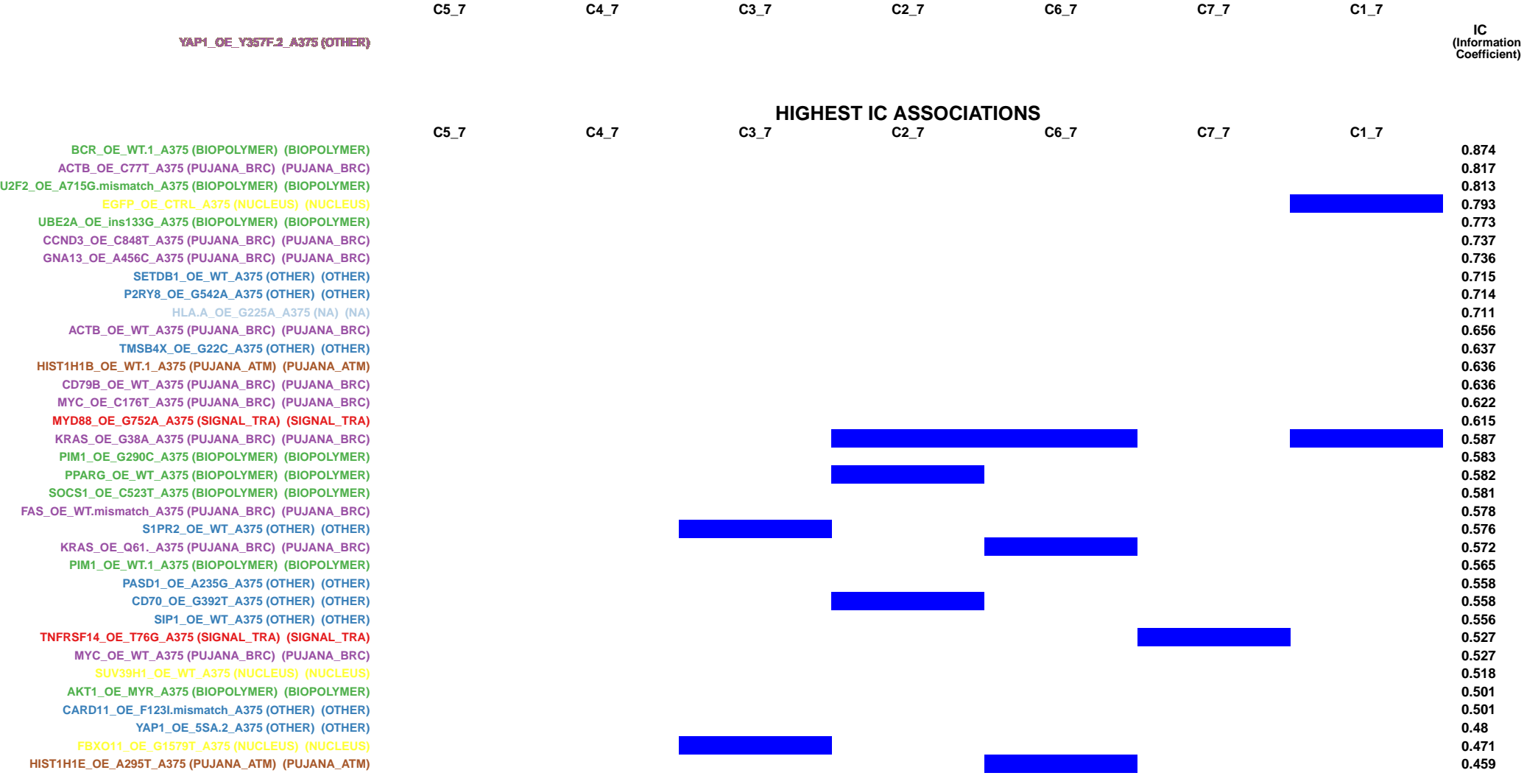
PIK3CA\_OE\_H1047R.2\_A375 (OTHER)



HDAC1\_OE\_WT\_A375 (PUJANA\_BRC)

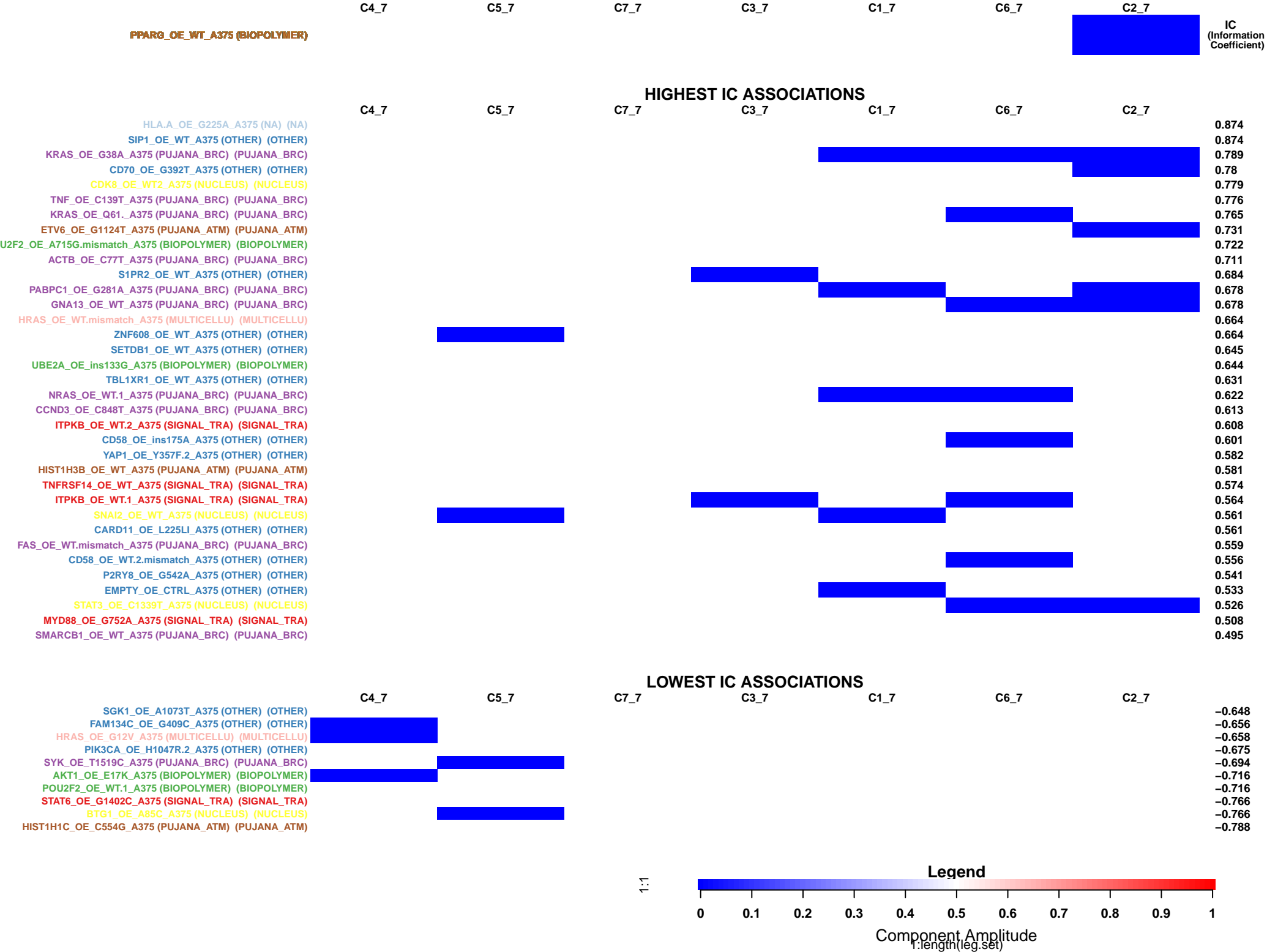


YAP1\_OE\_Y357F.2\_A375 (OTHER)



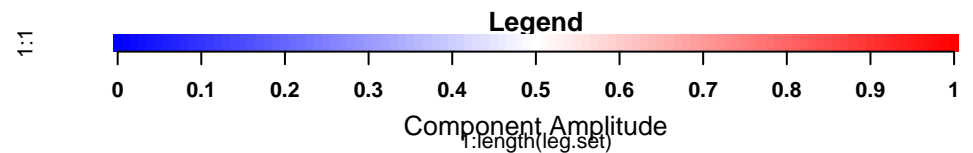


PPARG\_OE\_WT\_A375 (BIOPOLYMER)

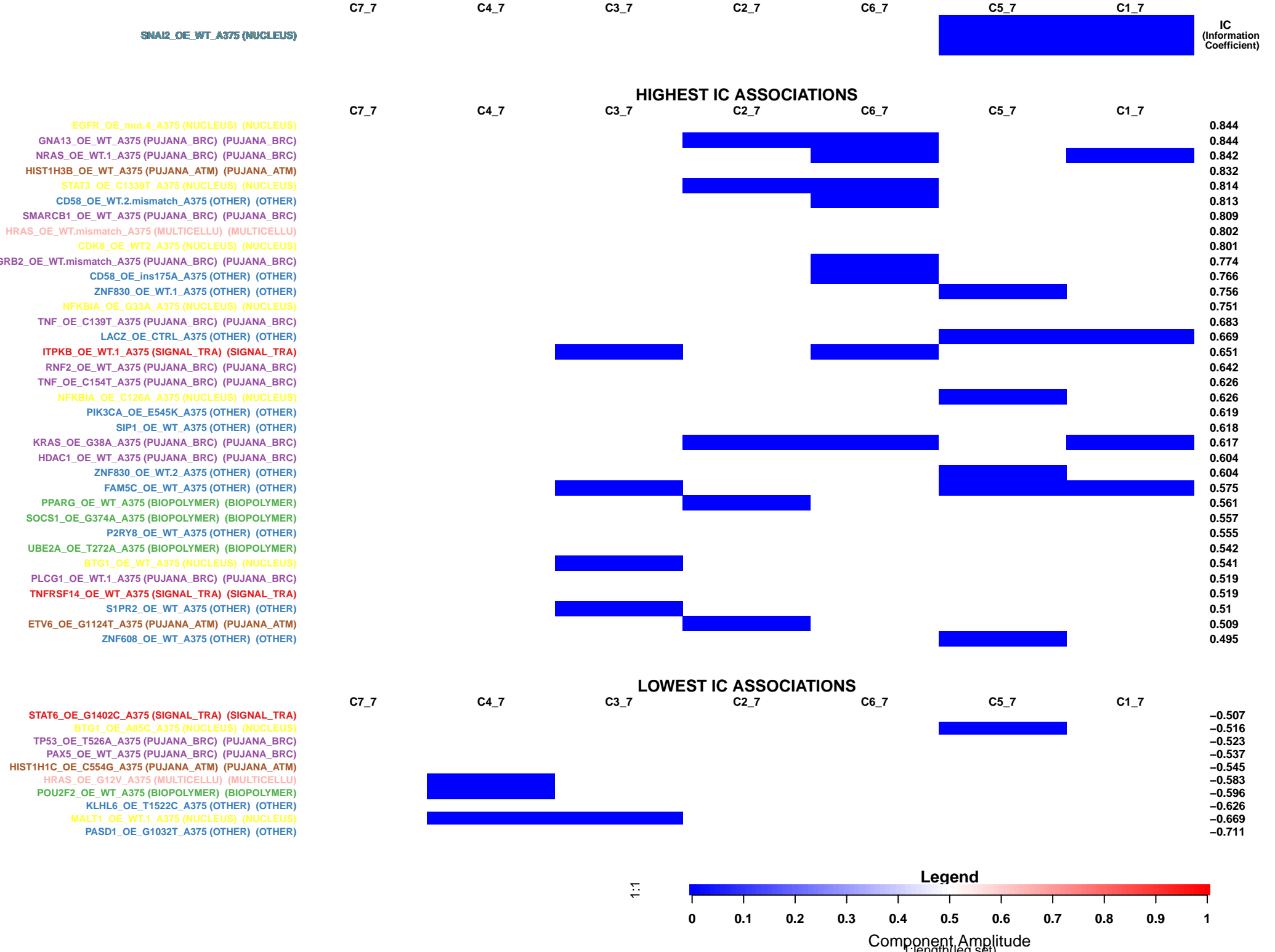


C5_7	C4_7	C3_7	C7_7	C1_7	C2_7	C6_7	IC (Information Coefficient)

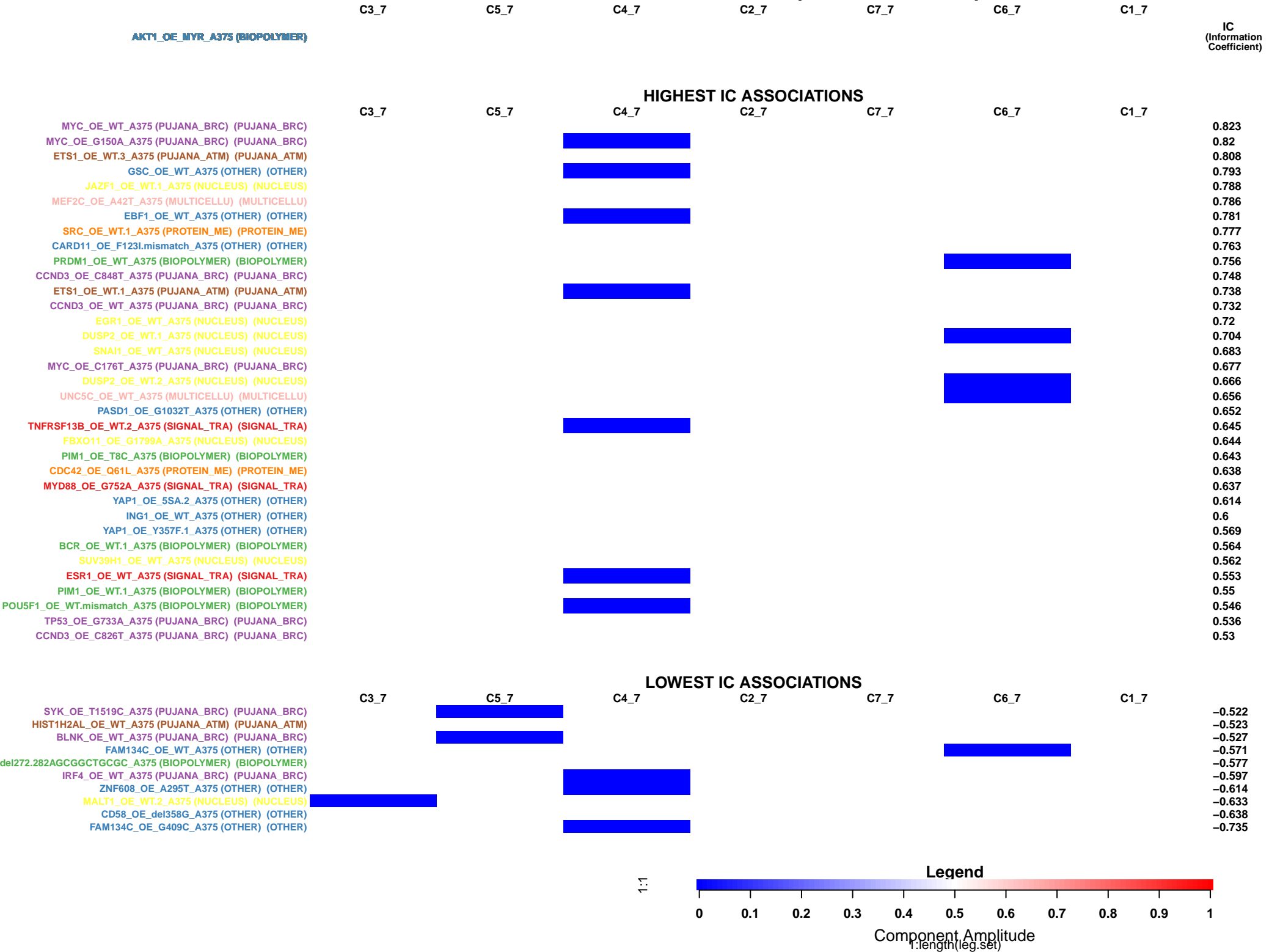
C5_7	C4_7	C3_7	C7_7	C1_7	C2_7	C6_7	
							0.842
							0.82
							0.788
							0.765
							0.721
							0.714
							0.691
							0.668
							0.643
							0.642
							0.639
							0.631
							0.624
							0.624
							0.623
							0.623
							0.61
							0.592
							0.592
							0.59
							0.589
							0.583
							0.577
							0.576
							0.575
							0.572
							0.57
							0.524
							0.524
							0.512
							0.499
							0.488
							0.484
							0.481
							0.475

[illegible]

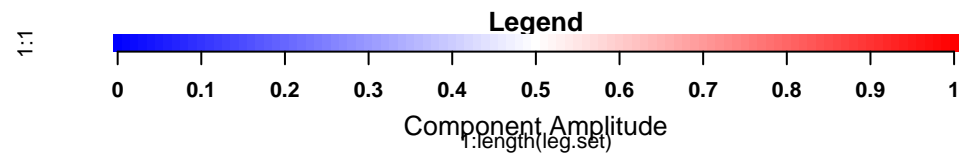
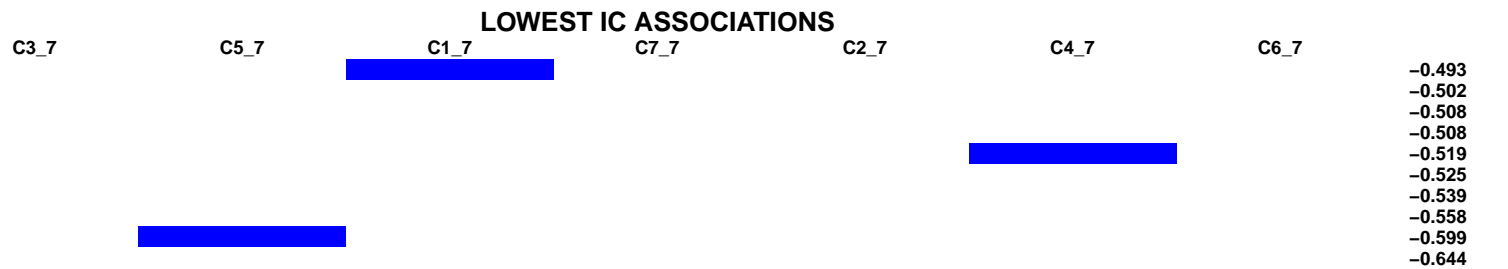
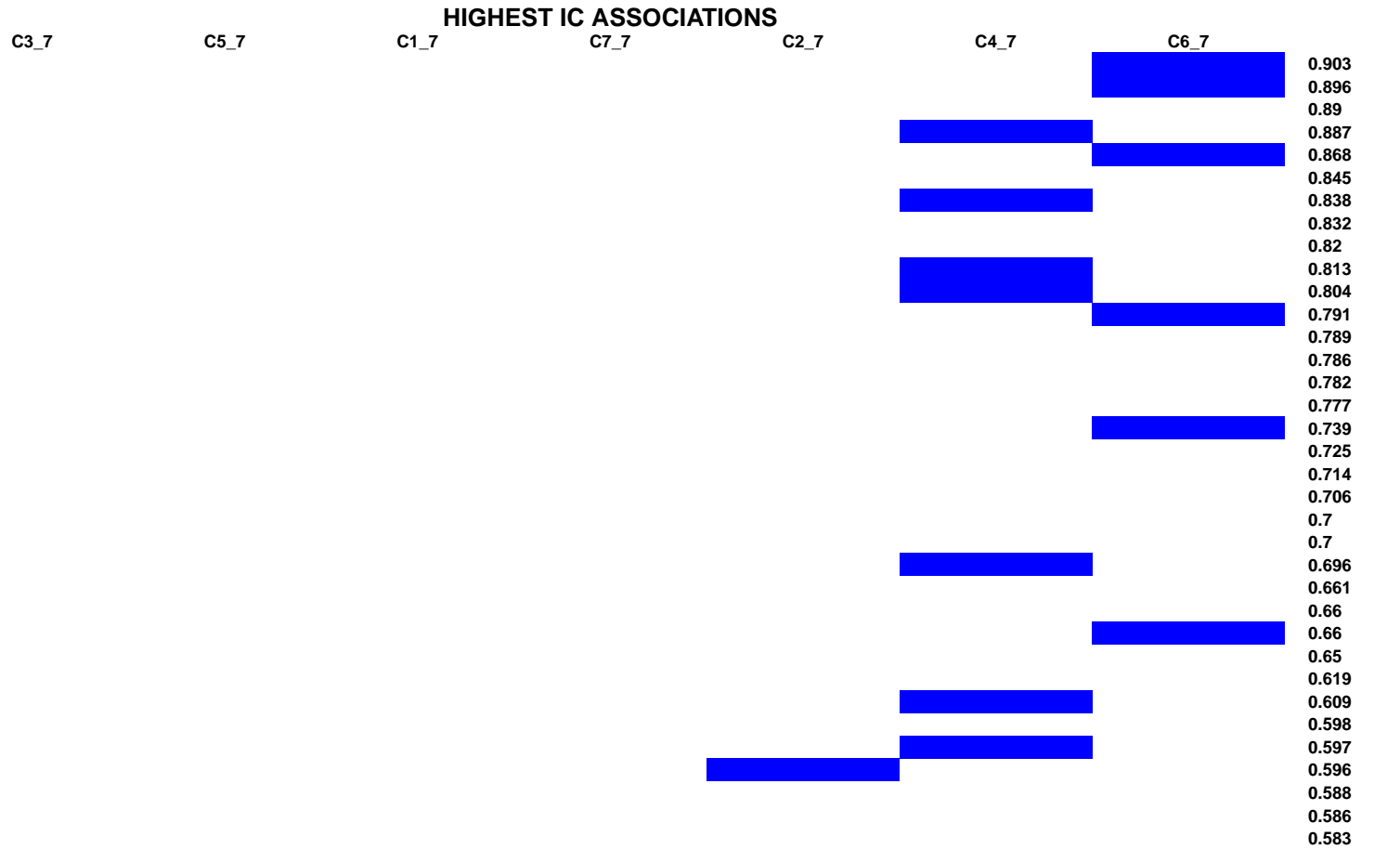
SNAI2\_OE\_WT\_A375 (NUCLEUS)



# AKT1\_OE\_MYR\_A375 (BIOPOLYMER)



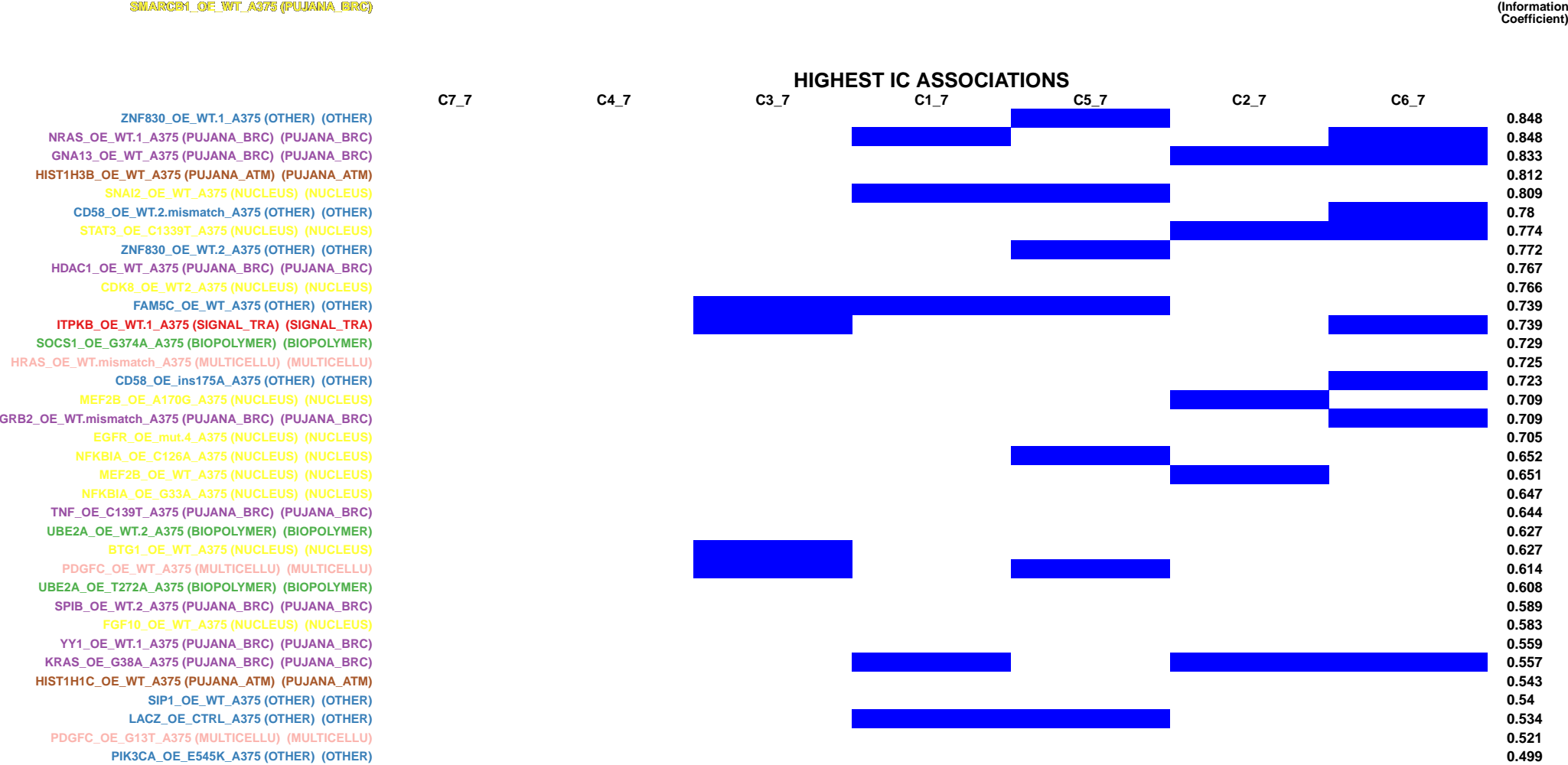
C3_7	C5_7	C1_7	C7_7	C2_7	C4_7	C6_7	IC (Information Coefficient)
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SMARCB1\_OE\_WT\_A375 (PUJANA\_BRC)

C7\_7 C4\_7 C3\_7 C1\_7 C5\_7 C2\_7 C6\_7

IC  
(Information  
Coefficient)

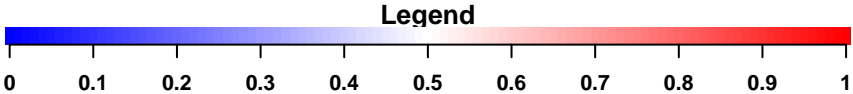


LOWEST IC ASSOCIATIONS

C7\_7 C4\_7 C3\_7 C1\_7 C5\_7 C2\_7 C6\_7



1:1



Component Amplitude

1:length(leg.set)

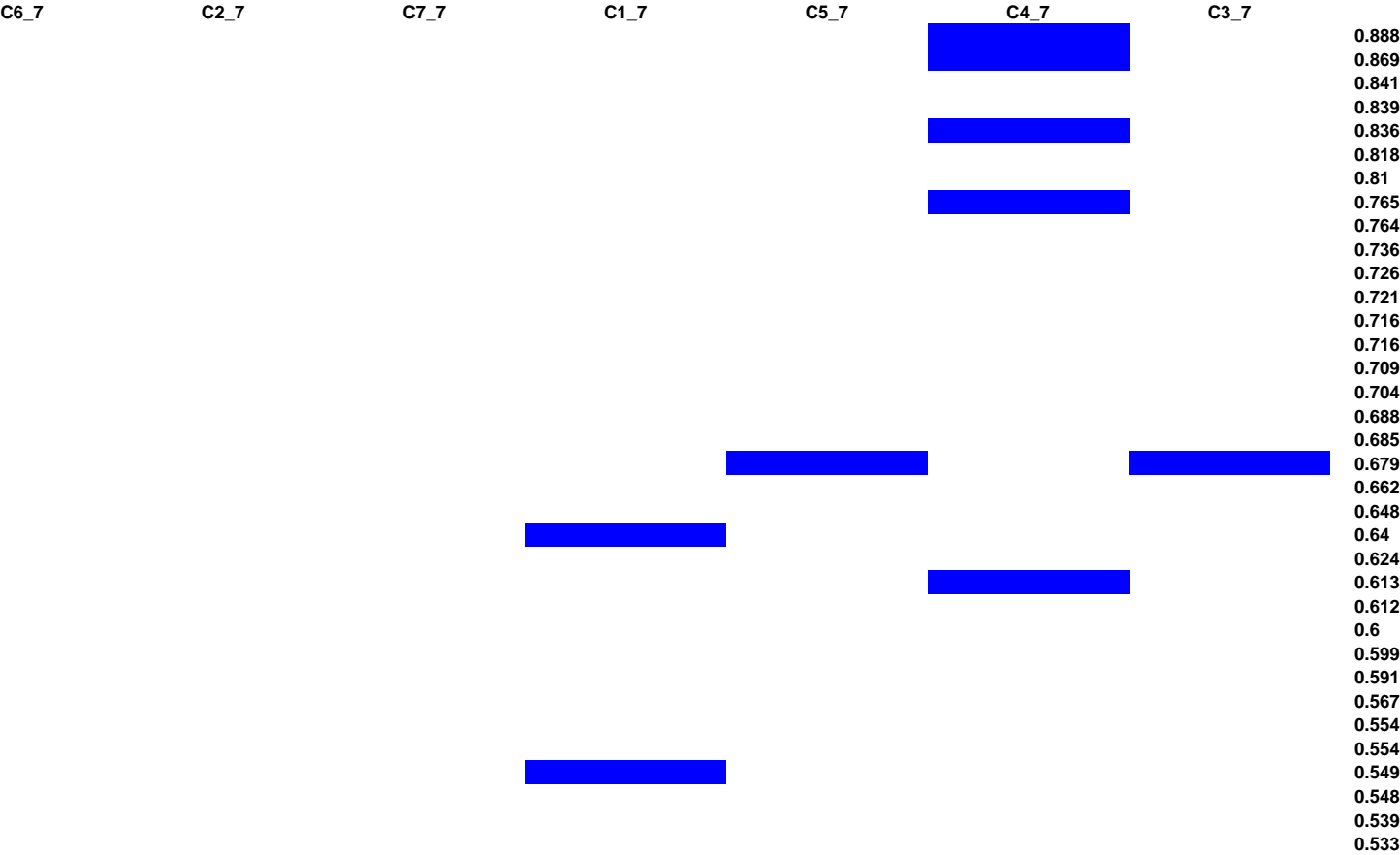
SOCS1\_OE\_del272.282AGCGGCTGCGC\_A375 (BIOPOLYMER)

C6\_7C2\_7C7\_7C1\_7C5\_7C4\_7C3\_7

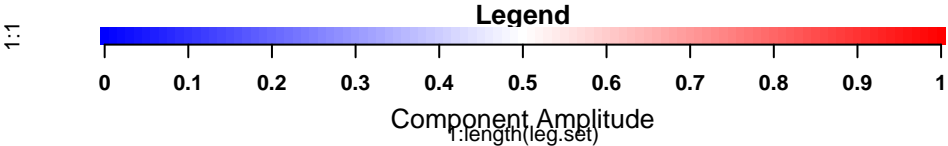
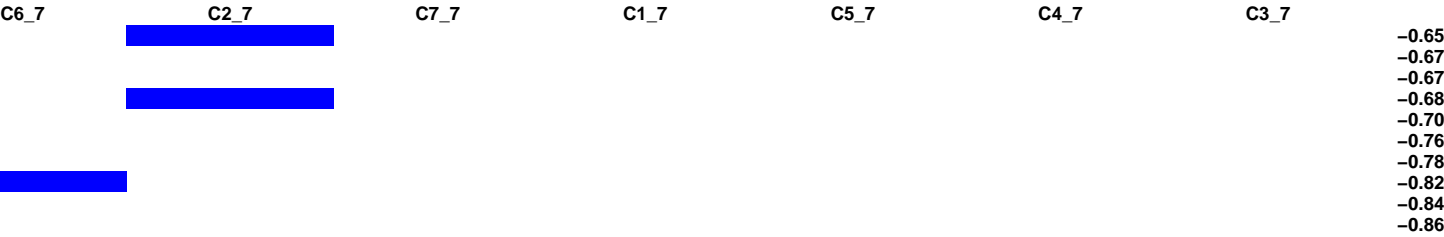
SOCS1\_OE\_del272.282AGCGGCTGCGC\_A375 (BIOPOLYMER)

IC  
(Information  
Coefficient)

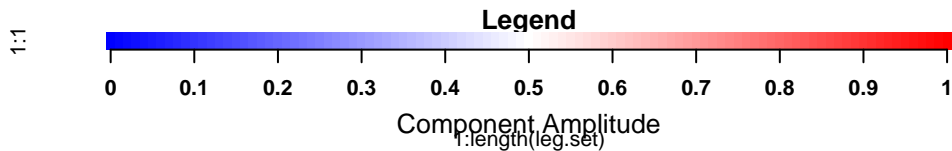
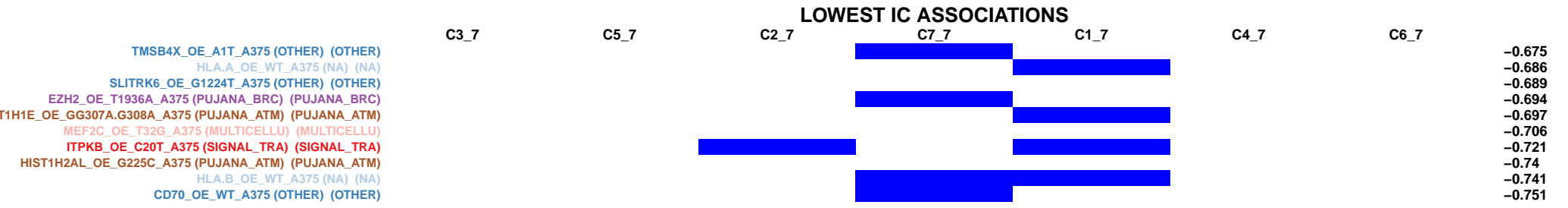
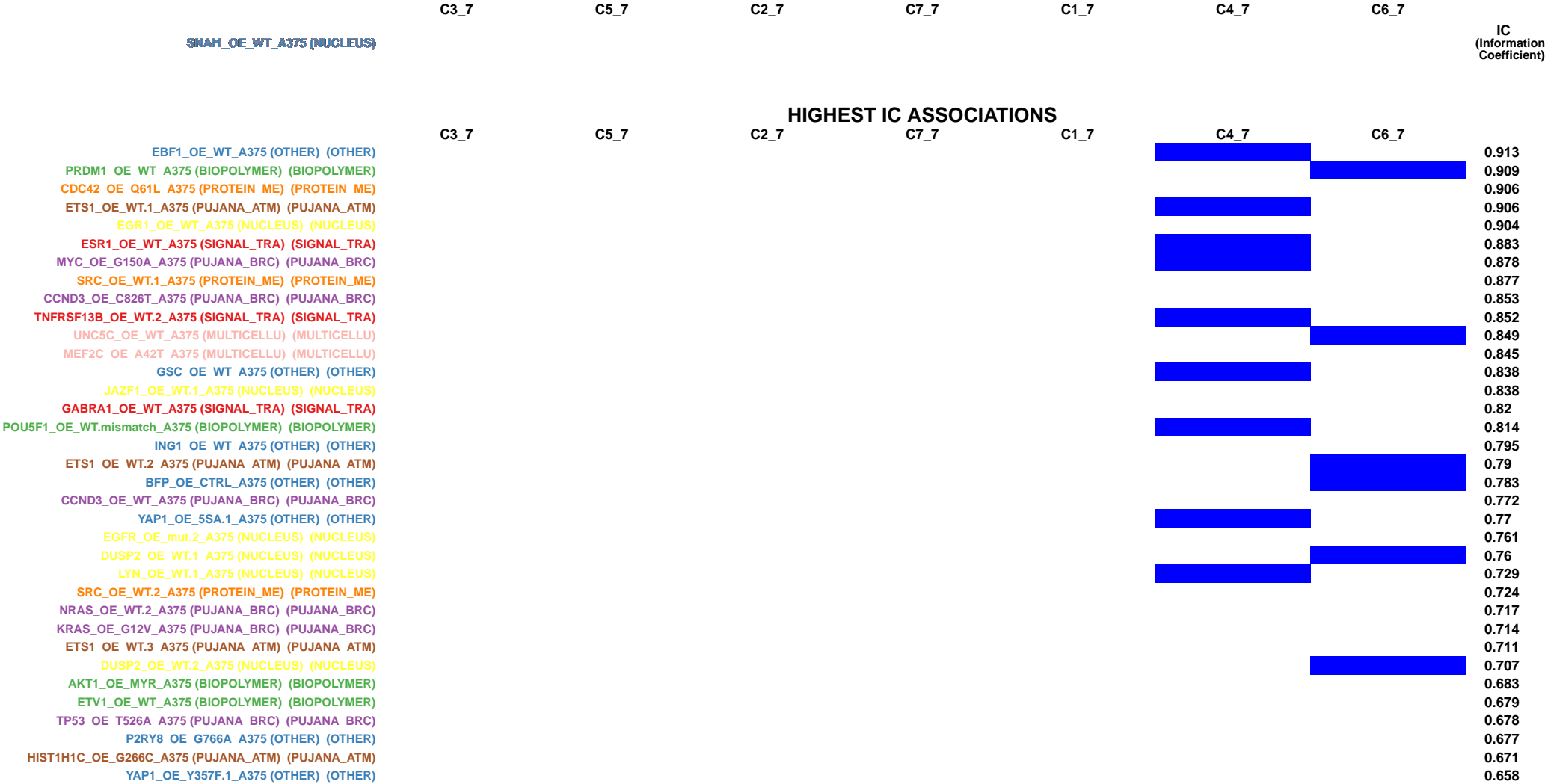
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS

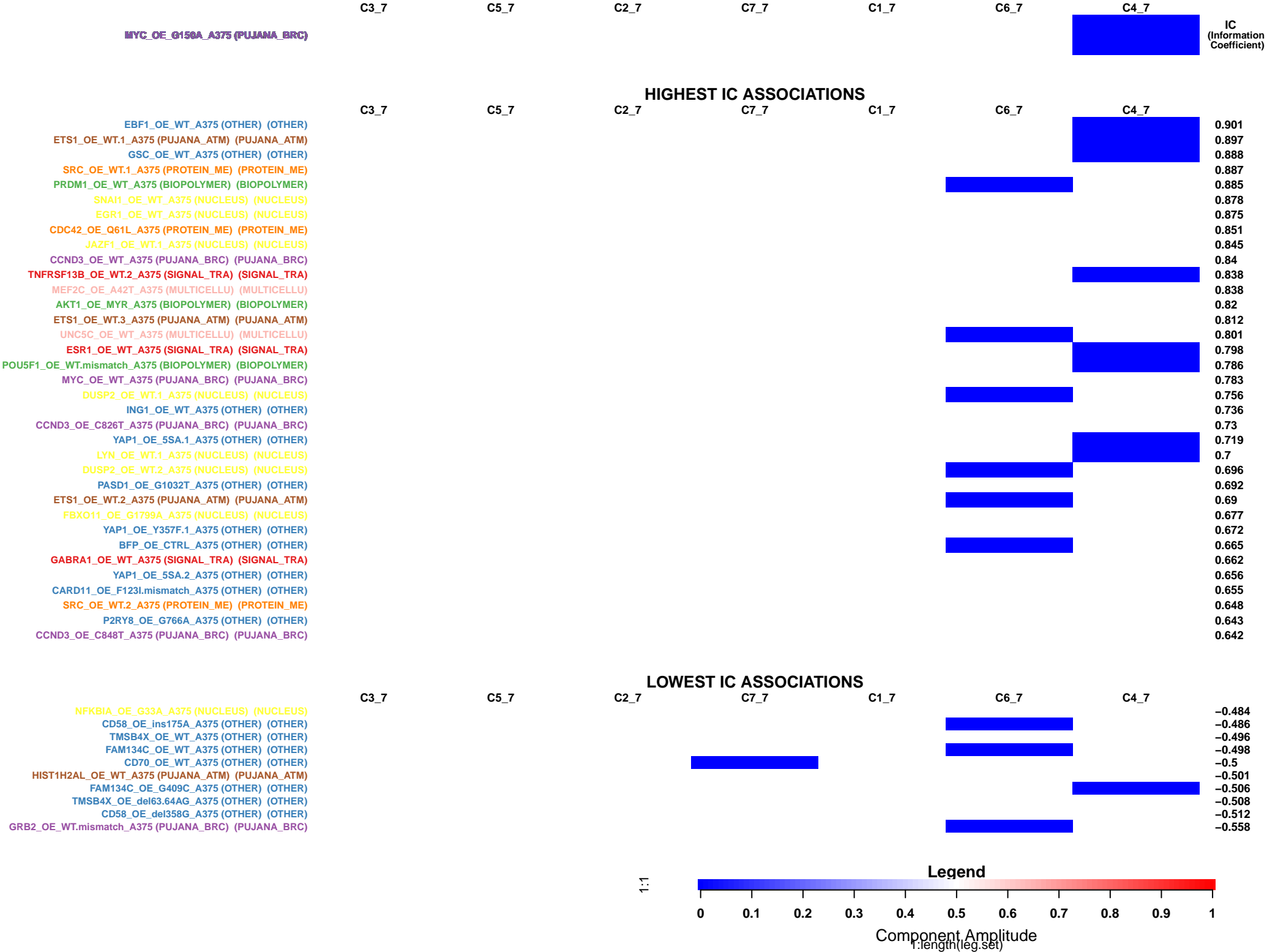


SNAI1\_OE\_WT\_A375 (NUCLEUS)

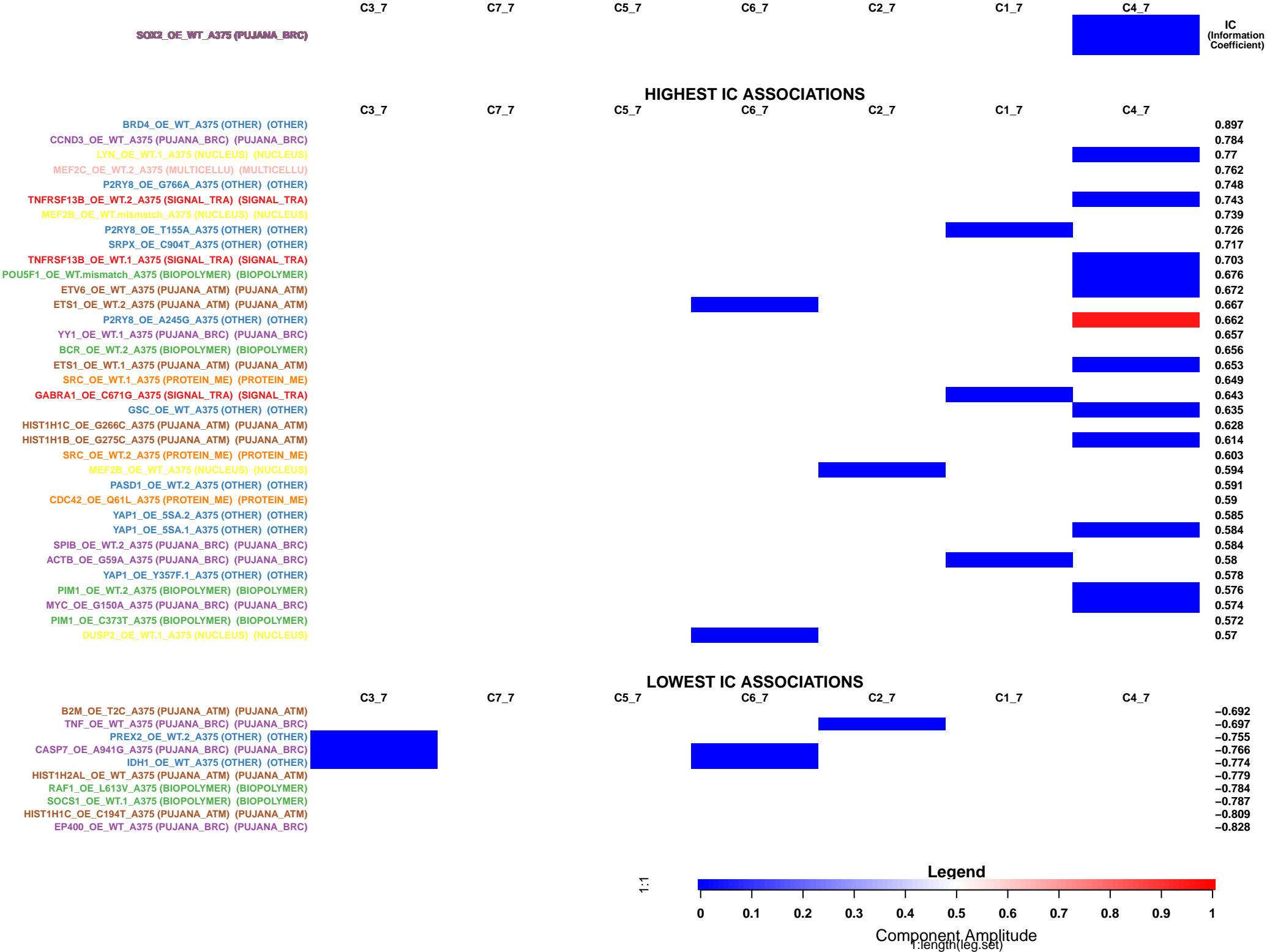




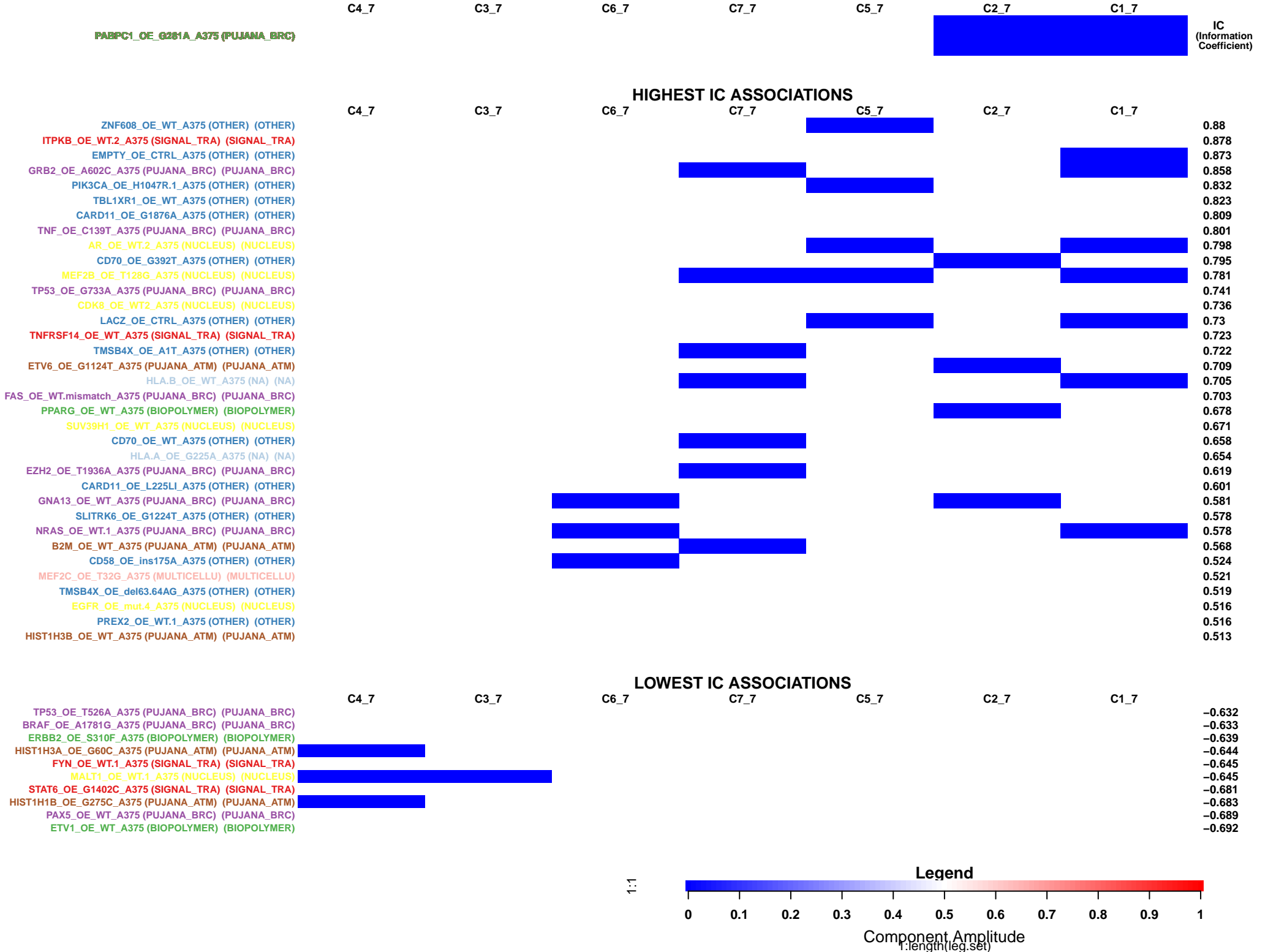
MYC\_OE\_G150A\_A375 (PUJANA\_BRC)



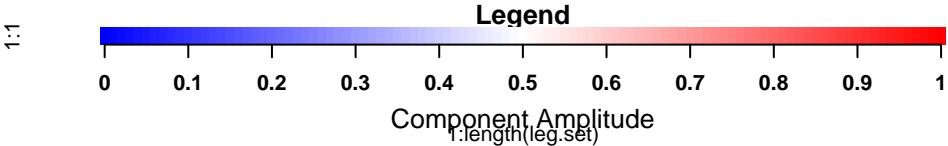
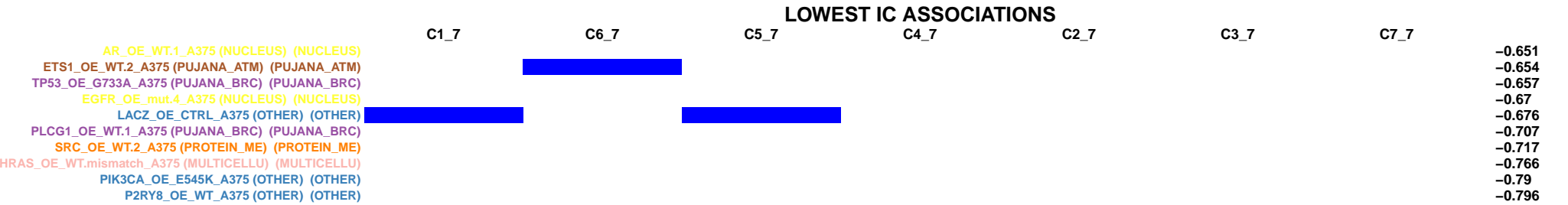
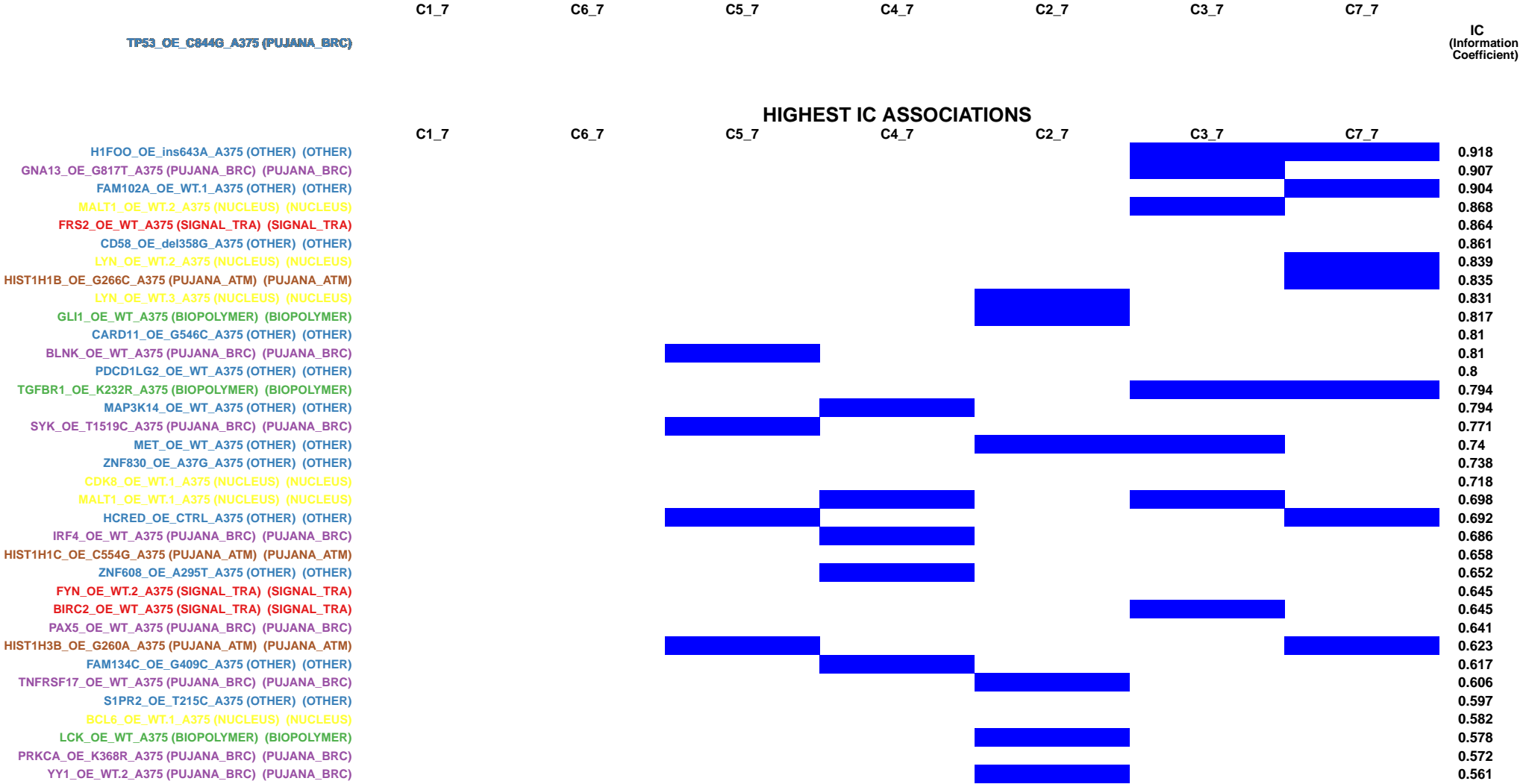
SOX2\_OE\_WT\_A375 (PUJANA\_BRC)



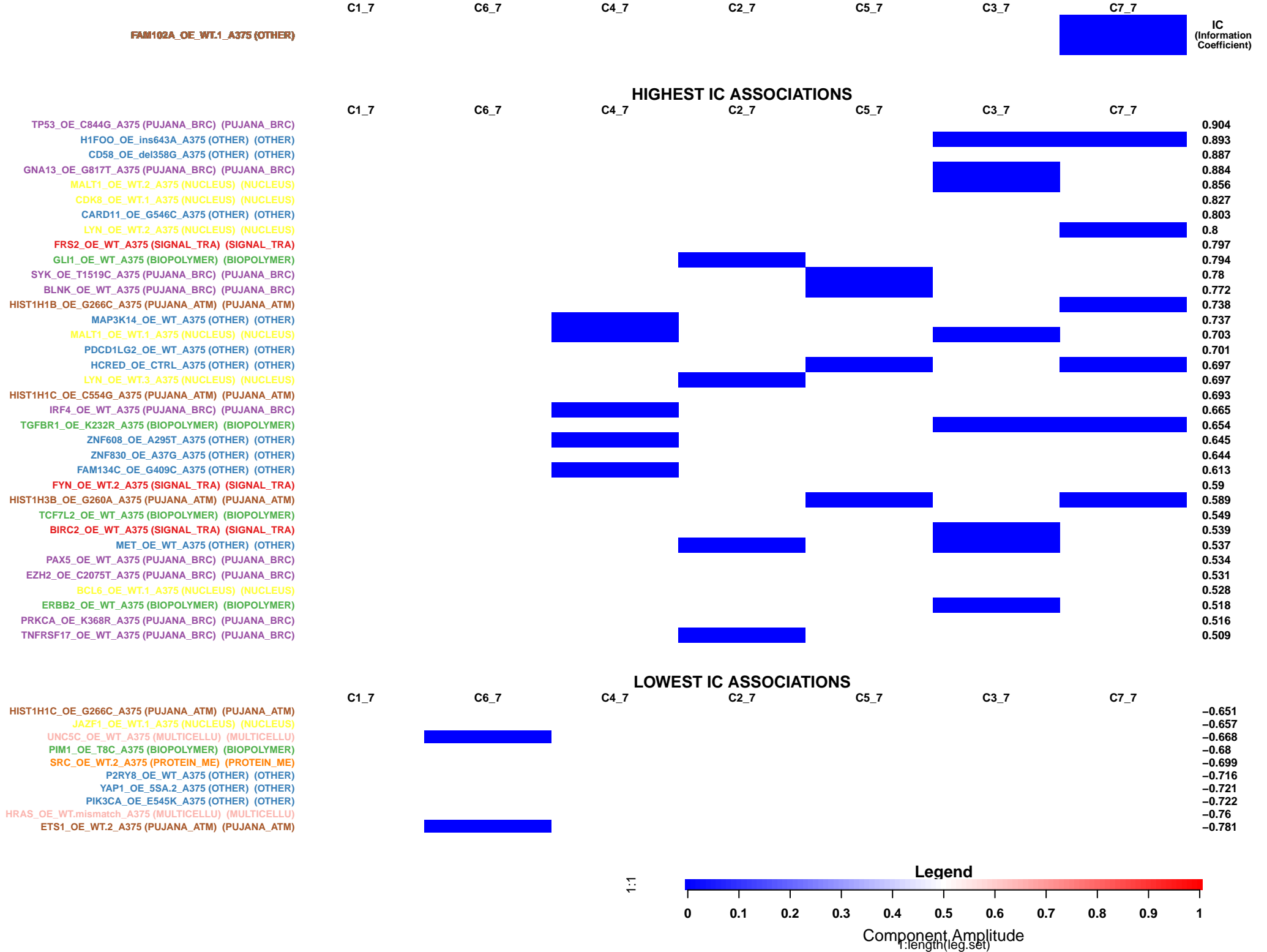
PABPC1\_OE\_G281A\_A375 (PUJANA\_BRC)



TP53\_OE\_C844G\_A375 (PUJANA\_BRC)



FAM102A\_OE\_WT.1\_A375 (OTHER)

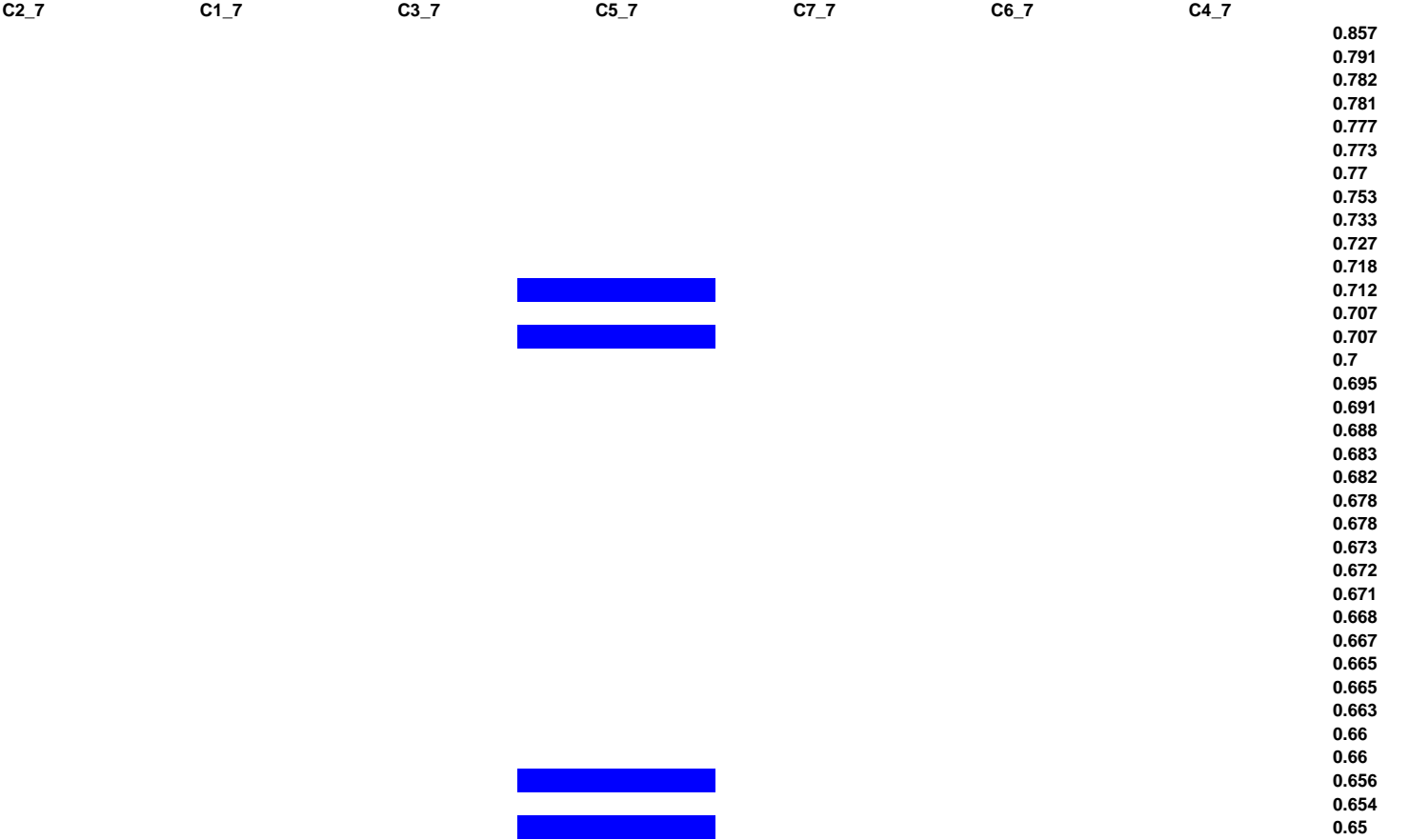


ZNF830\_OE\_C28T\_A375 (OTHER)

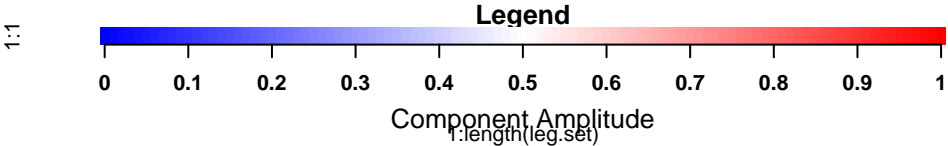
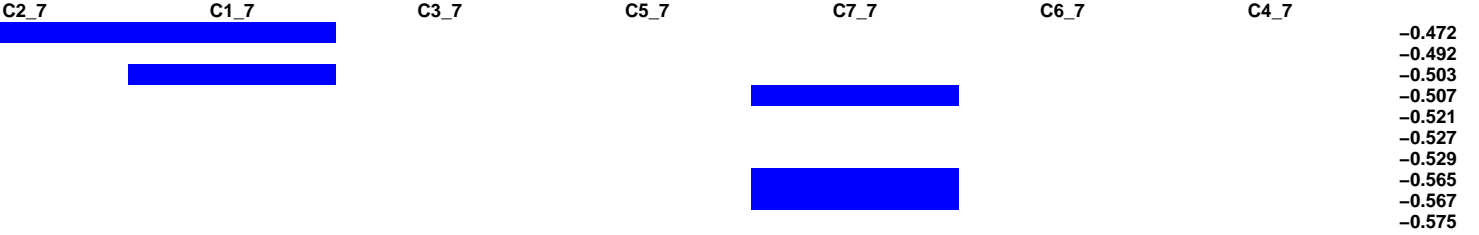
C2\_7 C1\_7 C3\_7 C5\_7 C7\_7 C6\_7 C4\_7 IC (Information Coefficient)

ZNF830\_OE\_C28T\_A375 (OTHER)

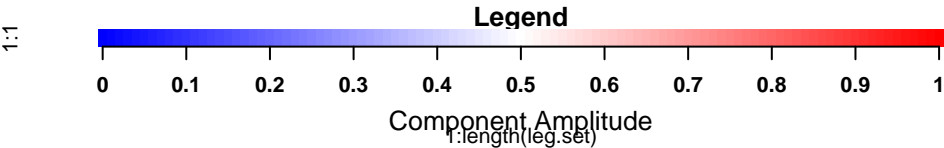
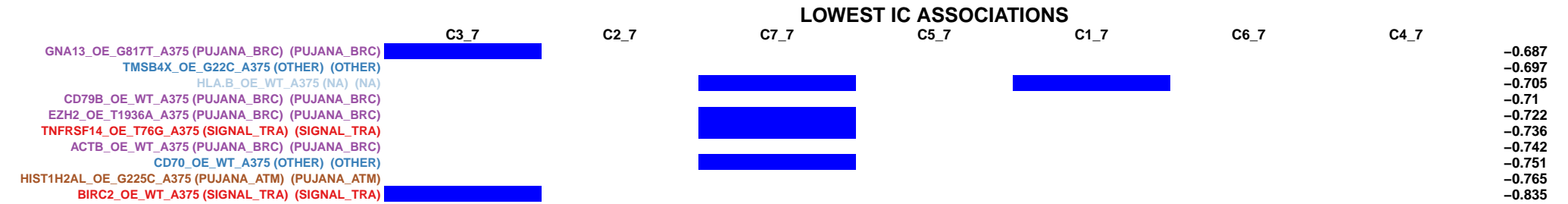
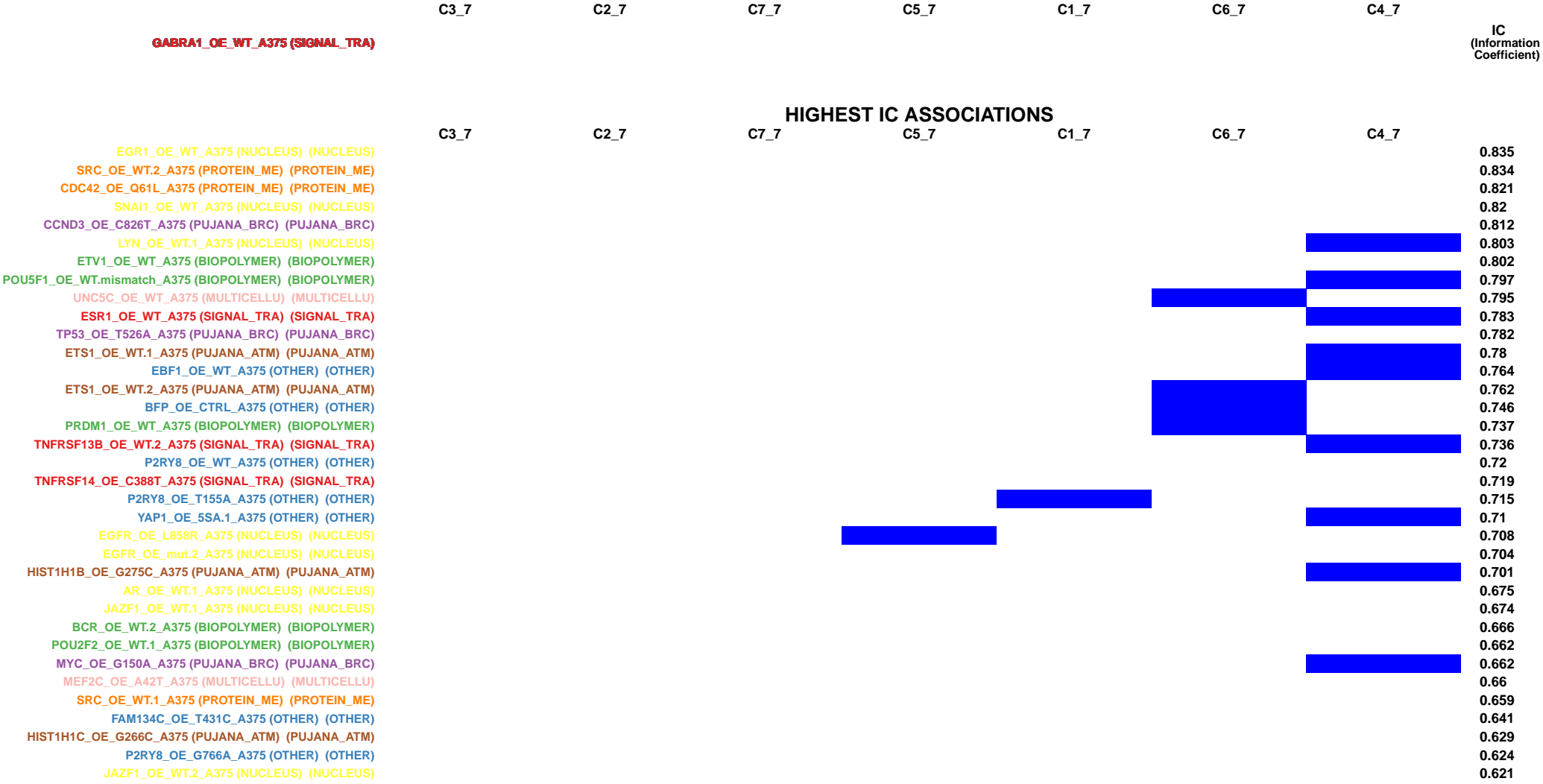
HIGHEST IC ASSOCIATIONS



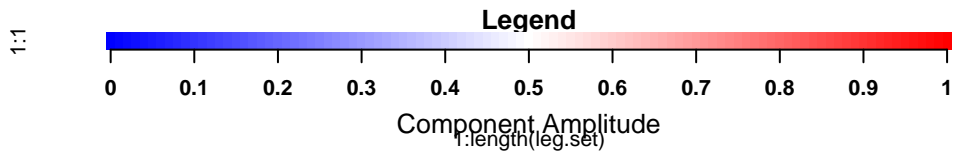
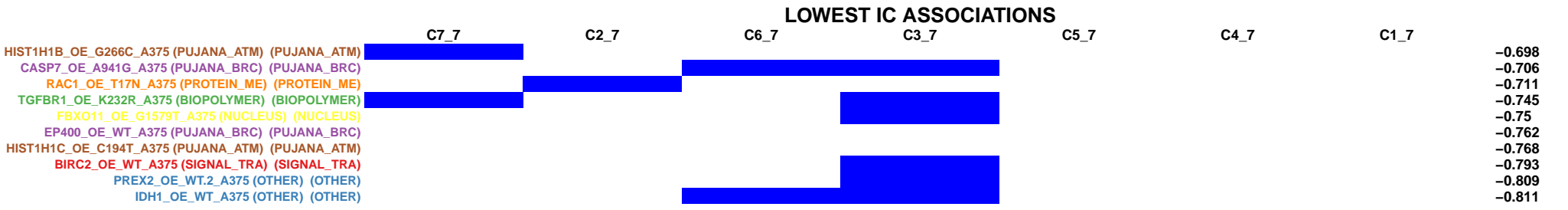
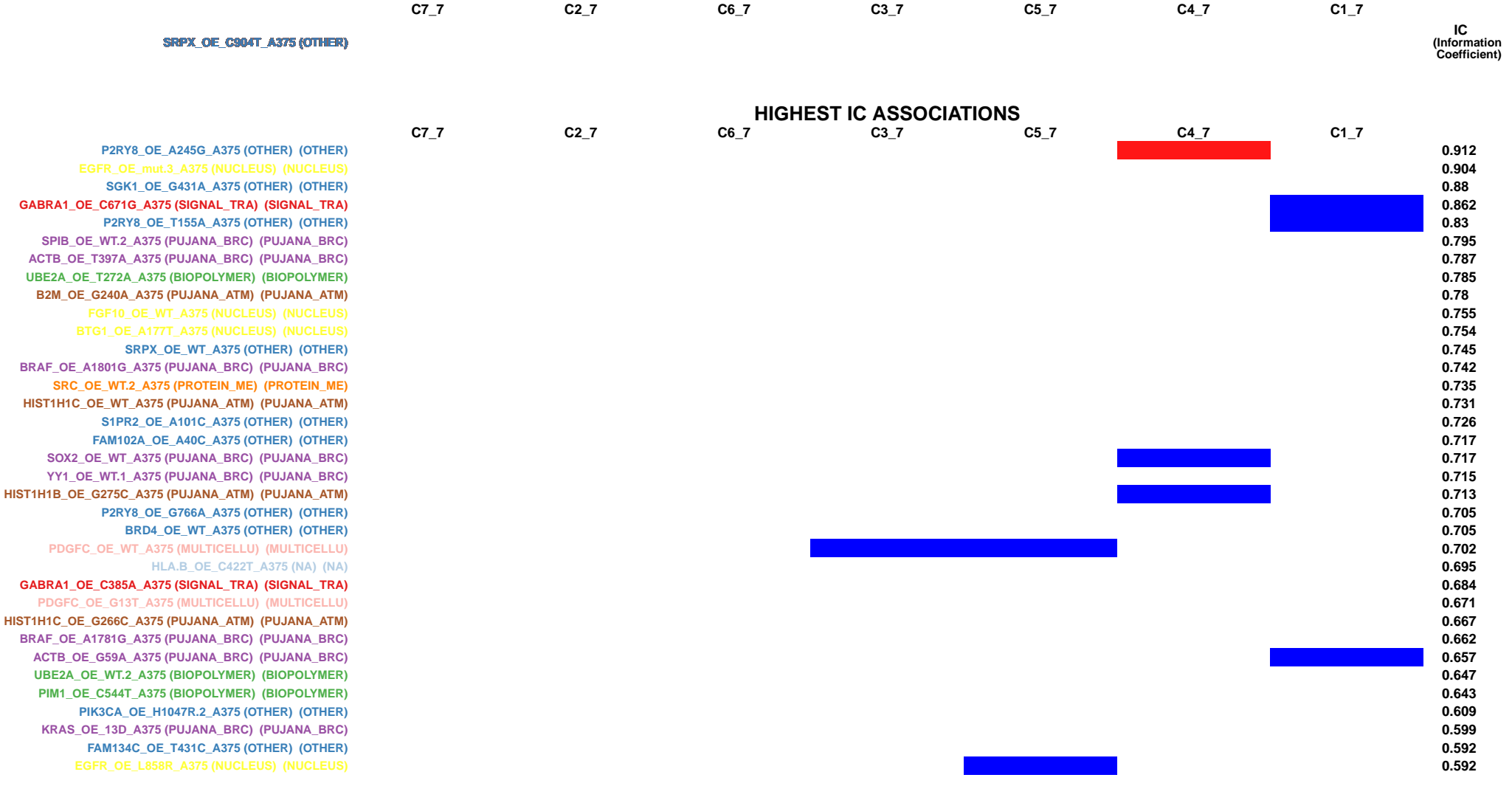
LOWEST IC ASSOCIATIONS



GABRA1\_OE\_WT\_A375 (SIGNAL\_TRA)

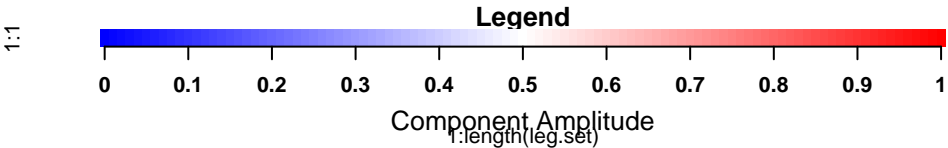
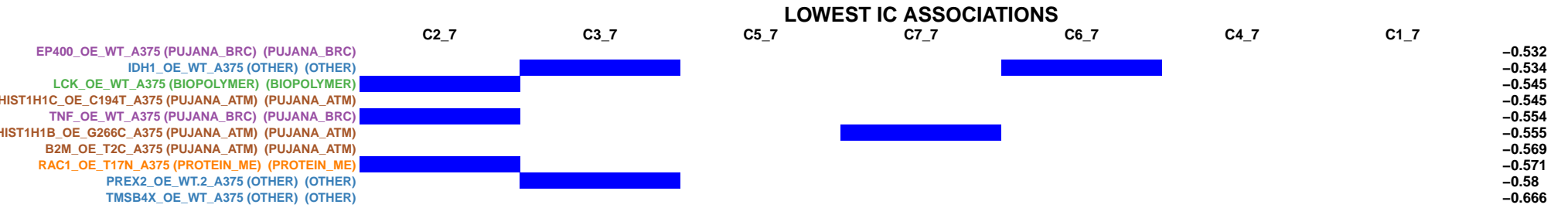
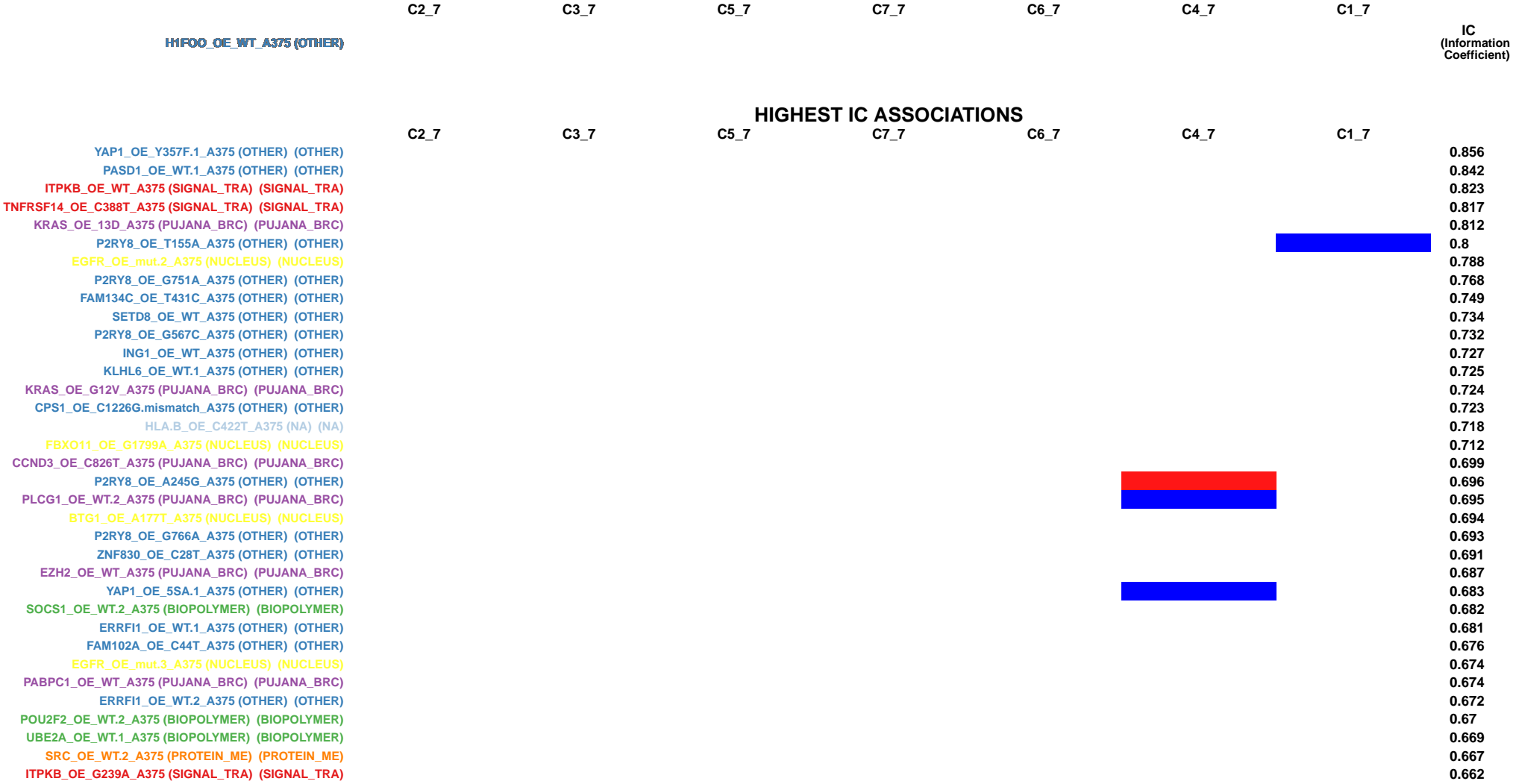


SRPX\_OE\_C904T\_A375 (OTHER)

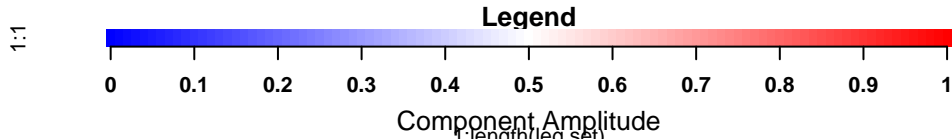
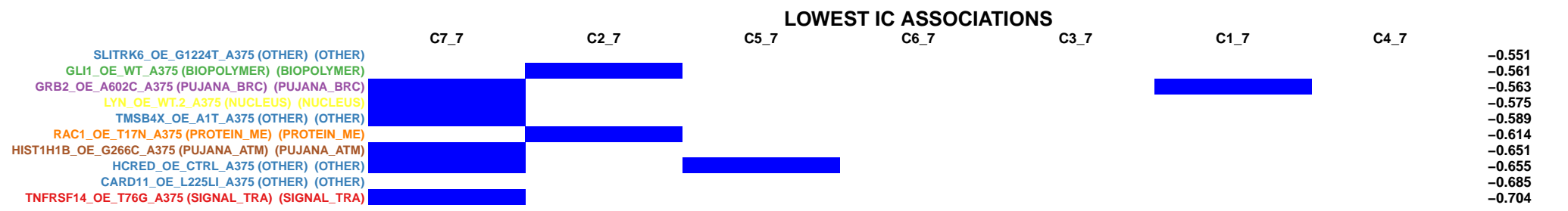
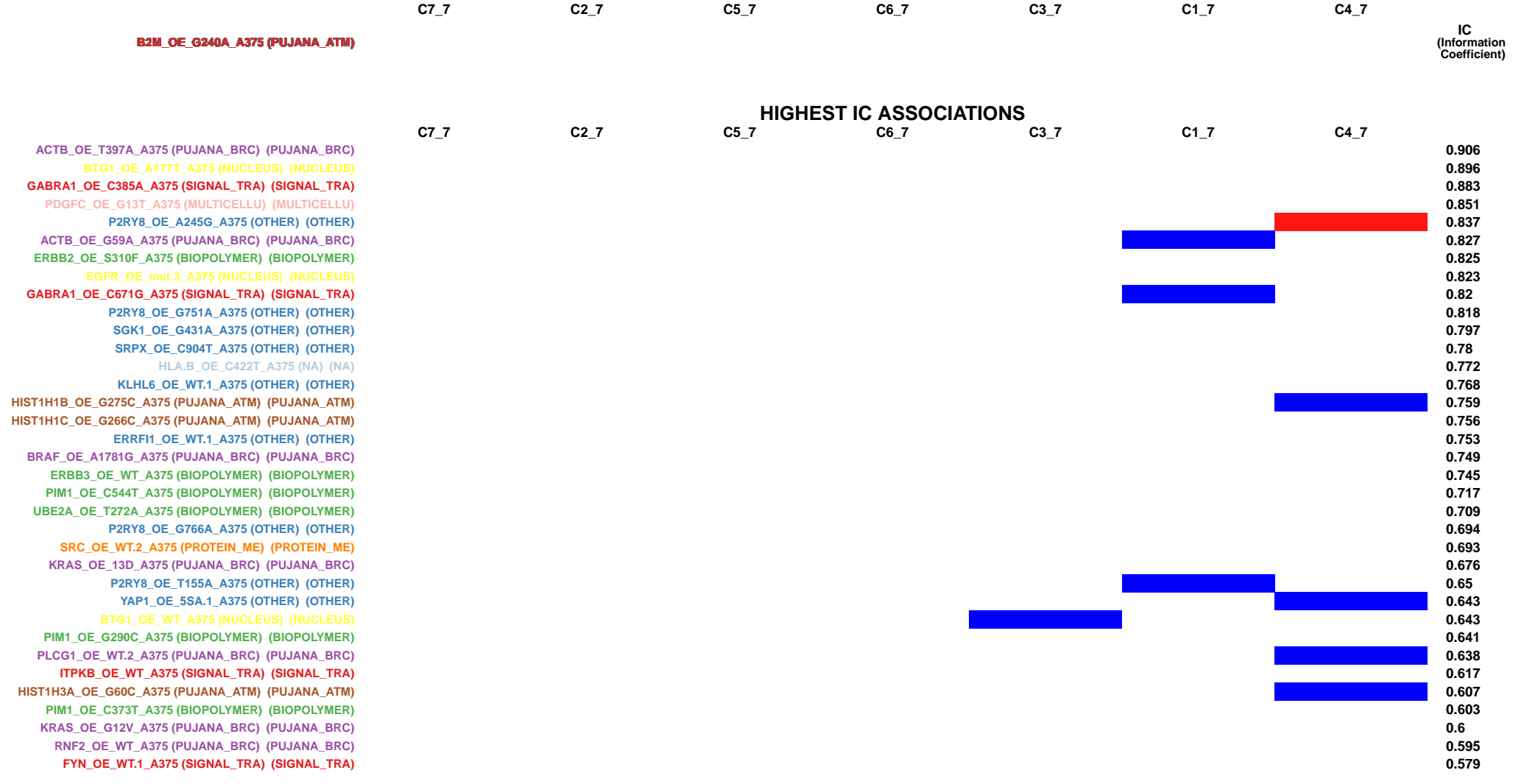




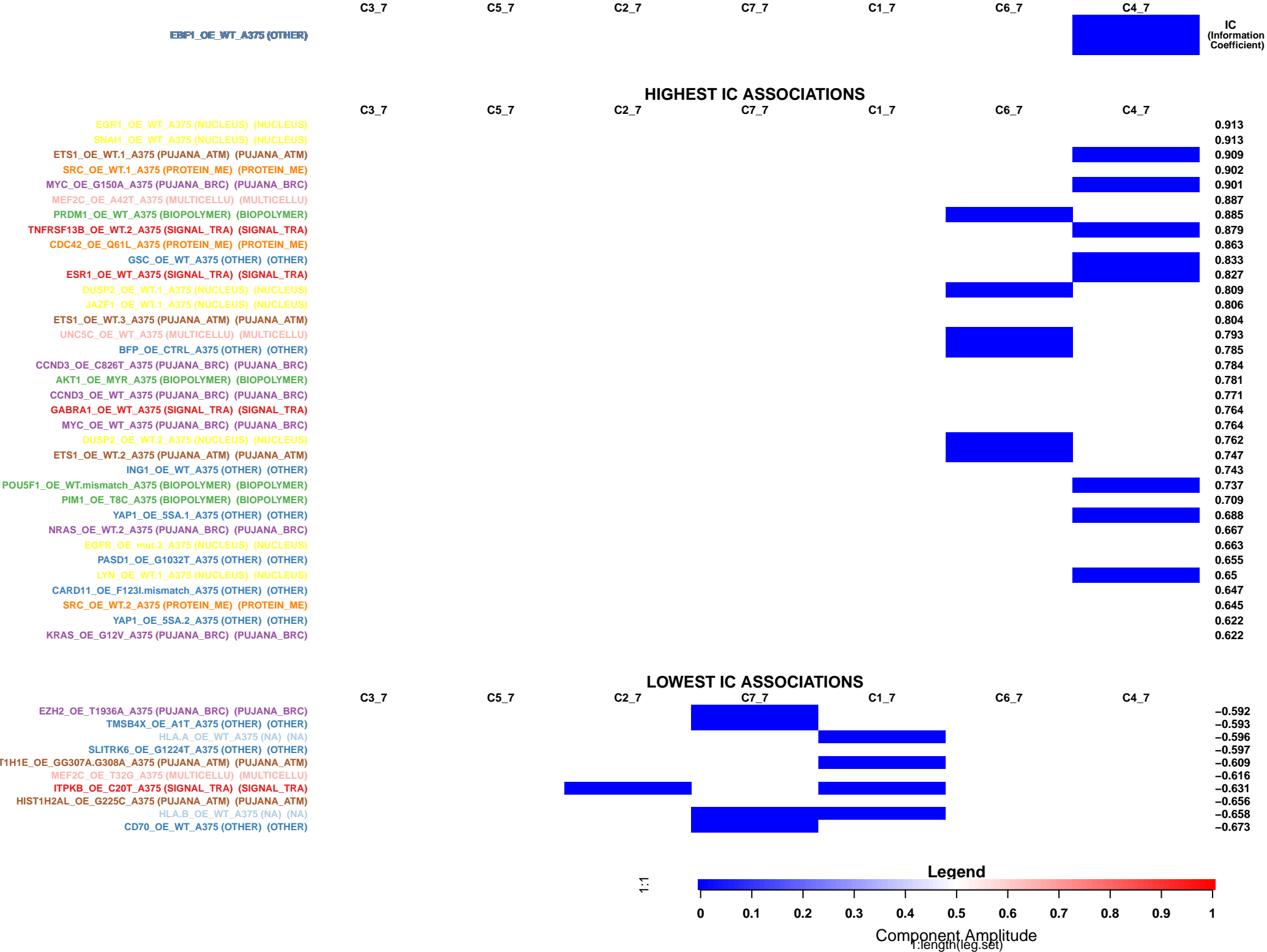
H1FOO\_OE\_WT\_A375 (OTHER)



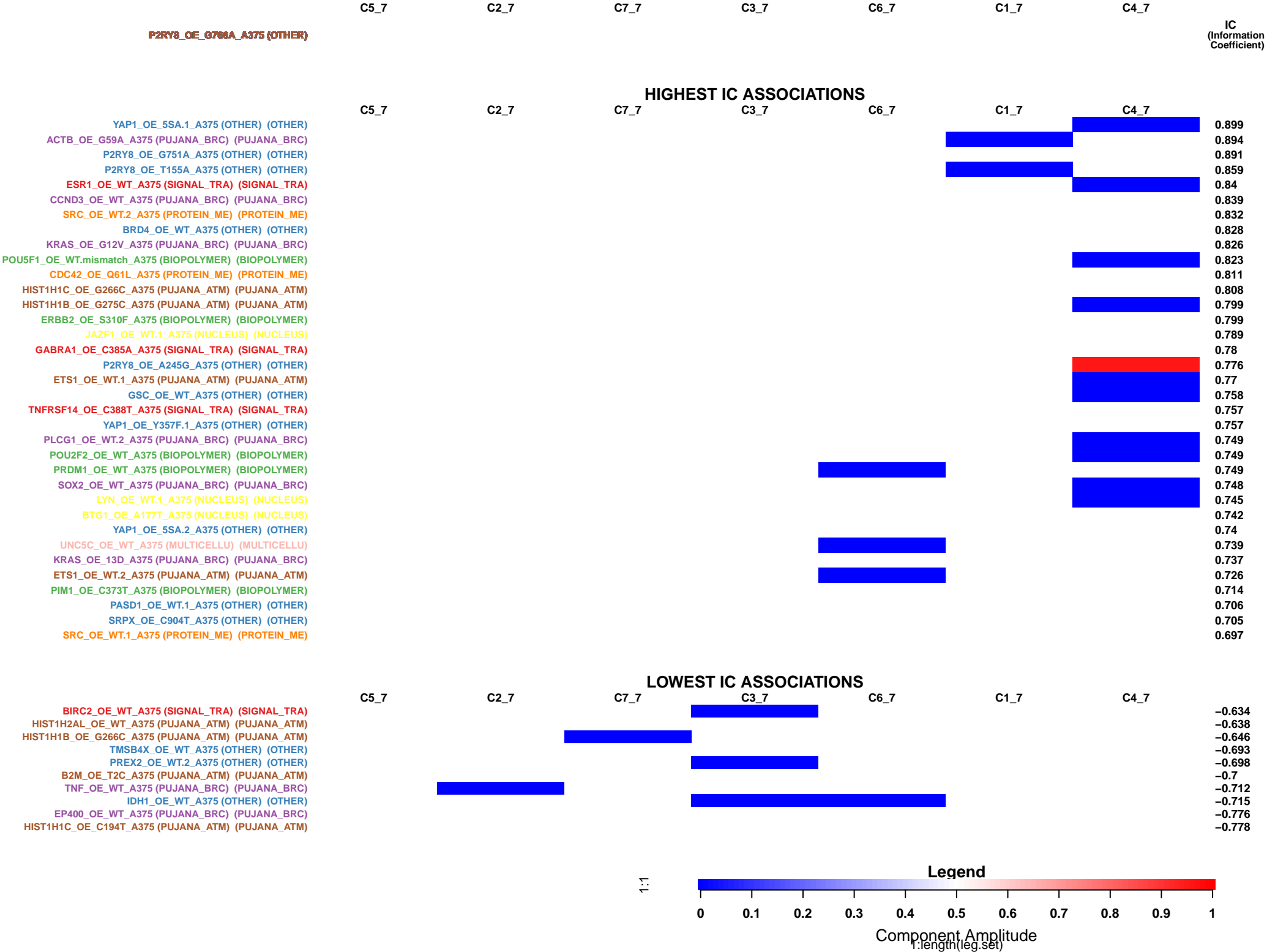
B2M\_OE\_G240A\_A375 (PUJANA\_ATM)



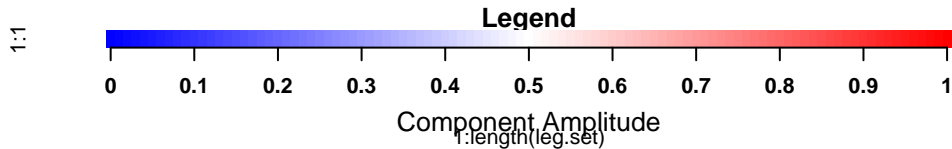
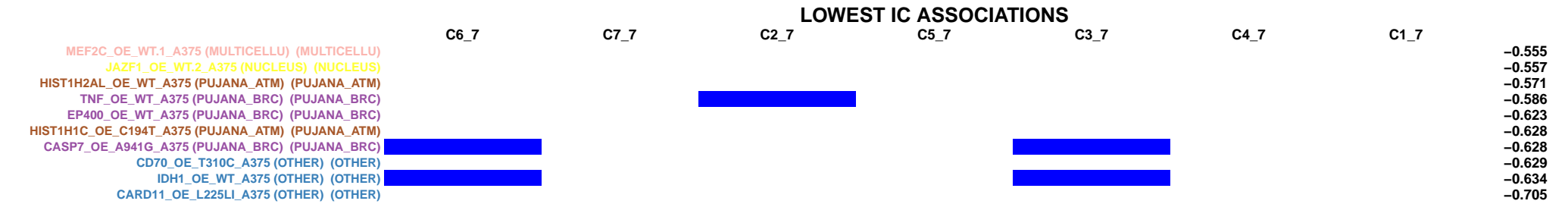
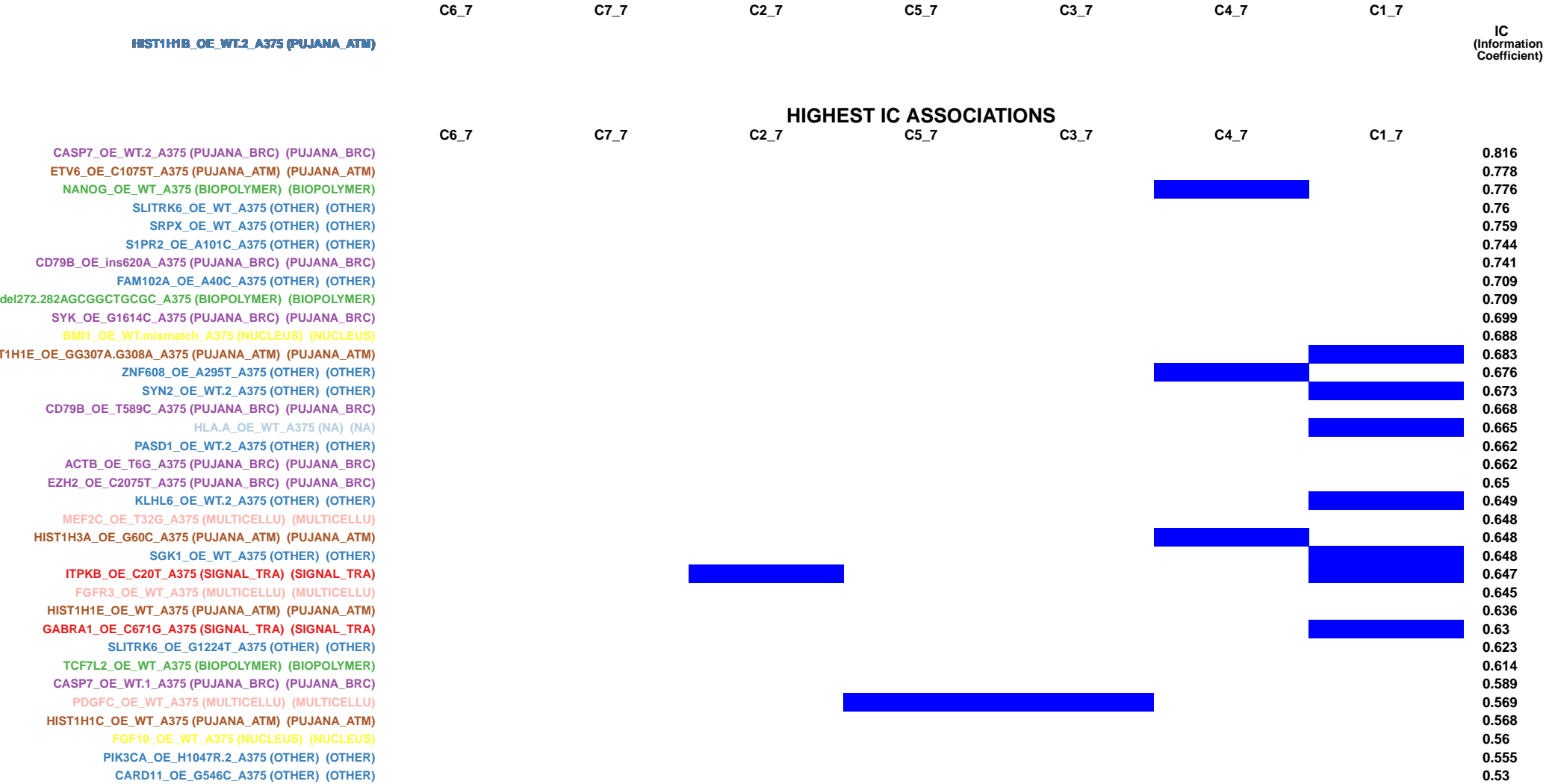
EBF1\_OE\_WT\_A375 (OTHER)



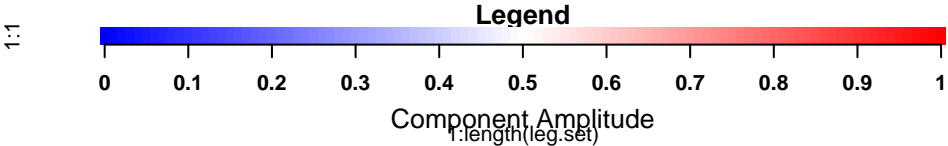
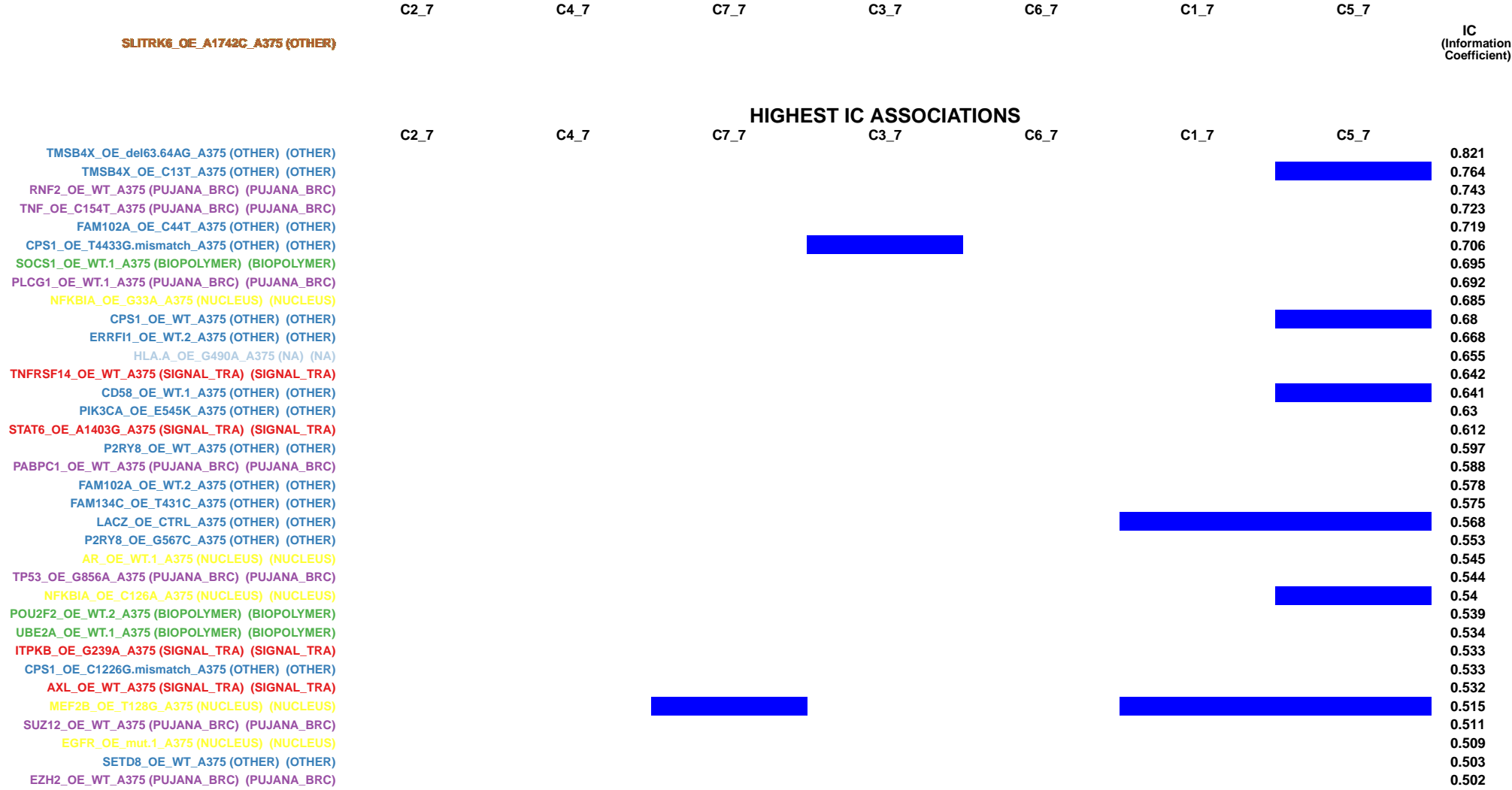
P2RY8\_OE\_G766A\_A375 (OTHER)



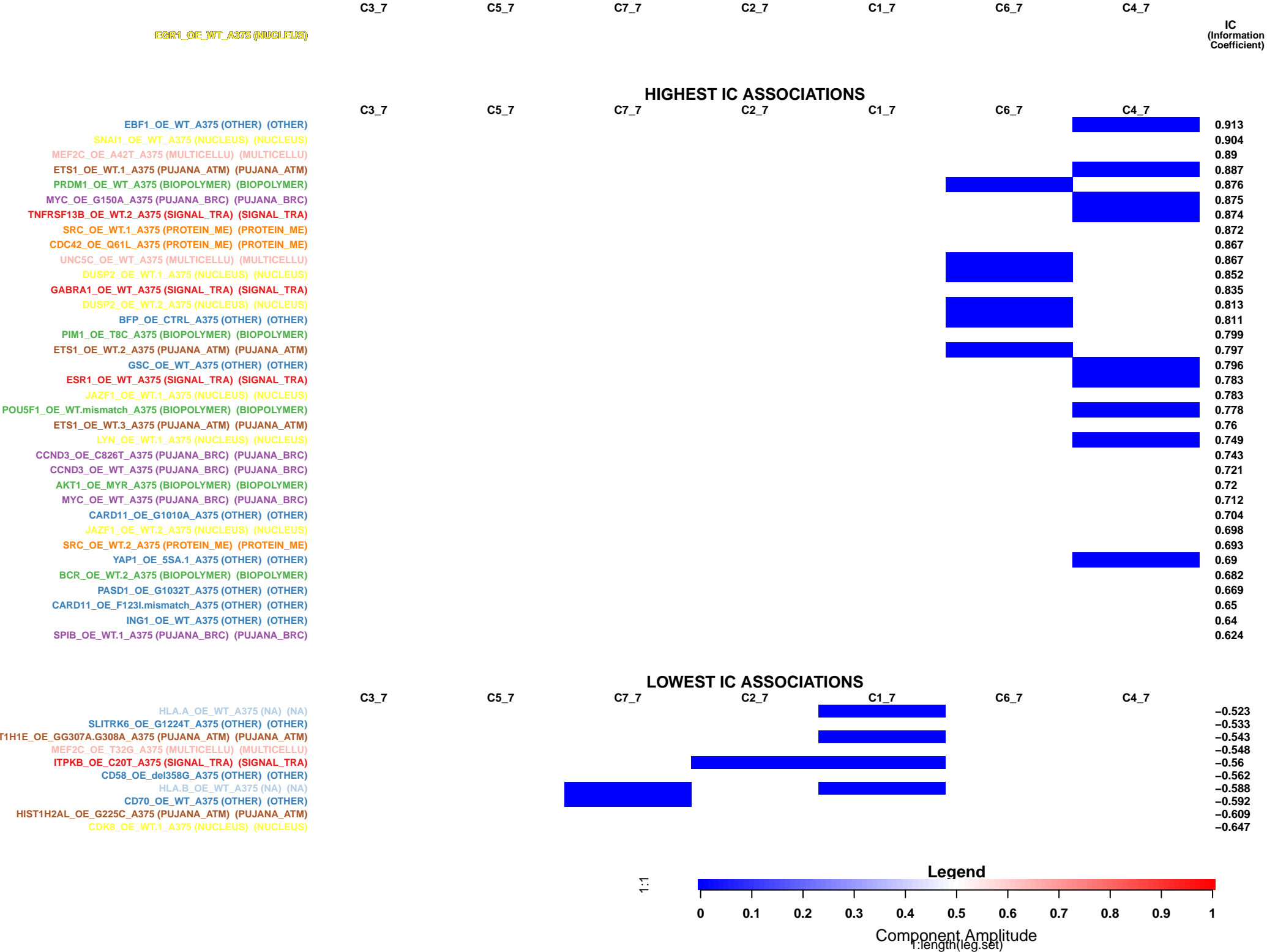
# HIST1H1B\_OE\_WT.2\_A375 (PUJANA\_ATM)



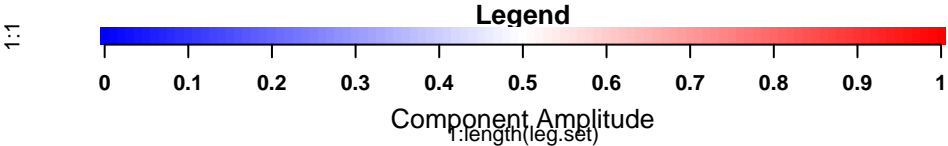
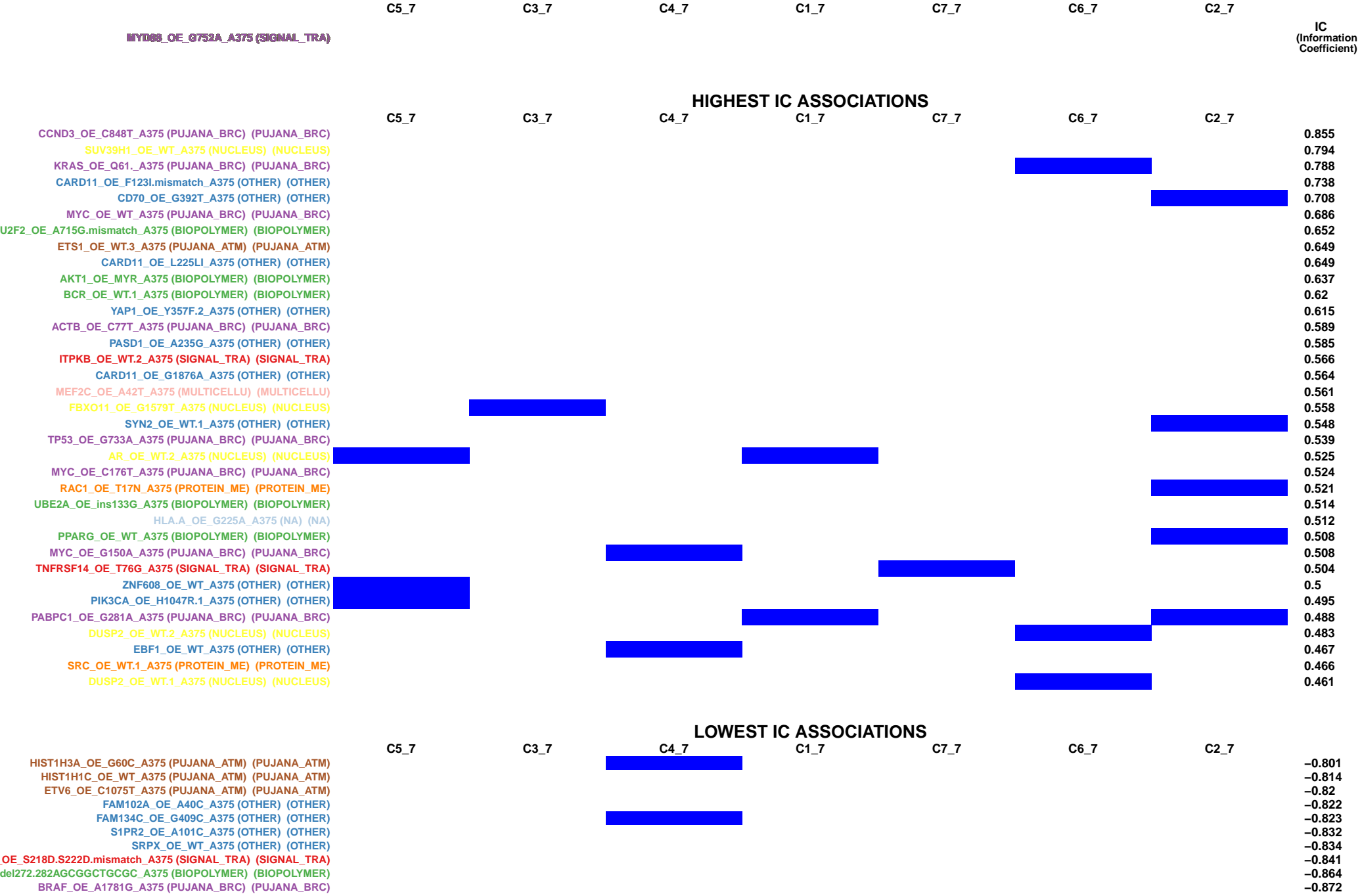
SLITRK6\_OE\_A1742C\_A375 (OTHER)



EGR1\_OE\_WT\_A375 (NUCLEUS)

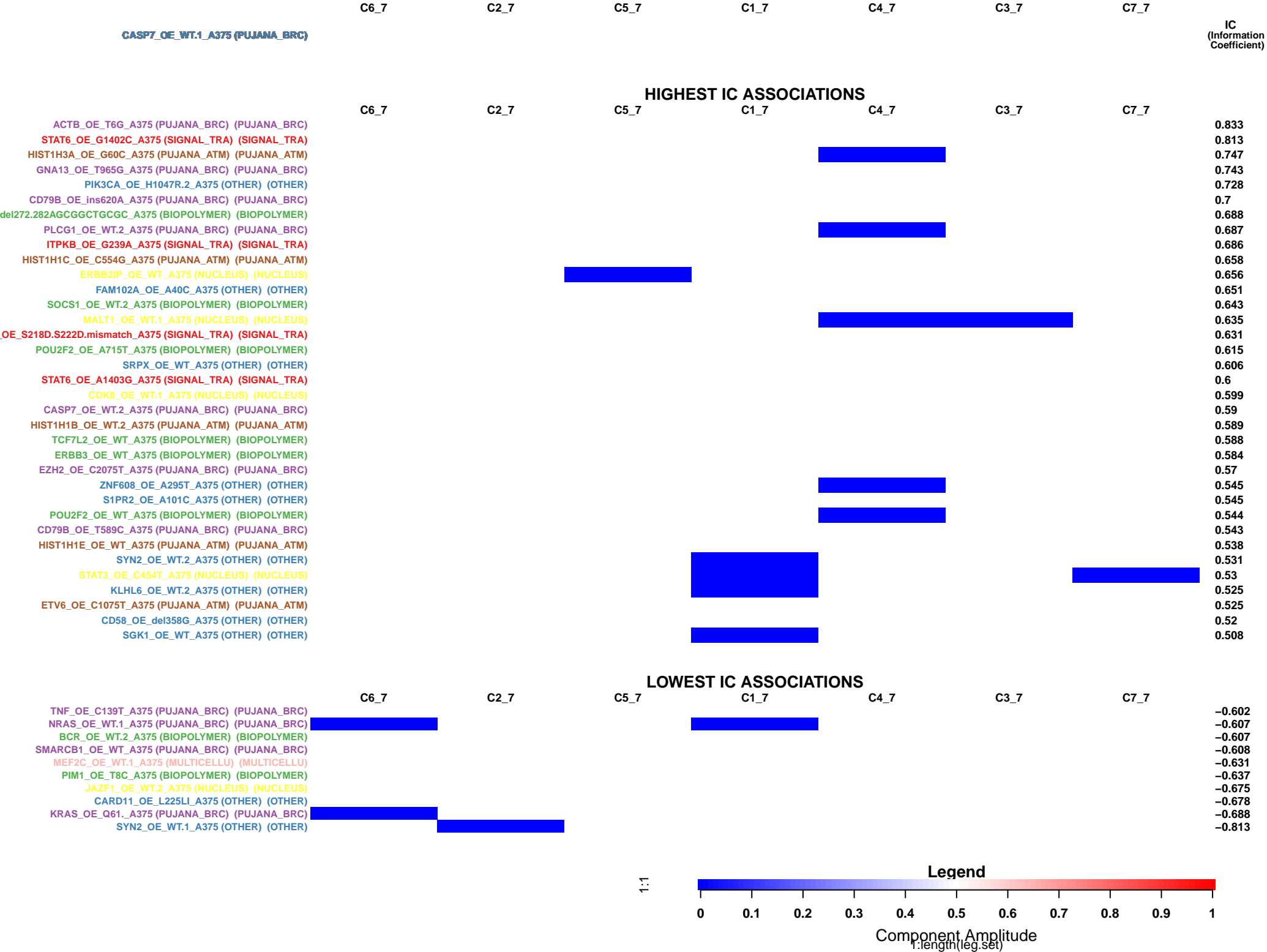


MYD88\_OE\_G752A\_A375 (SIGNAL\_TRA)

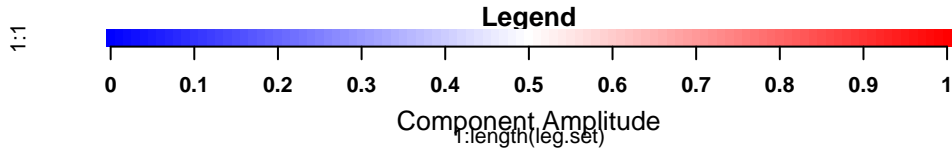
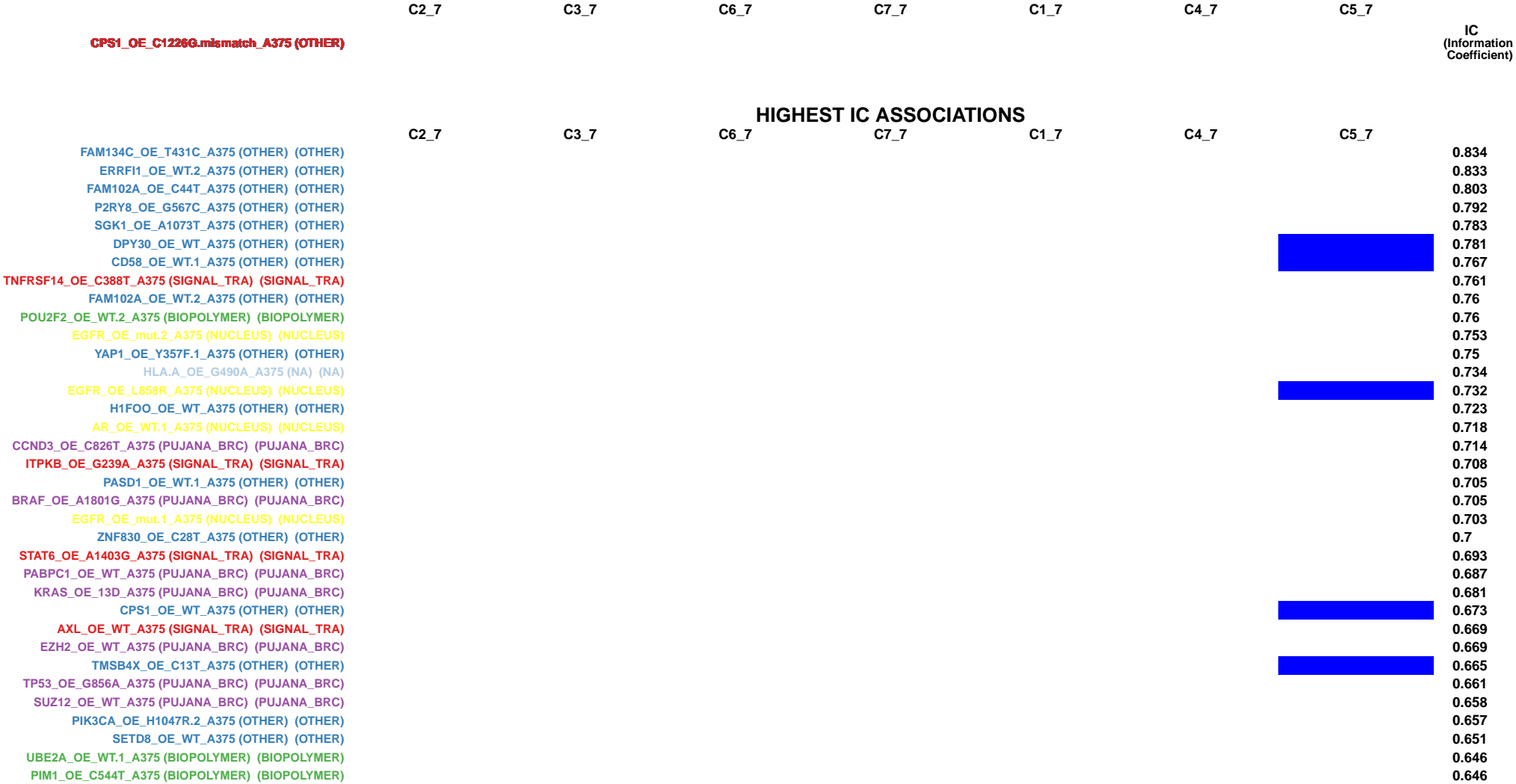




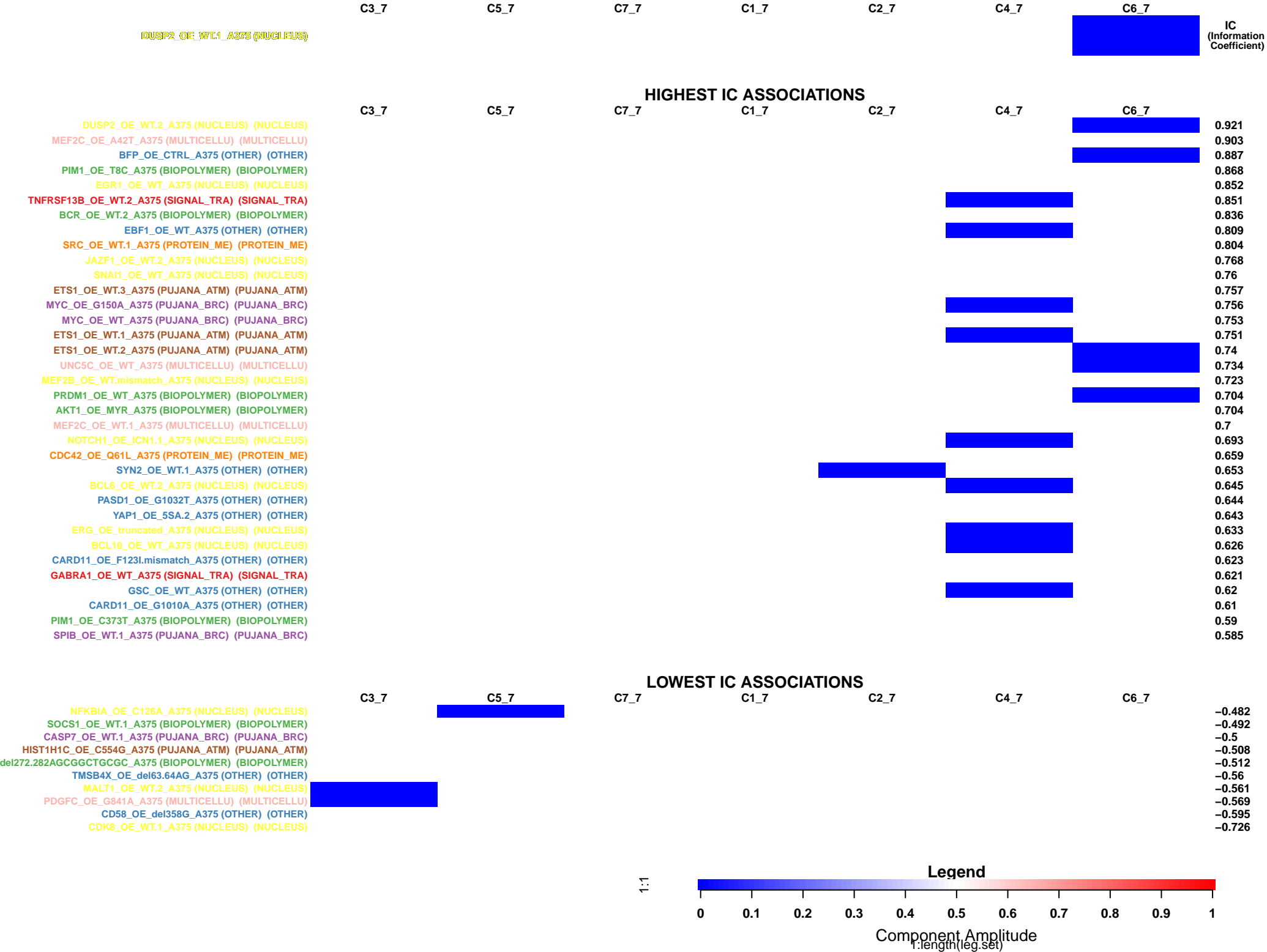
CASP7\_OE\_WT.1\_A375 (PUJANA\_BRC)



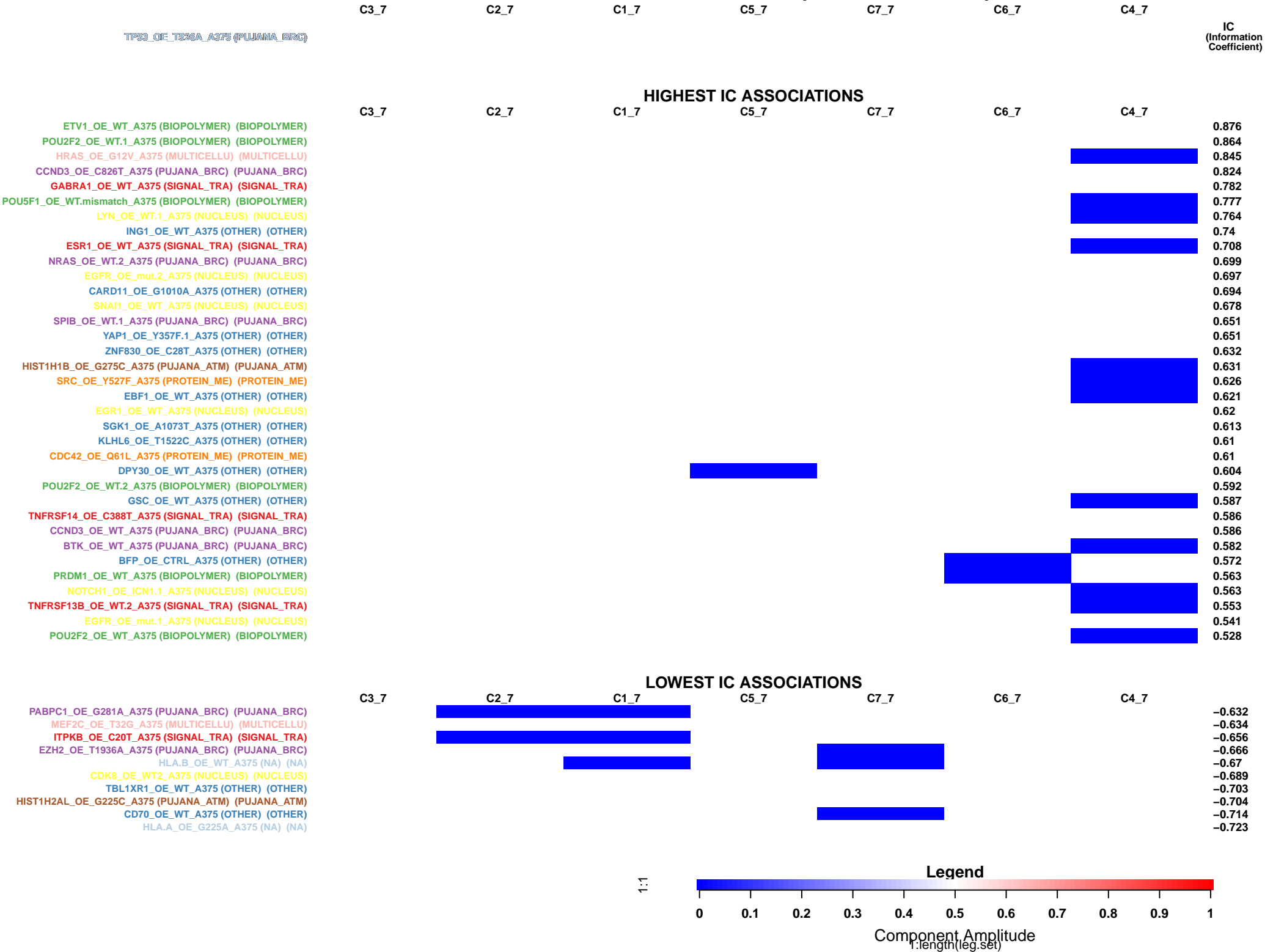
CPS1\_OE\_C1226G.mismatch\_A375 (OTHER)



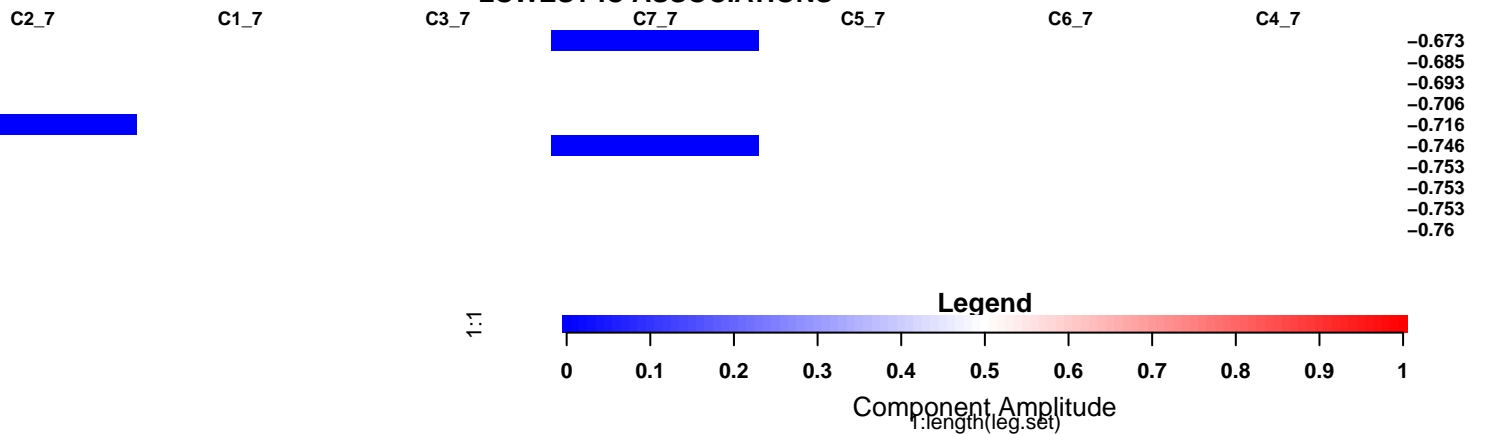
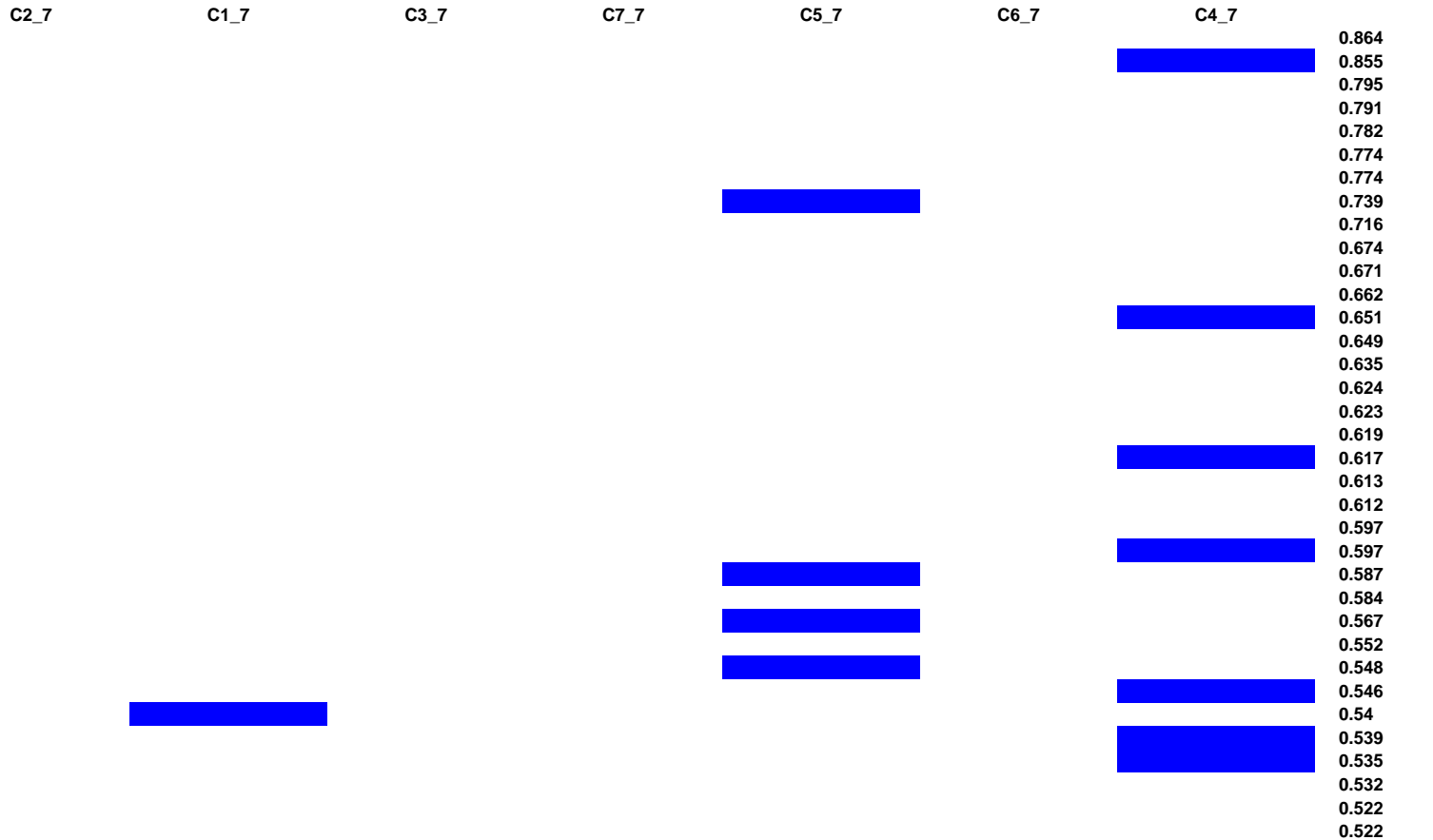
DUSP2\_OE\_WT.1\_A375 (NUCLEUS)



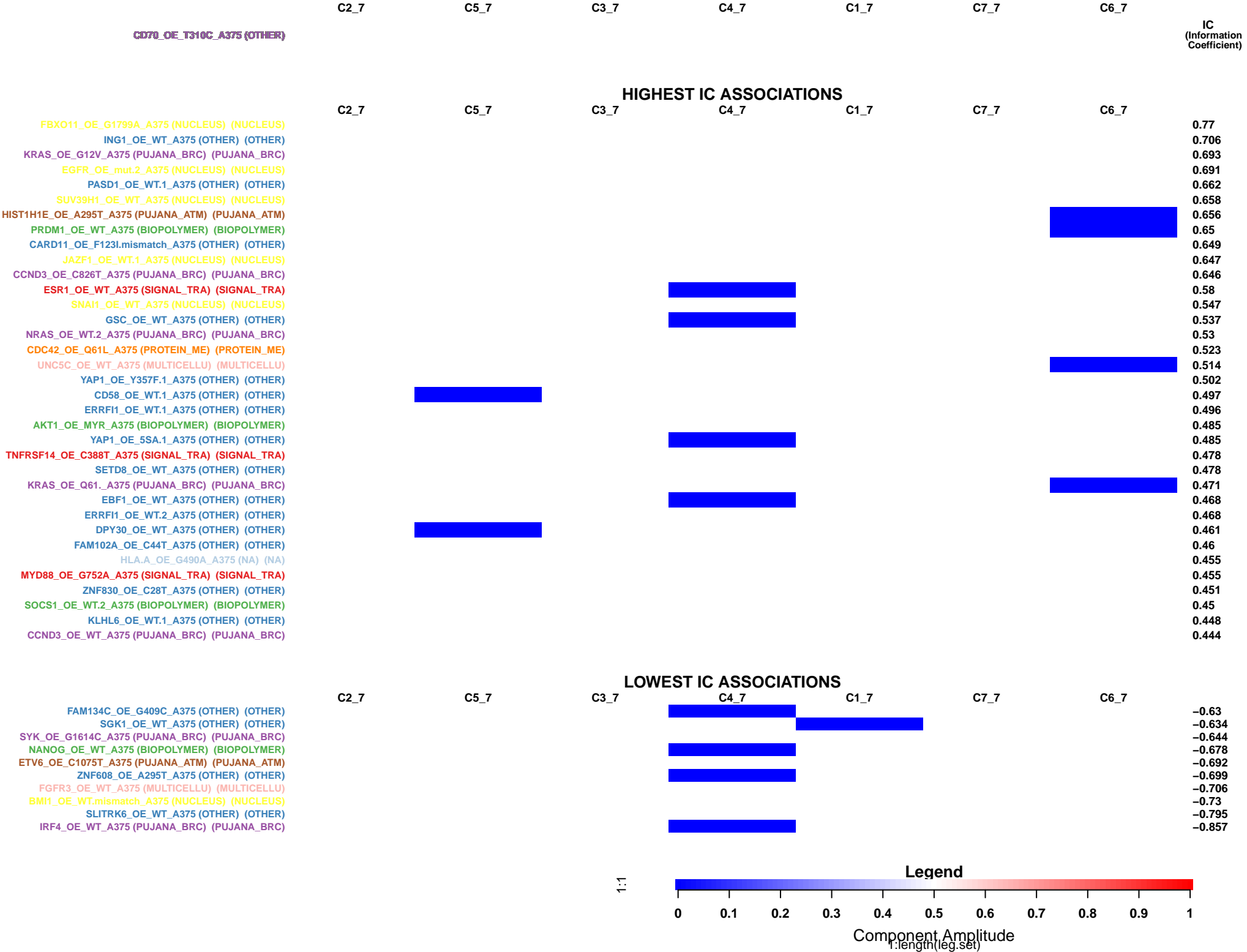
TP53\_OE\_T526A\_A375 (PUJANA\_BRC)



C2_7	C1_7	C3_7	C7_7	C5_7	C6_7	C4_7	IC (Information Coefficient)
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CD70\_OE\_T310C\_A375 (OTHER)



C2\_7                      C3\_7                      C7\_7                      C6\_7                      C1\_7                      C4\_7                      C5\_7

IC  
Information  
Coefficient)

C2\_7                      C3\_7                      C7\_7                      C6\_7                      C1\_7                      C4\_7                      C5\_7



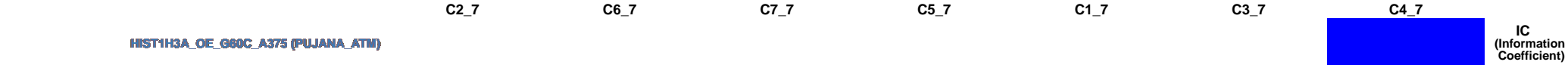
**C2\_7                      C3\_7                      C7\_7                      C6\_7                      C1\_7                      C4\_7                      C5\_7**



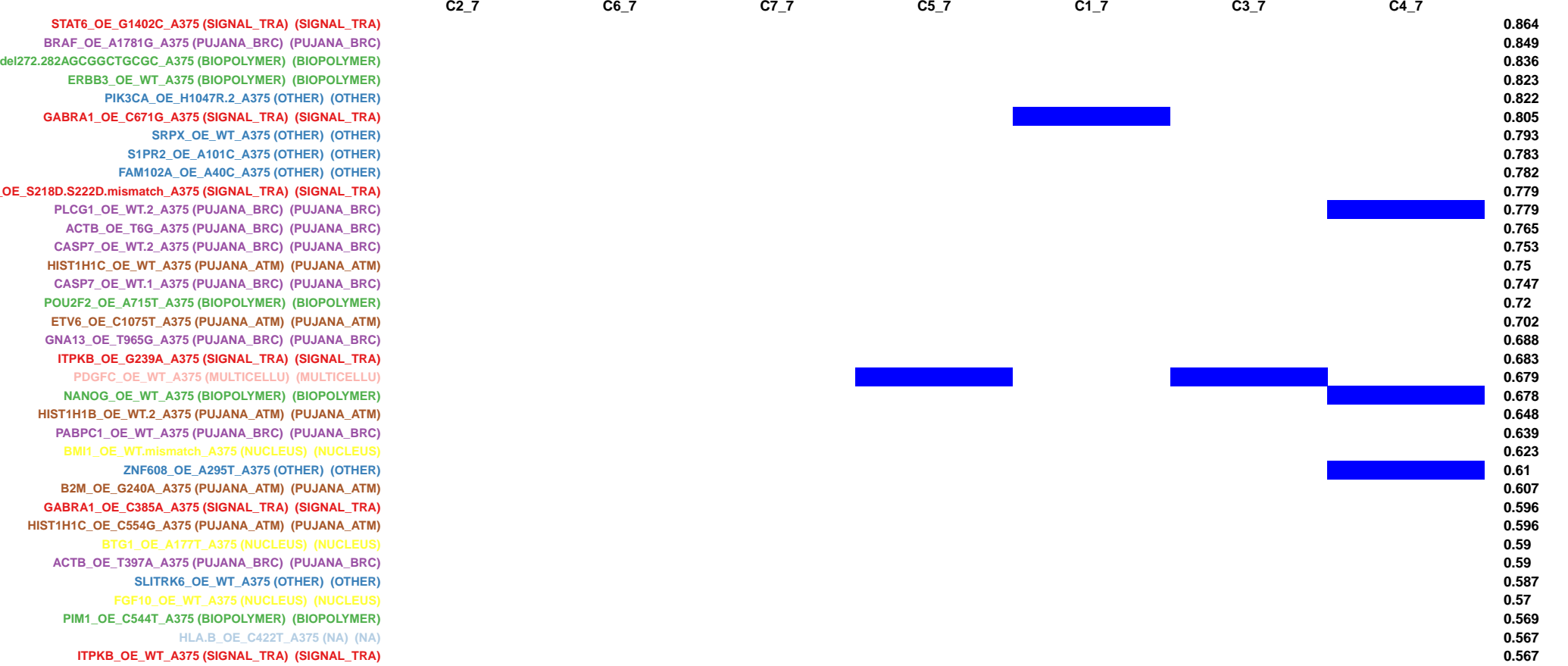
Legend

1: length (lég. söt)

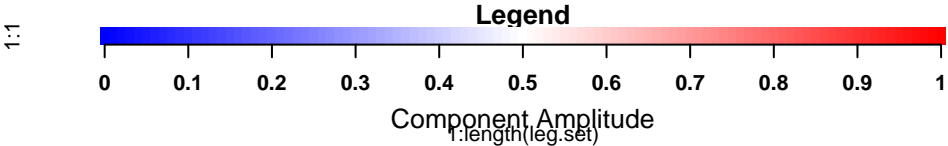
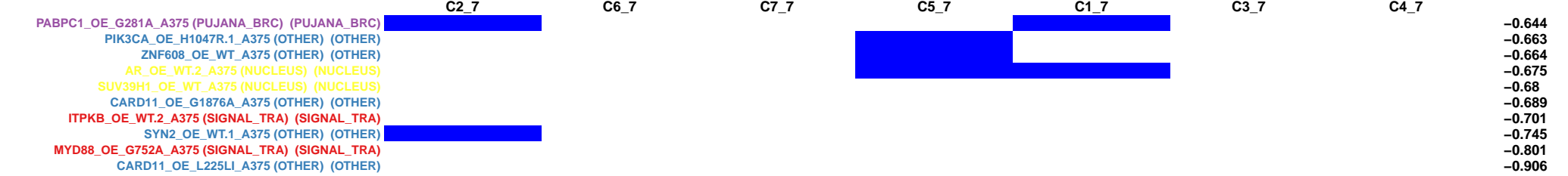
HIST1H3A\_OE\_G60C\_A375 (PUJANA\_ATM)



HIGHEST IC ASSOCIATIONS

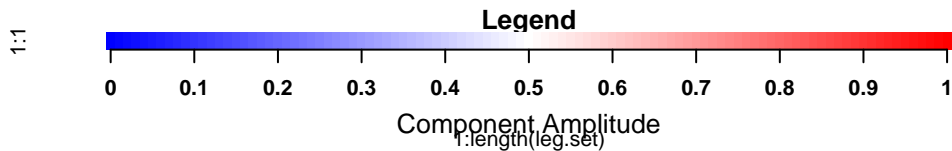
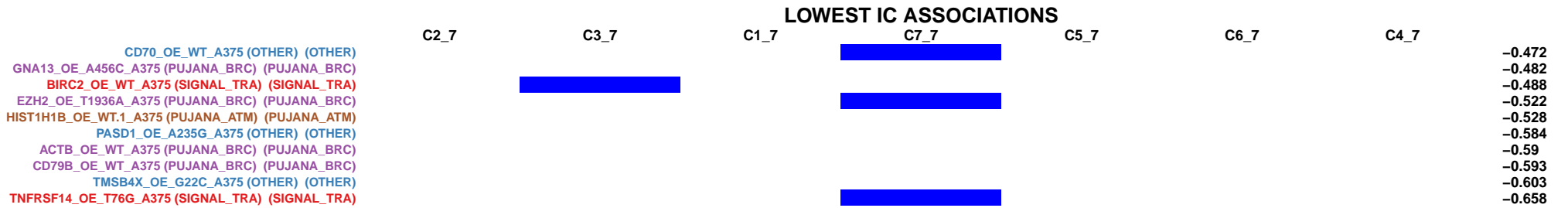
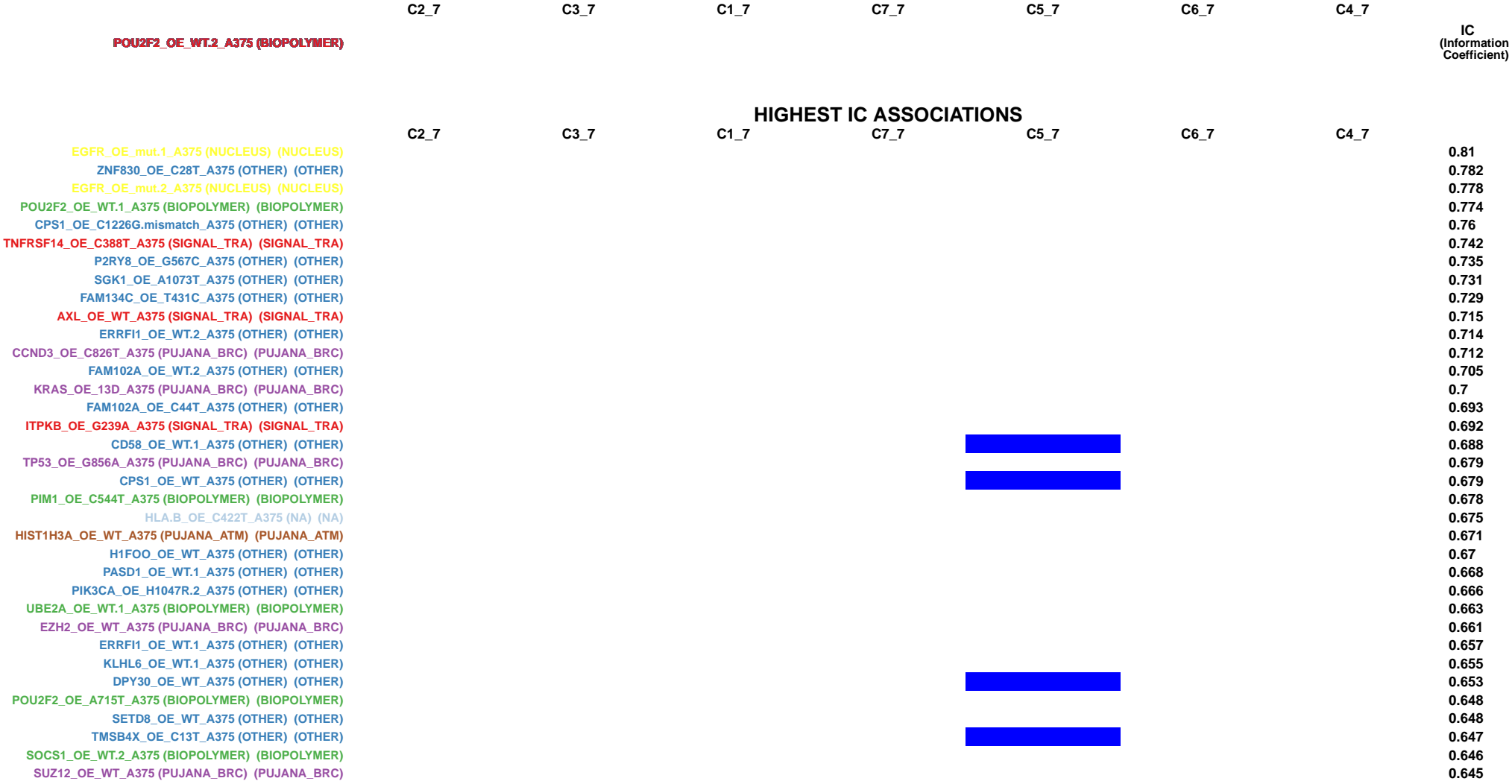


LOWEST IC ASSOCIATIONS

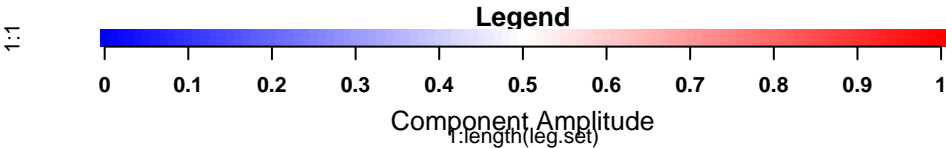
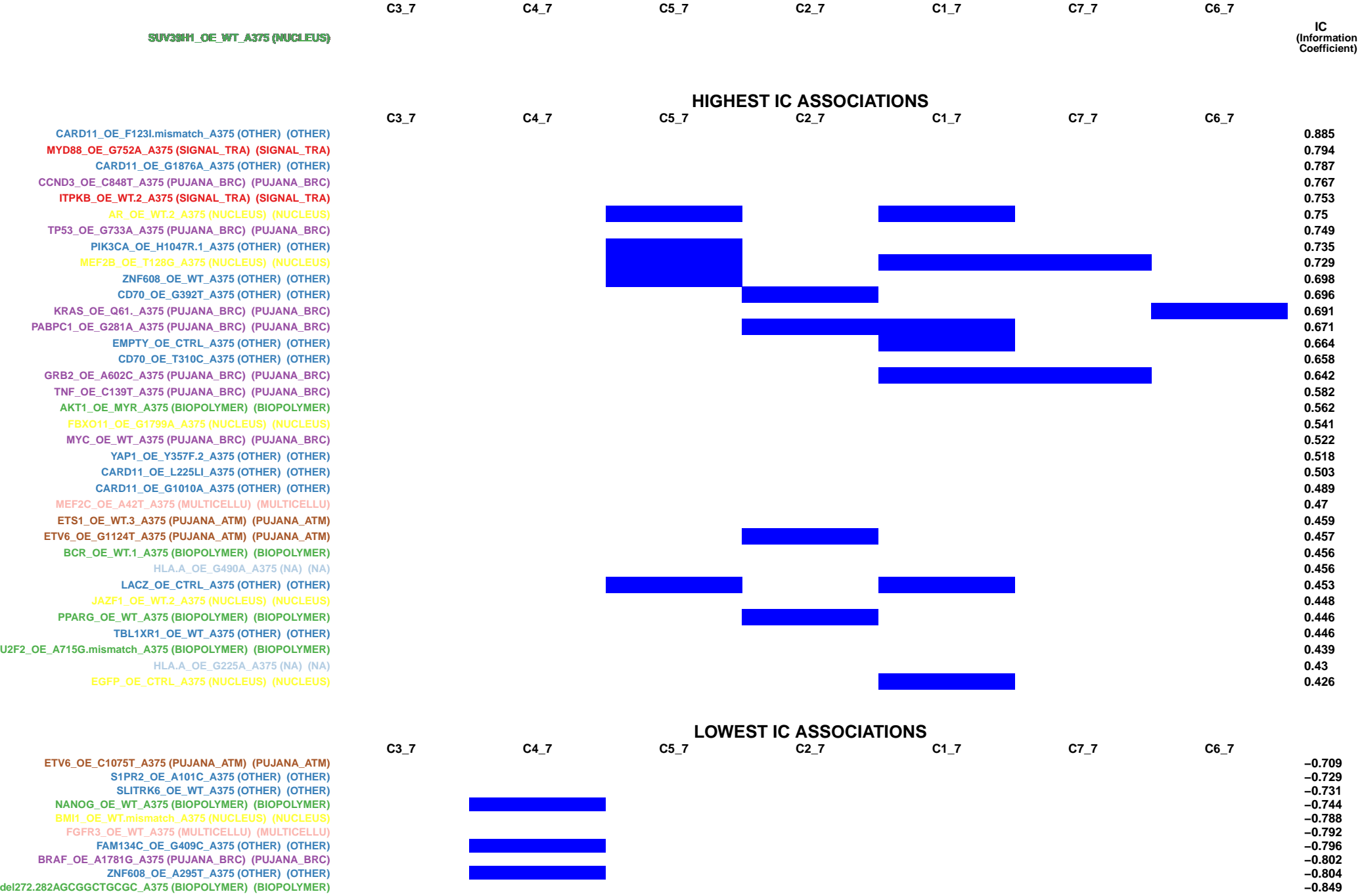




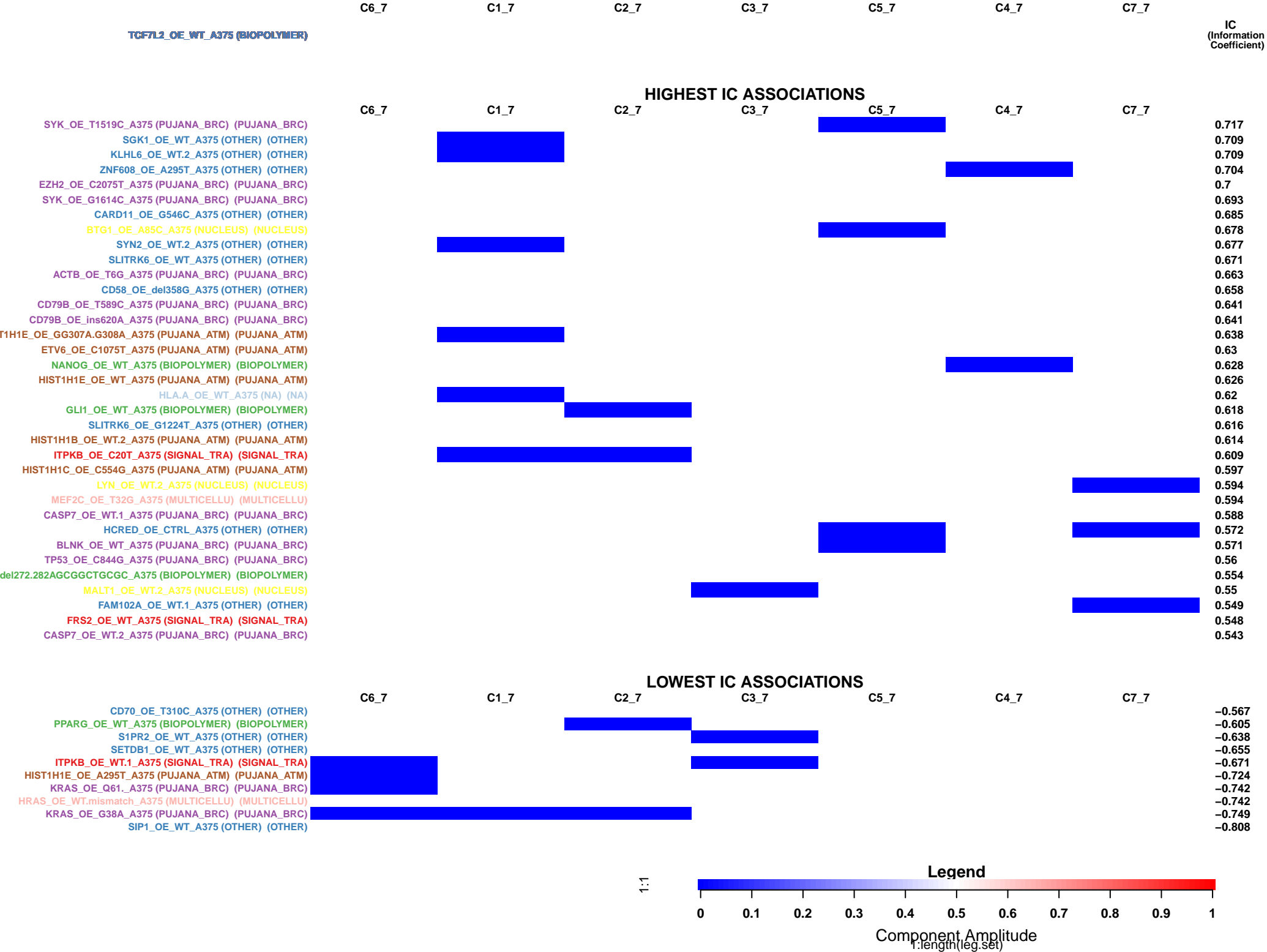
POU2F2\_OE\_WT.2\_A375 (BIOPOLYMER)



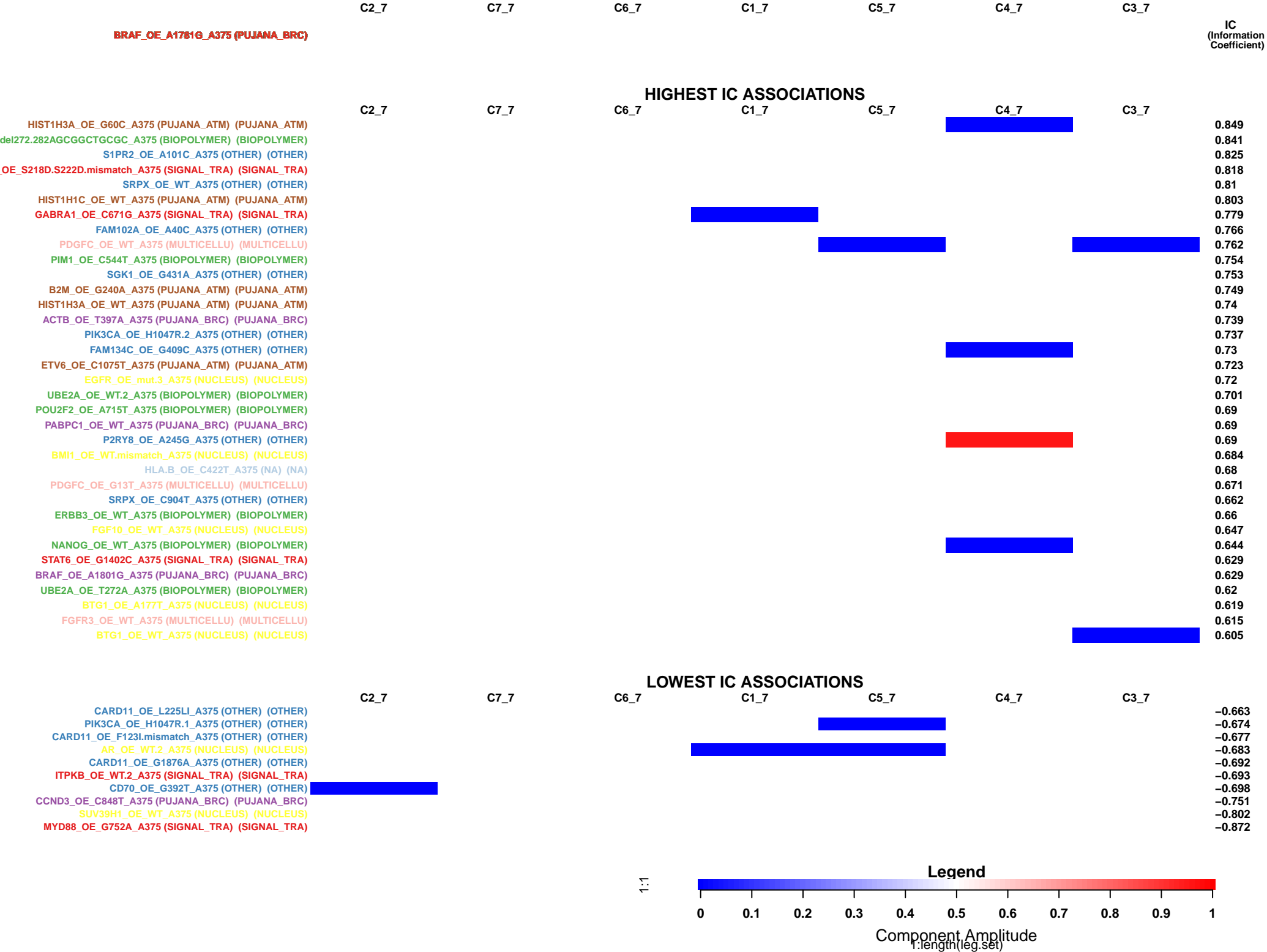
SUV39H1\_OE\_WT\_A375 (NUCLEUS)



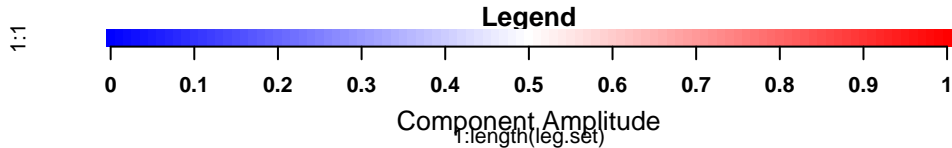
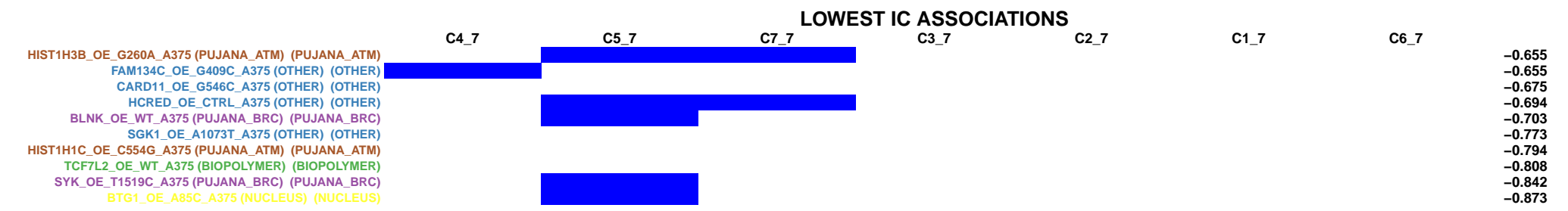
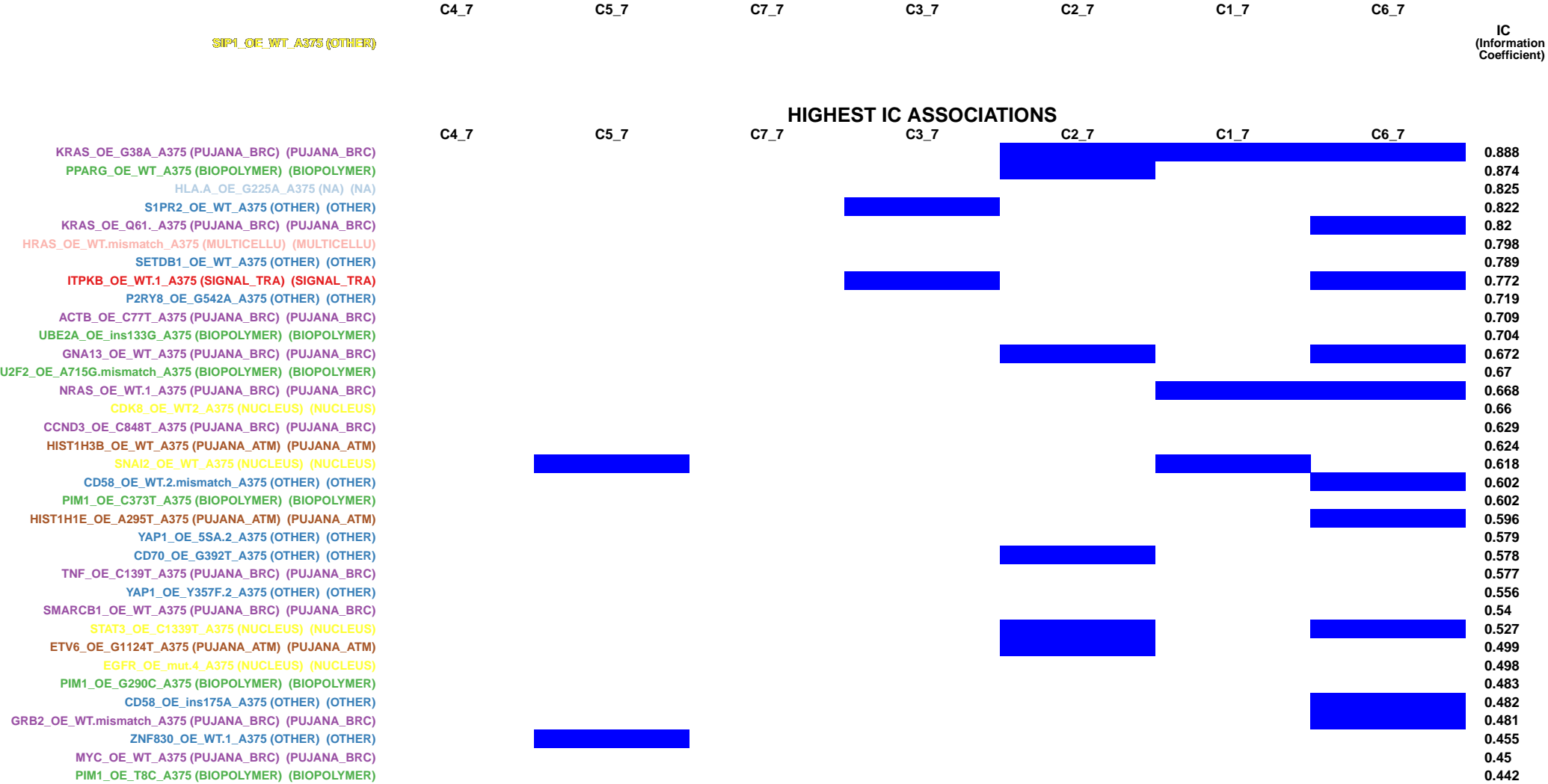
TCF7L2\_OE\_WT\_A375 (BIOPOLYMER)



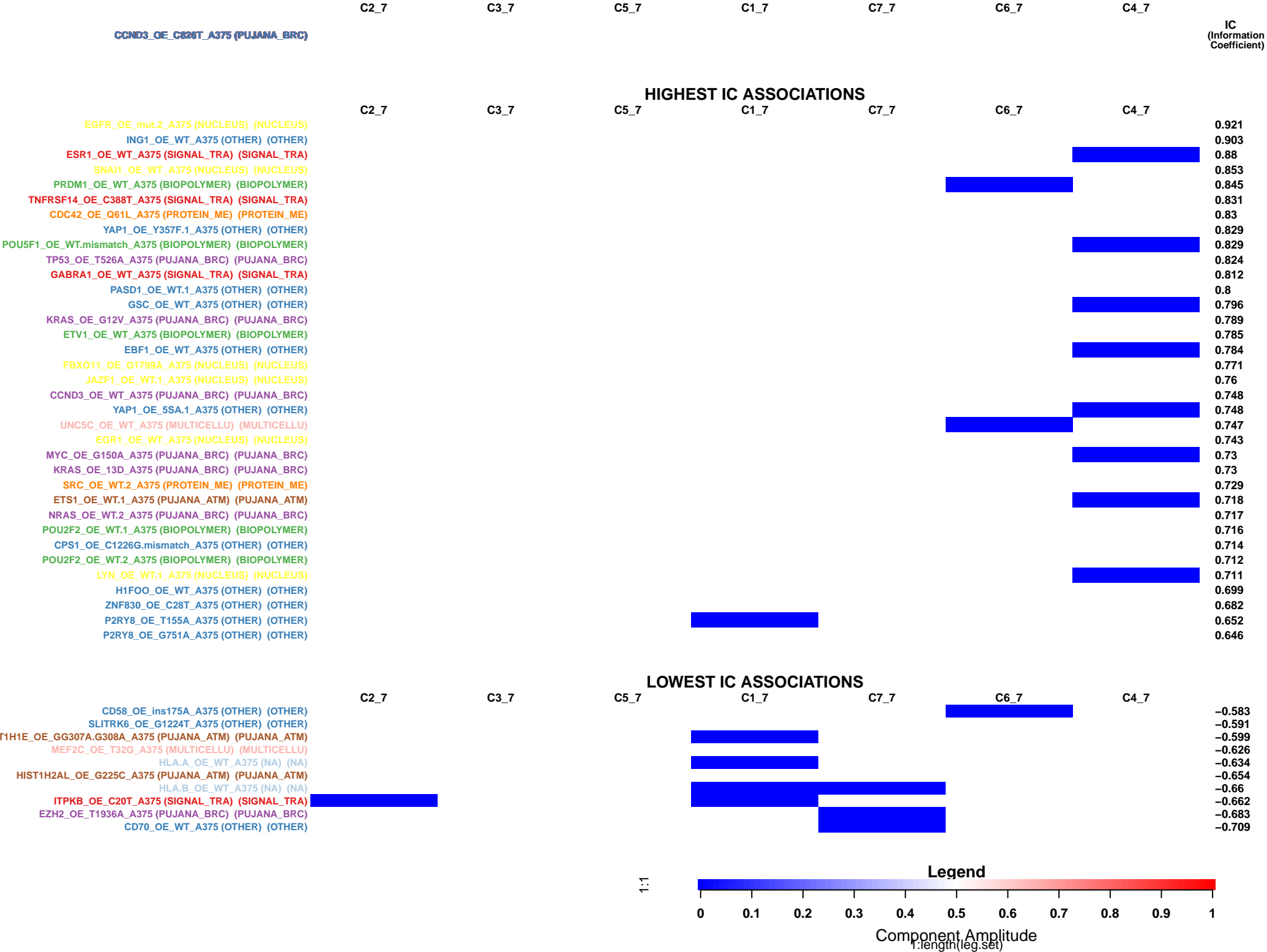
BRAF\_OE\_A1781G\_A375 (PUJANA\_BRC)



SIP1\_OE\_WT\_A375 (OTHER)



CCND3\_OE\_C826T\_A375 (PUJANA\_BRC)



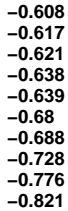
C5_7	C7_7	C3_7	C2_7	C4_7	C6_7	C1_7
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0.901

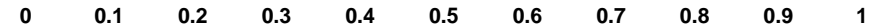
C5\_7                      C7\_7                      C3\_7                      C2\_7                      C4\_7                      C6\_7                      C1\_7



C5_7	C7_7	C3_7	C2_7	C4_7	C6_7	C1_7
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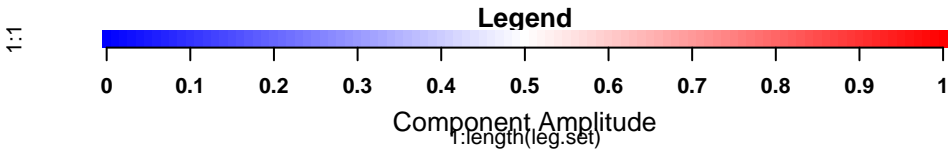
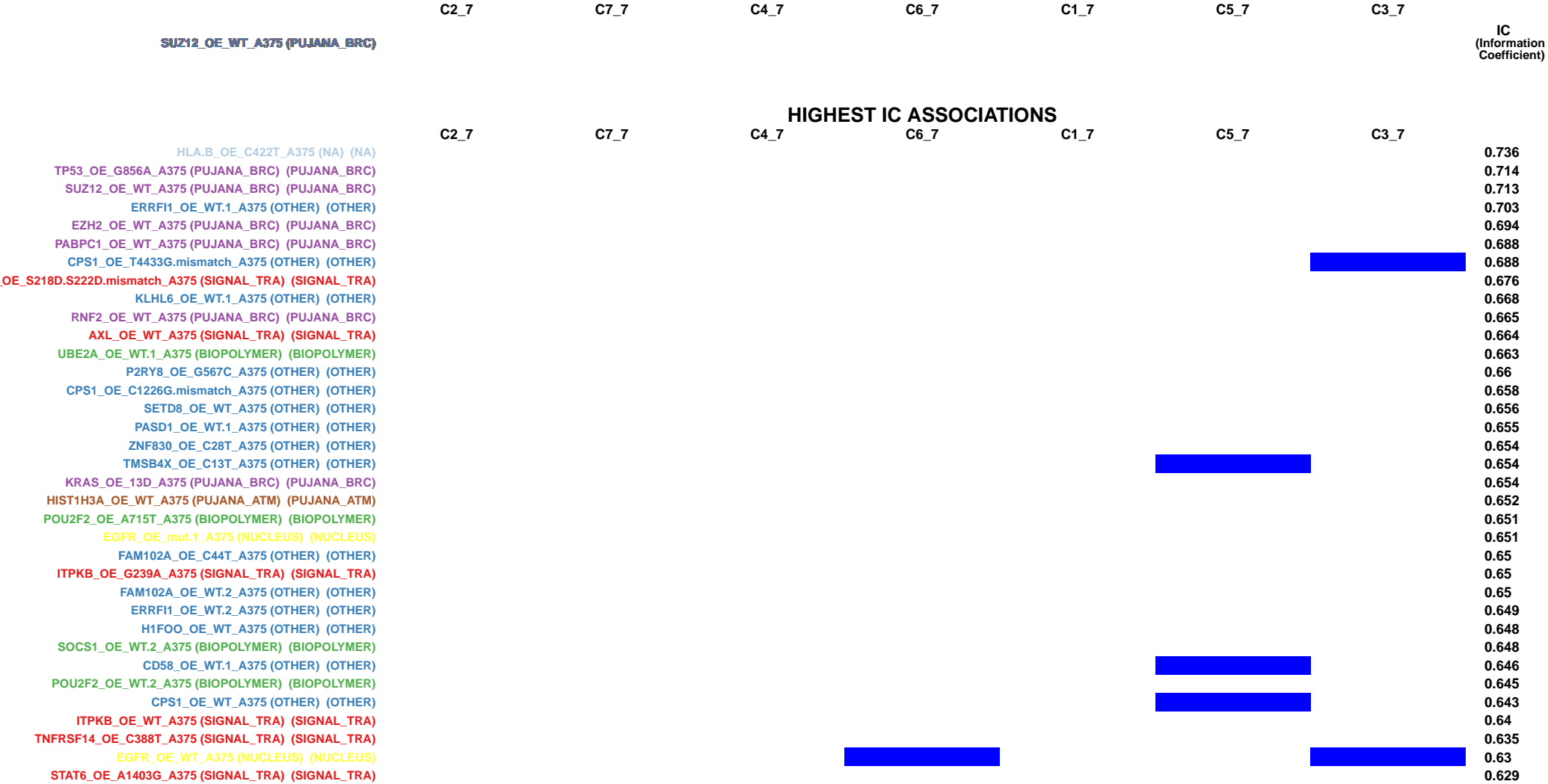


### Legend



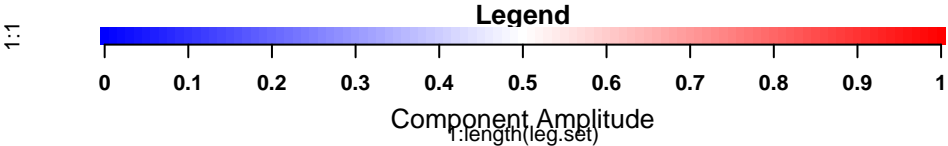
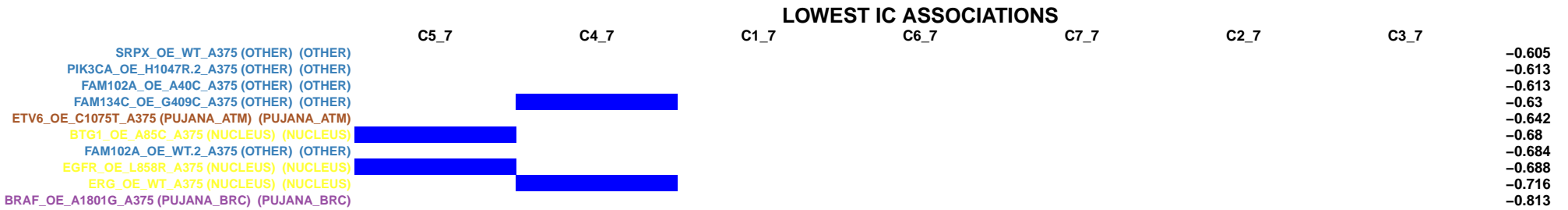
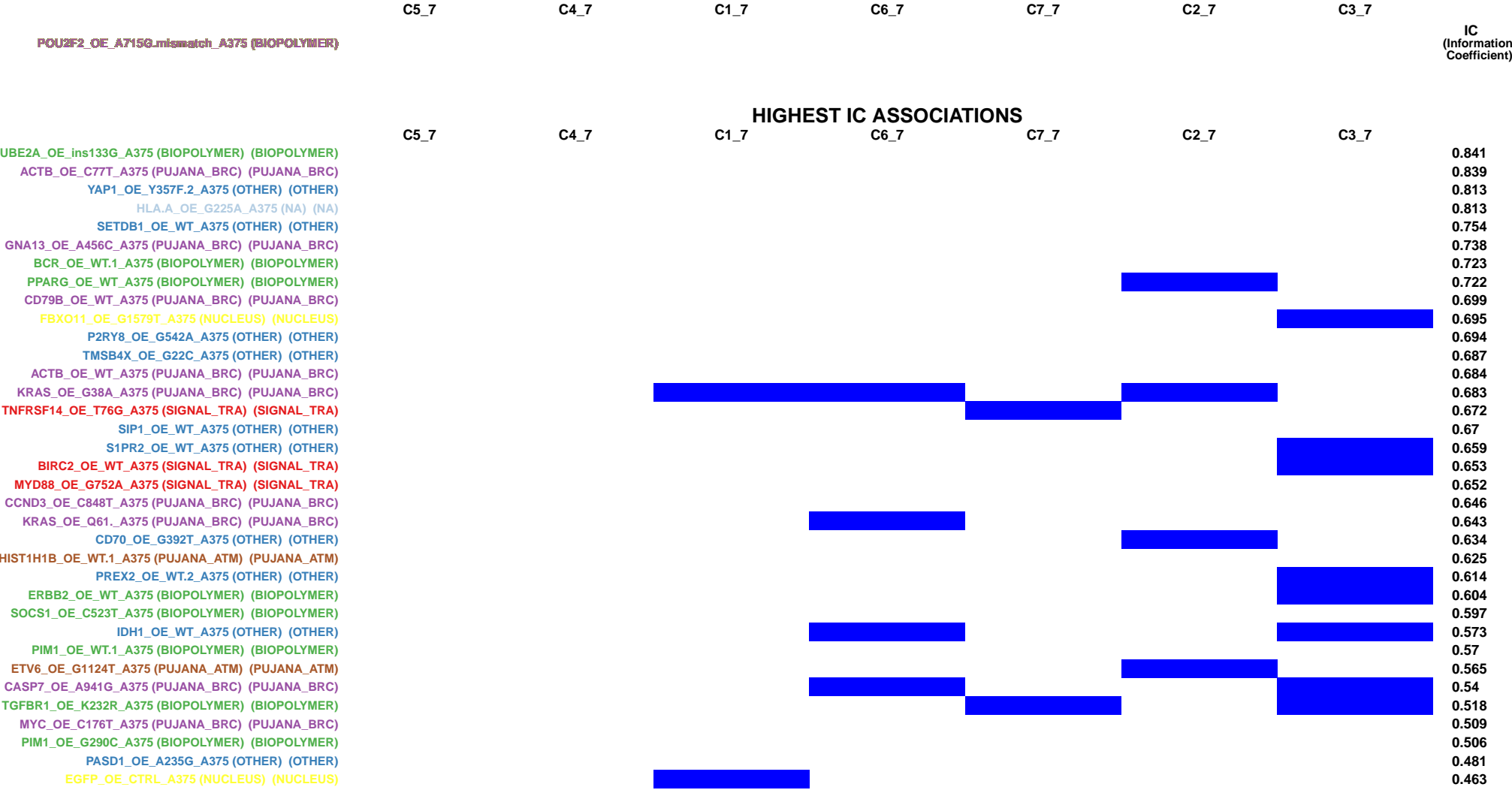
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1:length(leg.set)
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SUZ12\_OE\_WT\_A375 (PUJANA\_BRC)





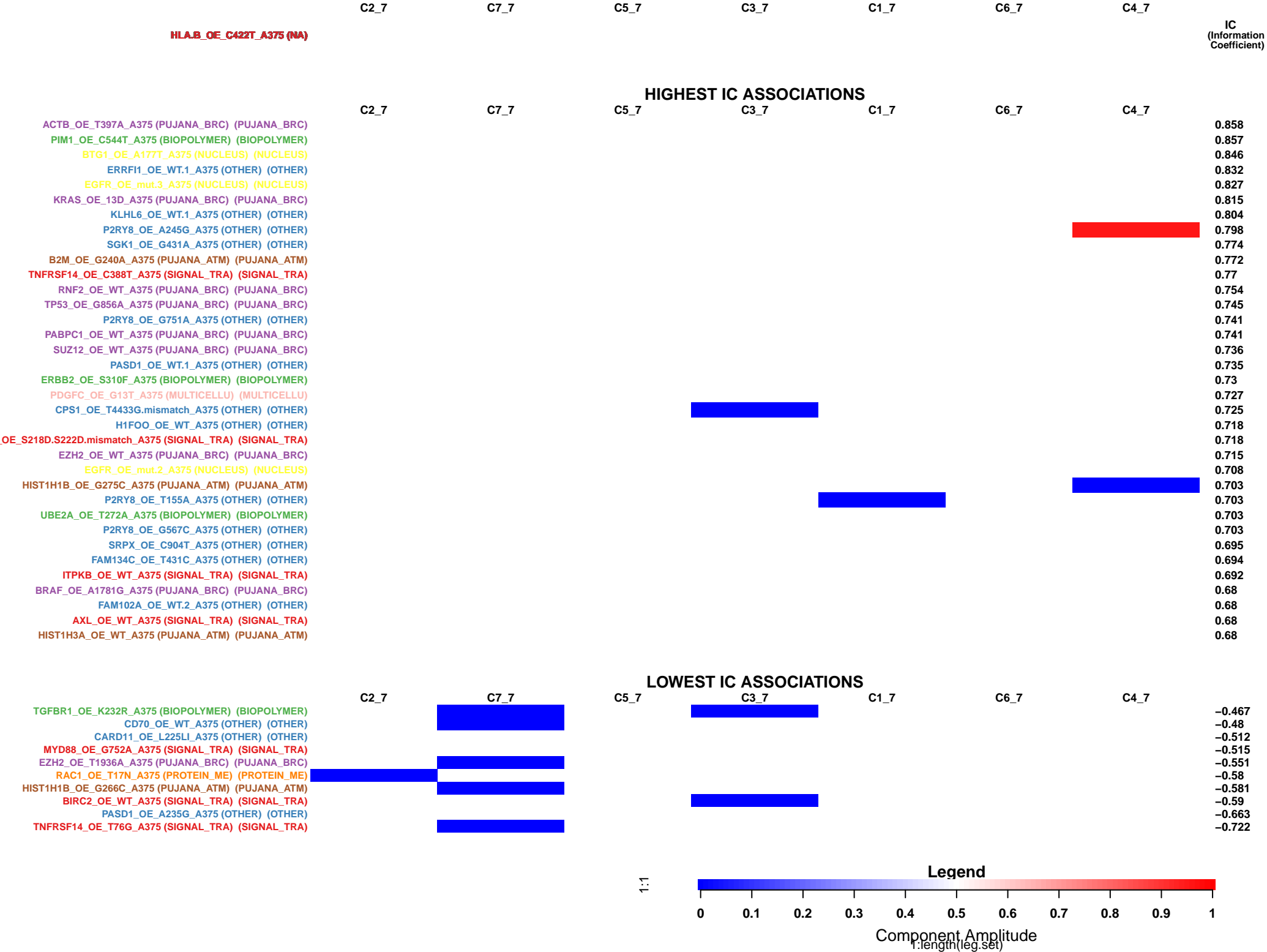
POU2F2\_OE\_A715G.mismatch\_A375 (BIOPOLYMER)



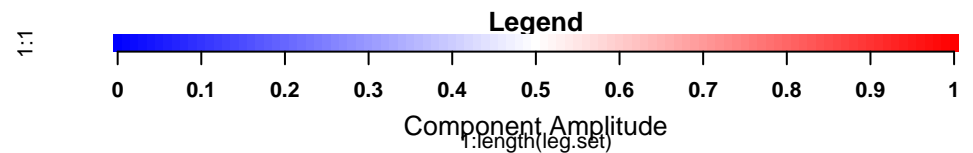
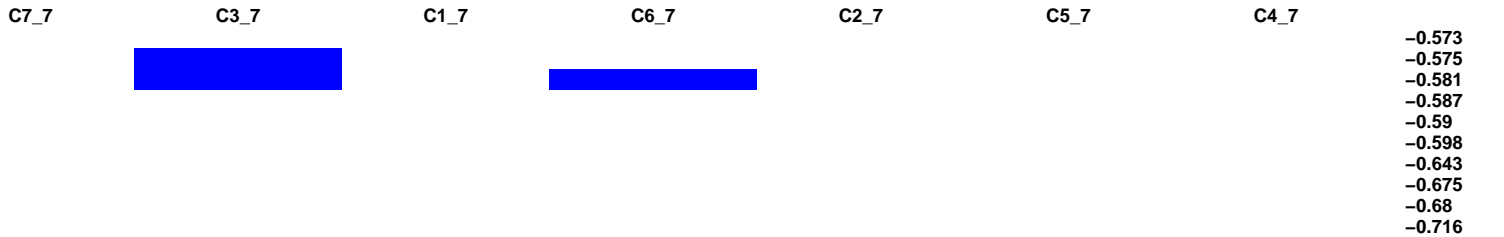
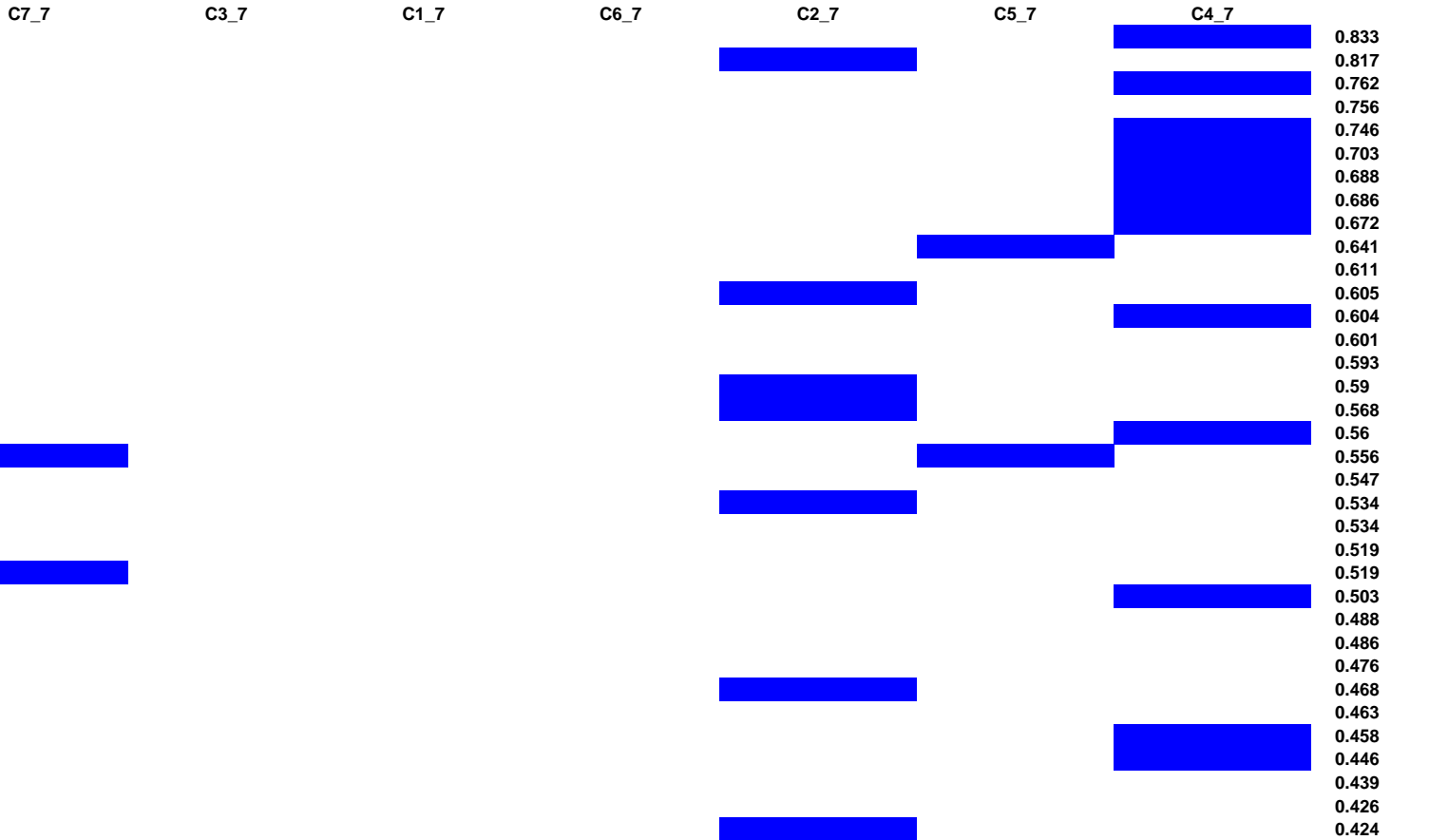
## 1:1



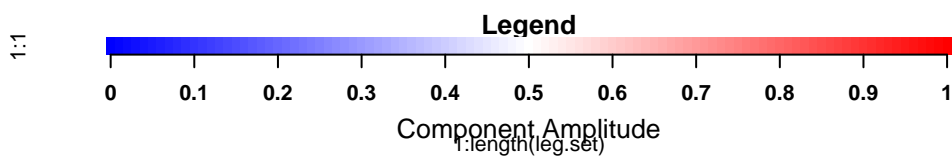
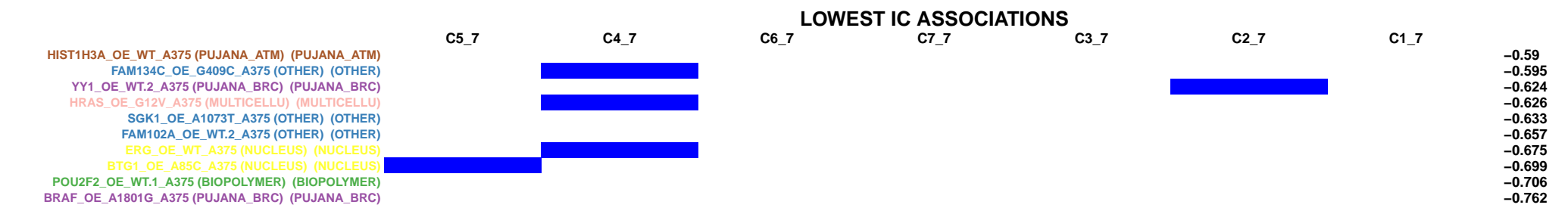
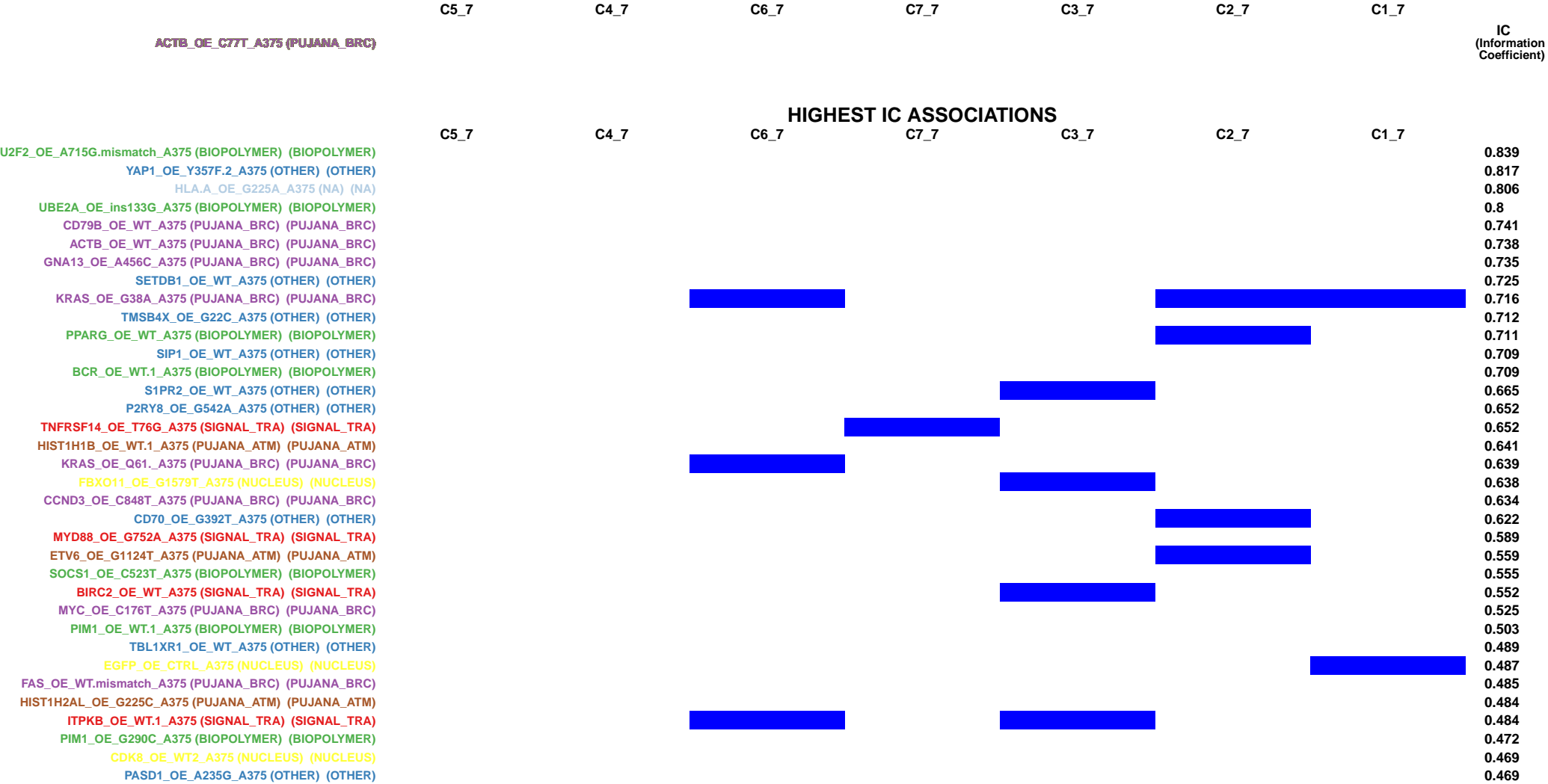
HLA.B\_OE\_C422T\_A375 (NA)



C7_7	C3_7	C1_7	C6_7	C2_7	C5_7	C4_7	IC (Information Coefficient)

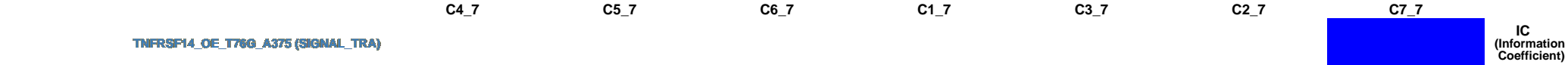


ACTB\_OE\_C77T\_A375 (PUJANA\_BRC)

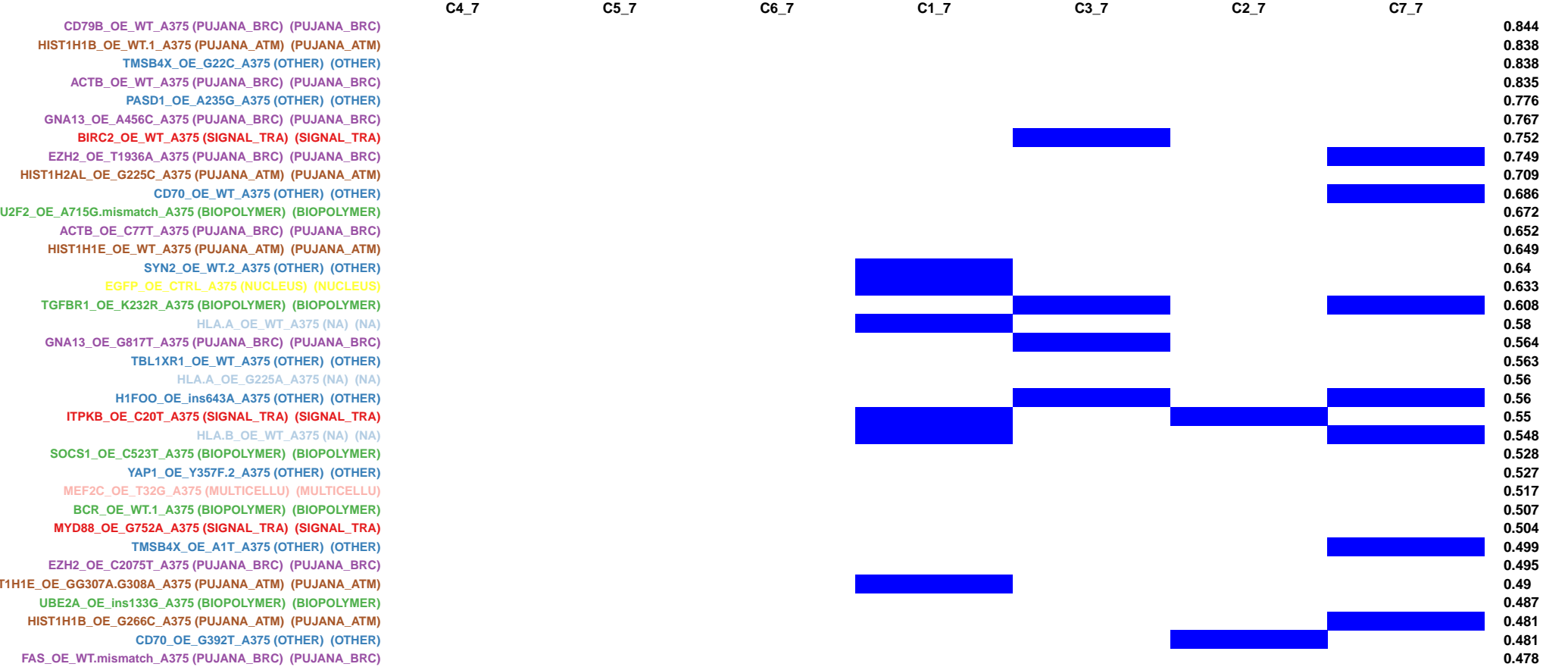




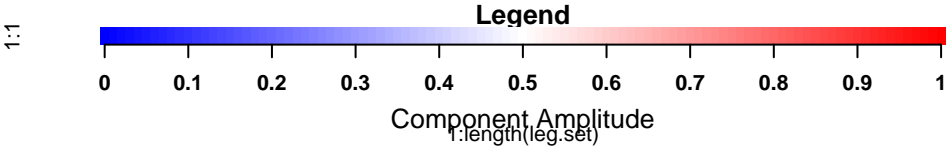
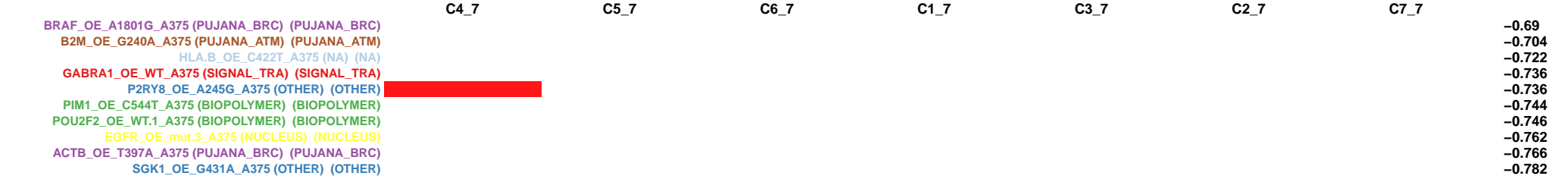
TNFRSF14\_OE\_T76G\_A375 (SIGNAL\_TRA)



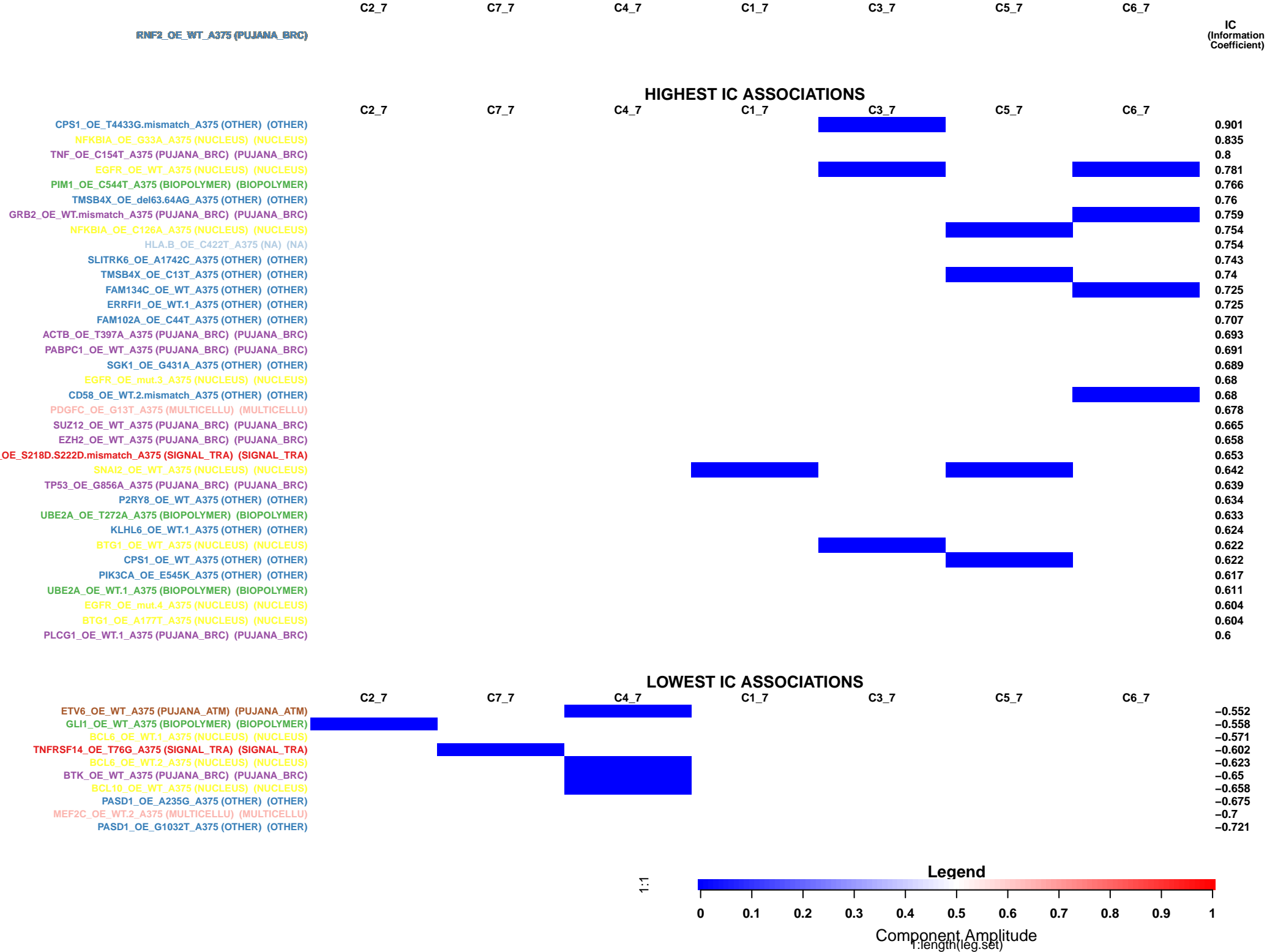
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS



RNF2\_OE\_WT\_A375 (PUJANA\_BRC)

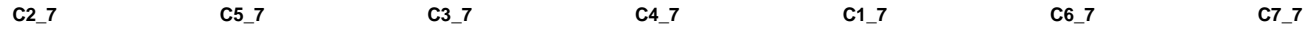




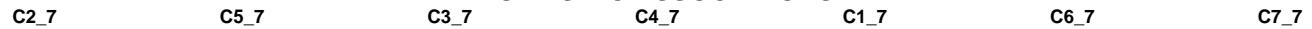
C2_7	C5_7	C3_7	C4_7	C1_7	C6_7	C7_7
------	------	------	------	------	------	------

**FBXO11\_OE\_G1799A\_A375 (NUCLEUS)**

C2\_7                      C5\_7                      C3\_7                      C4\_7                      C1\_7                      C6\_7                      C7\_7



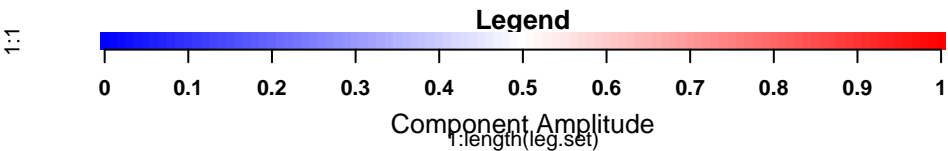
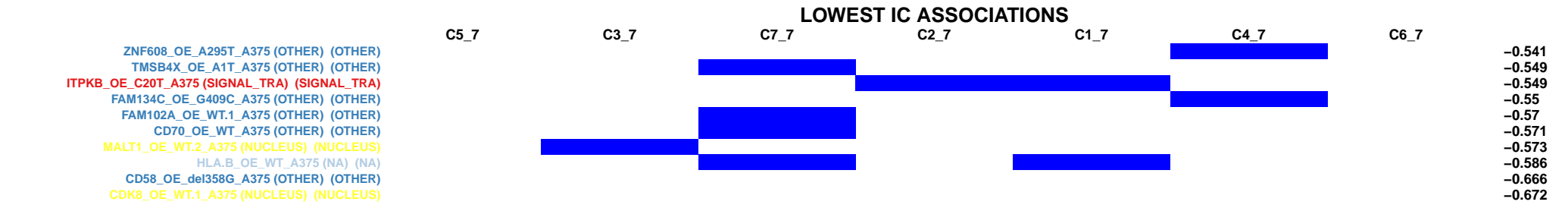
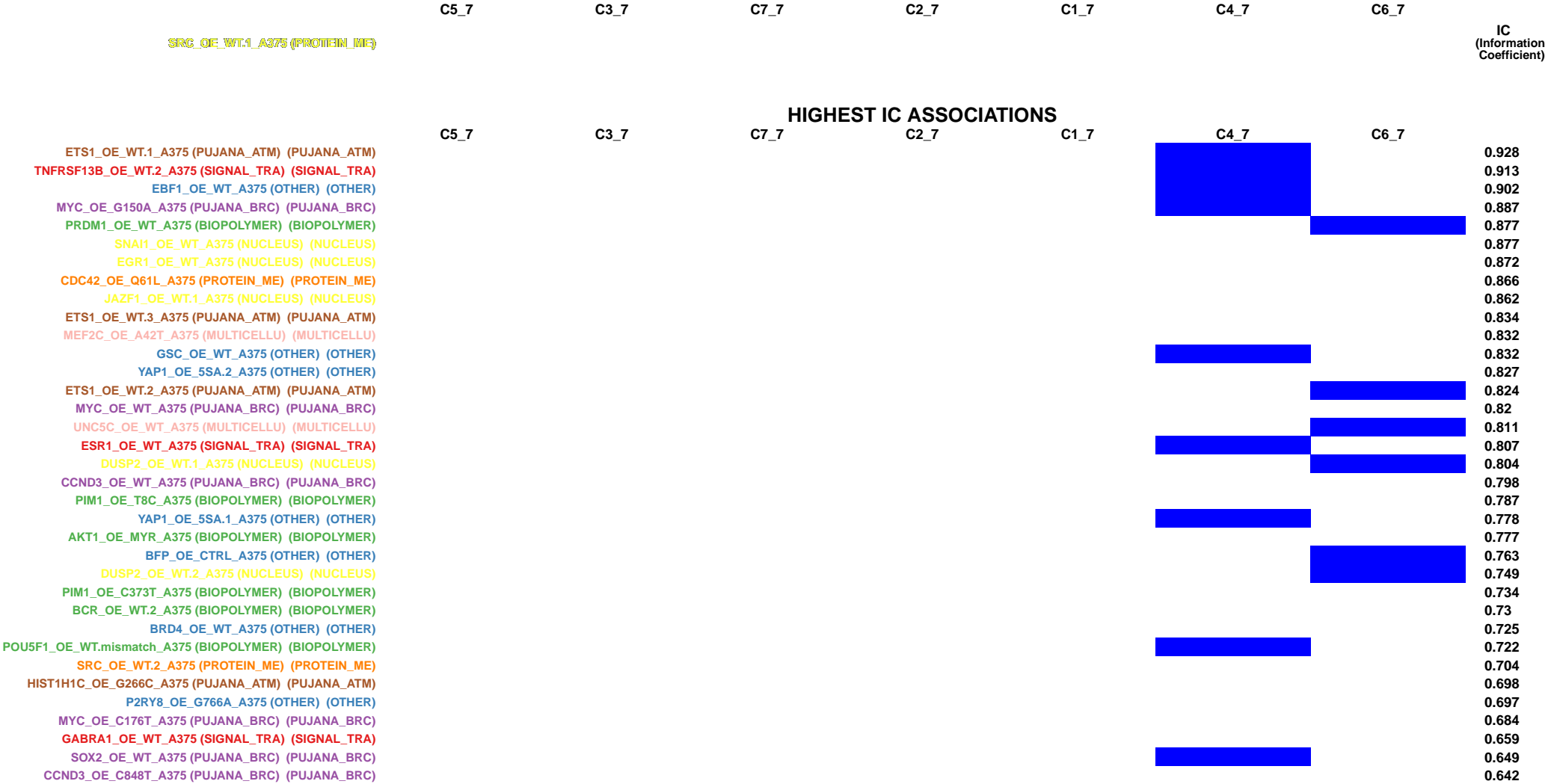
C2_7	C5_7	C3_7	C4_7	C1_7	C6_7	C7_7
------	------	------	------	------	------	------



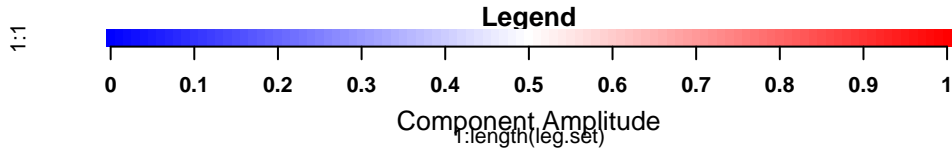
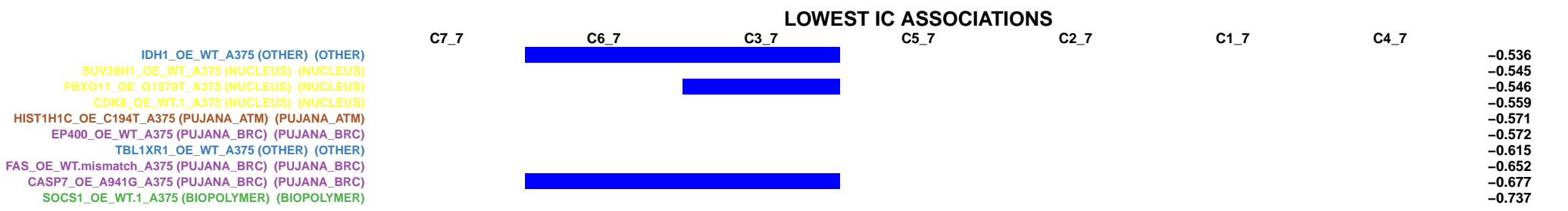
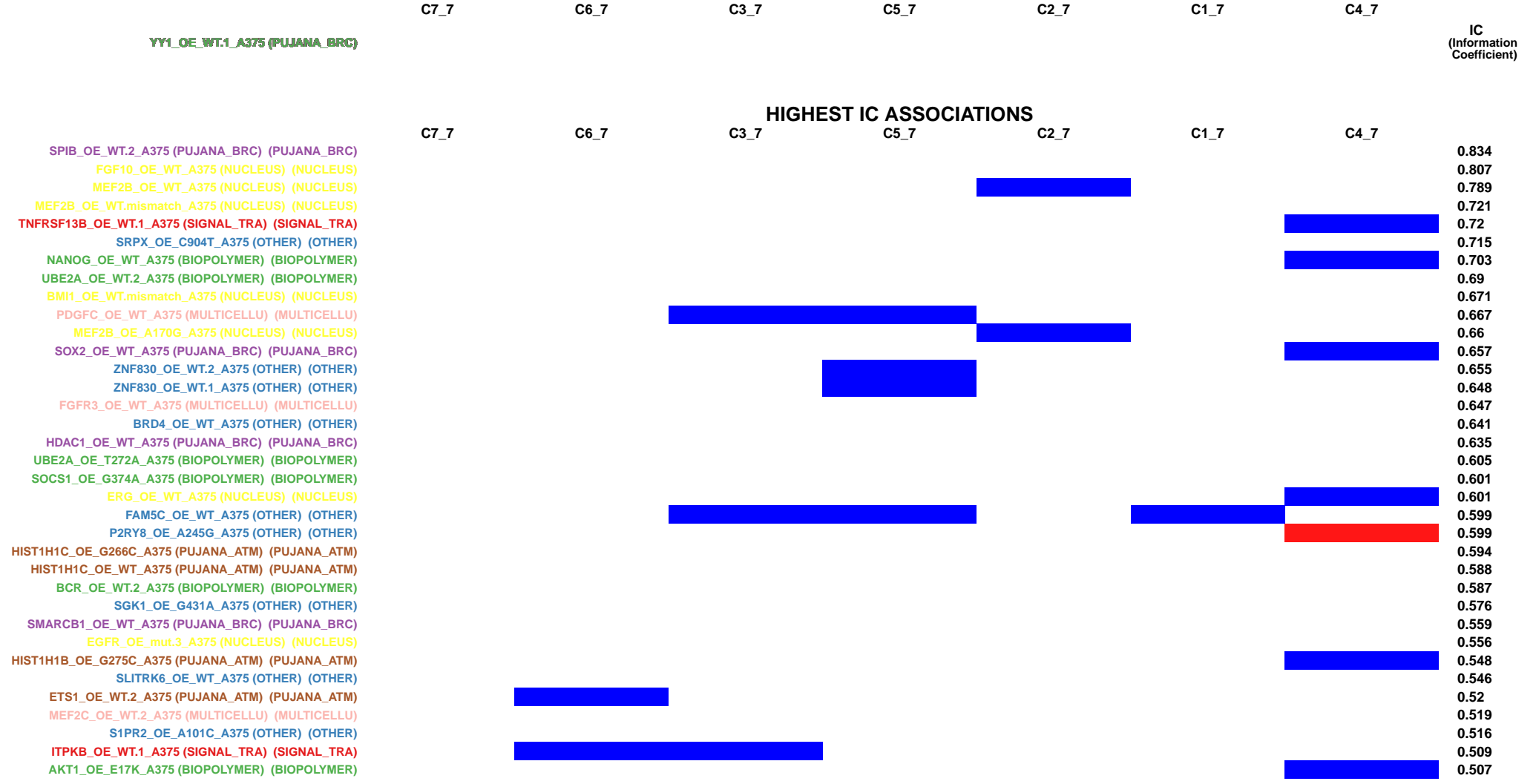
### Legend

0      0.1      0.2      0.3      0.4      0.5      0.6      0.7      0.8      0.9      1

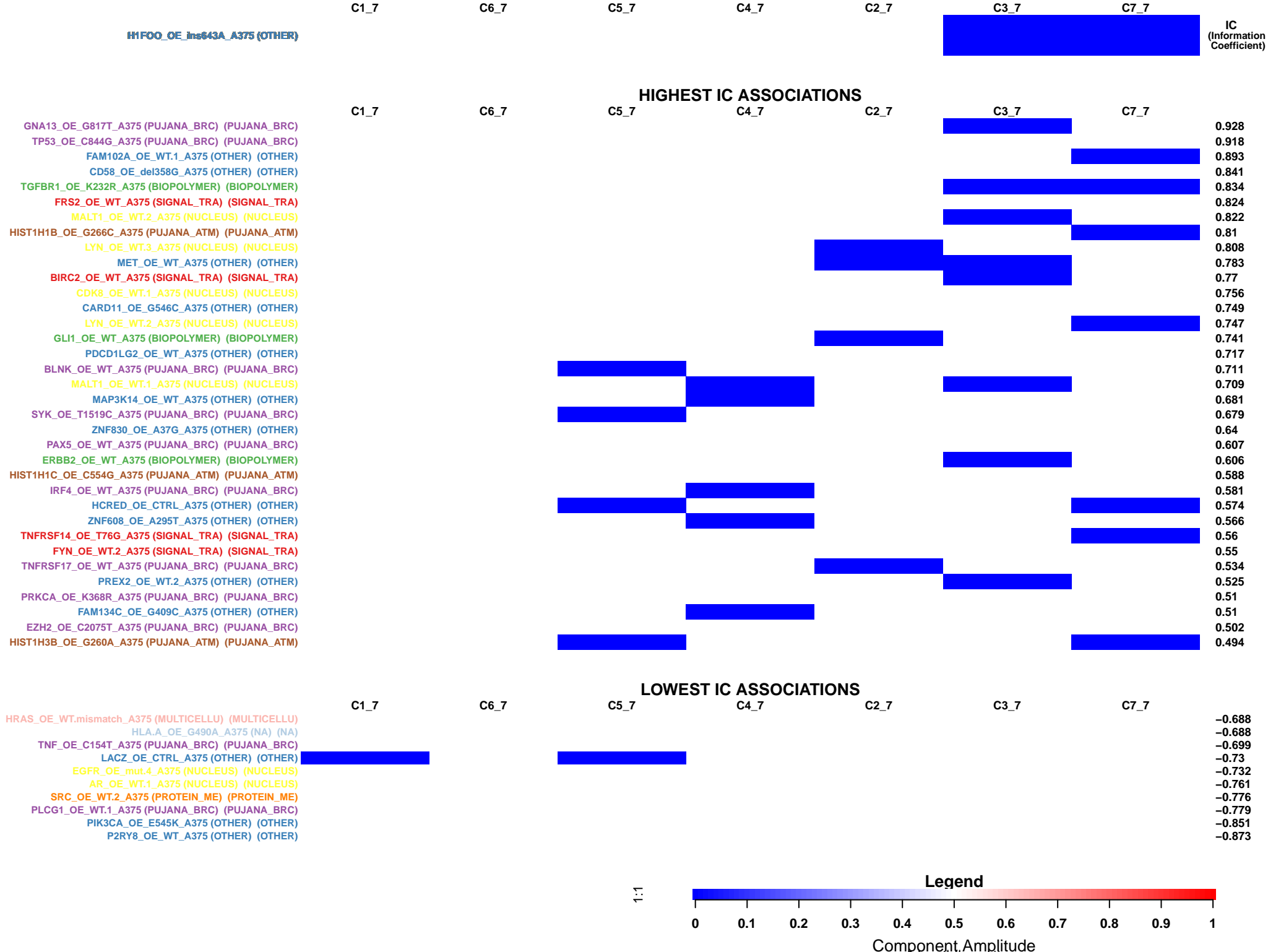
SRC\_OE\_WT.1\_A375 (PROTEIN\_ME)



YY1\_OE\_WT.1\_A375 (PUJANA\_BRC)



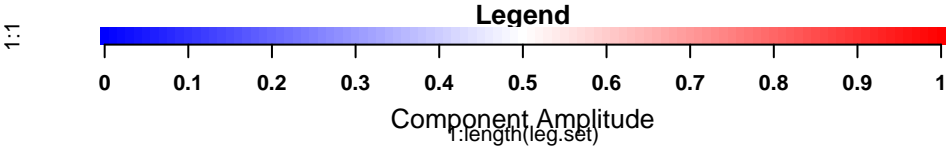
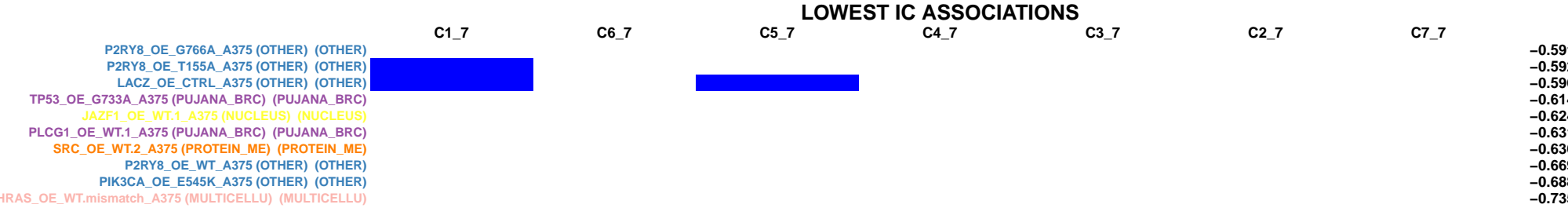
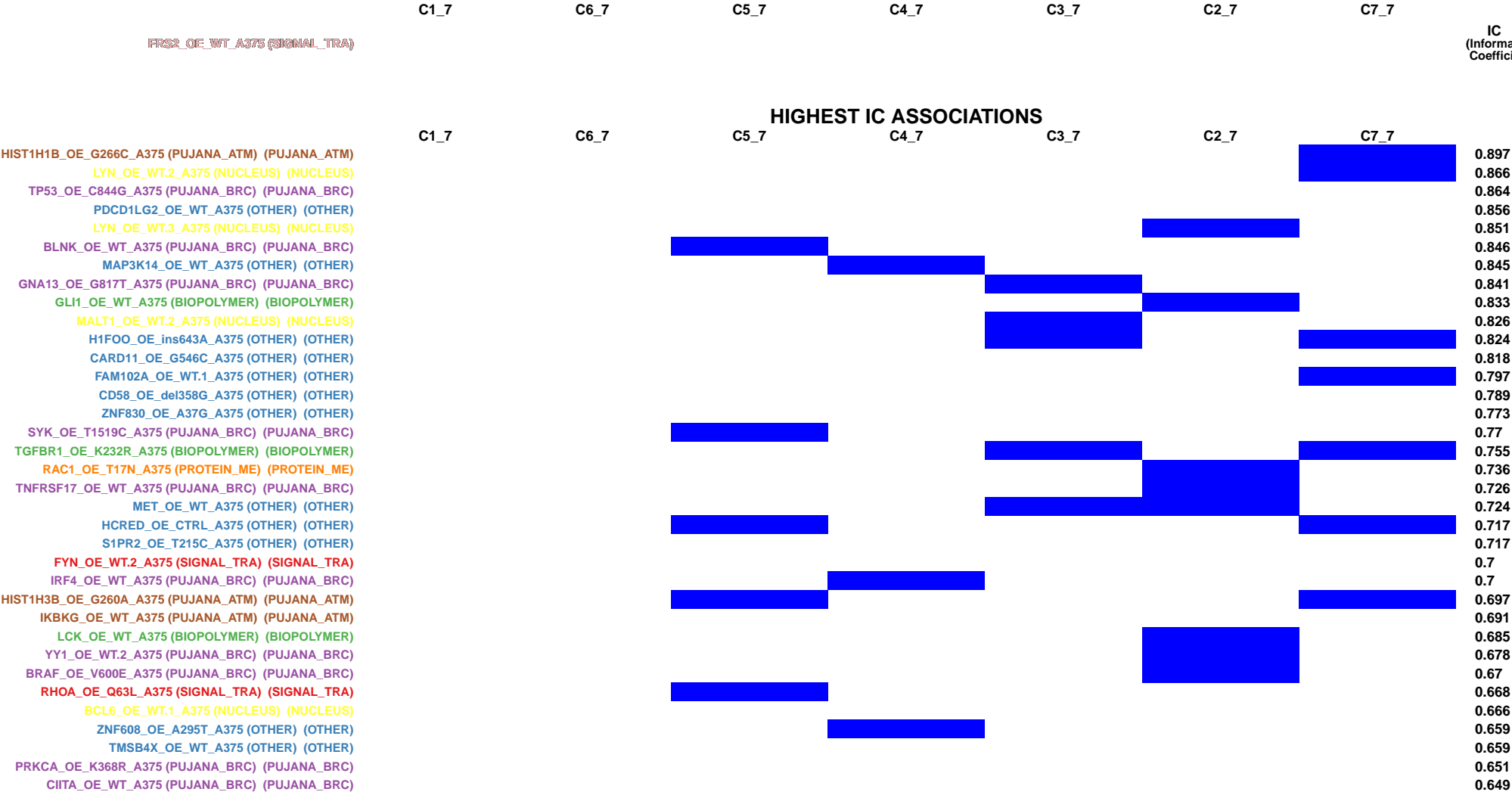
H1FOO\_OE\_ins643A\_A375 (OTHER)



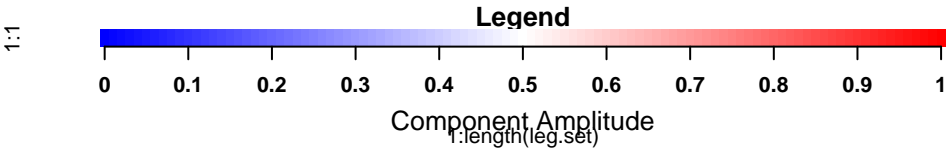
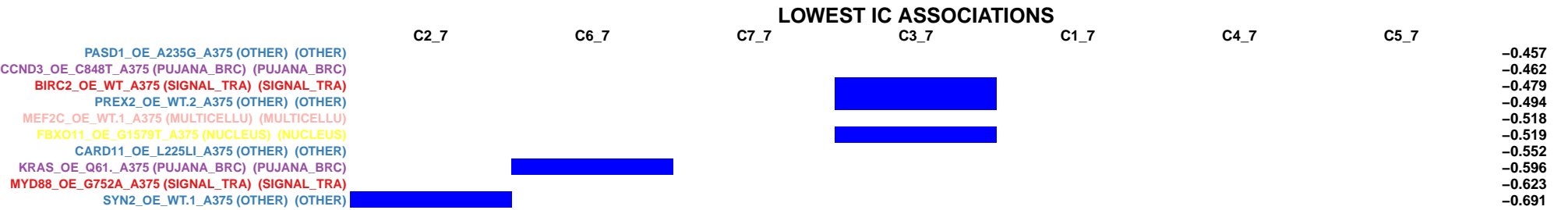
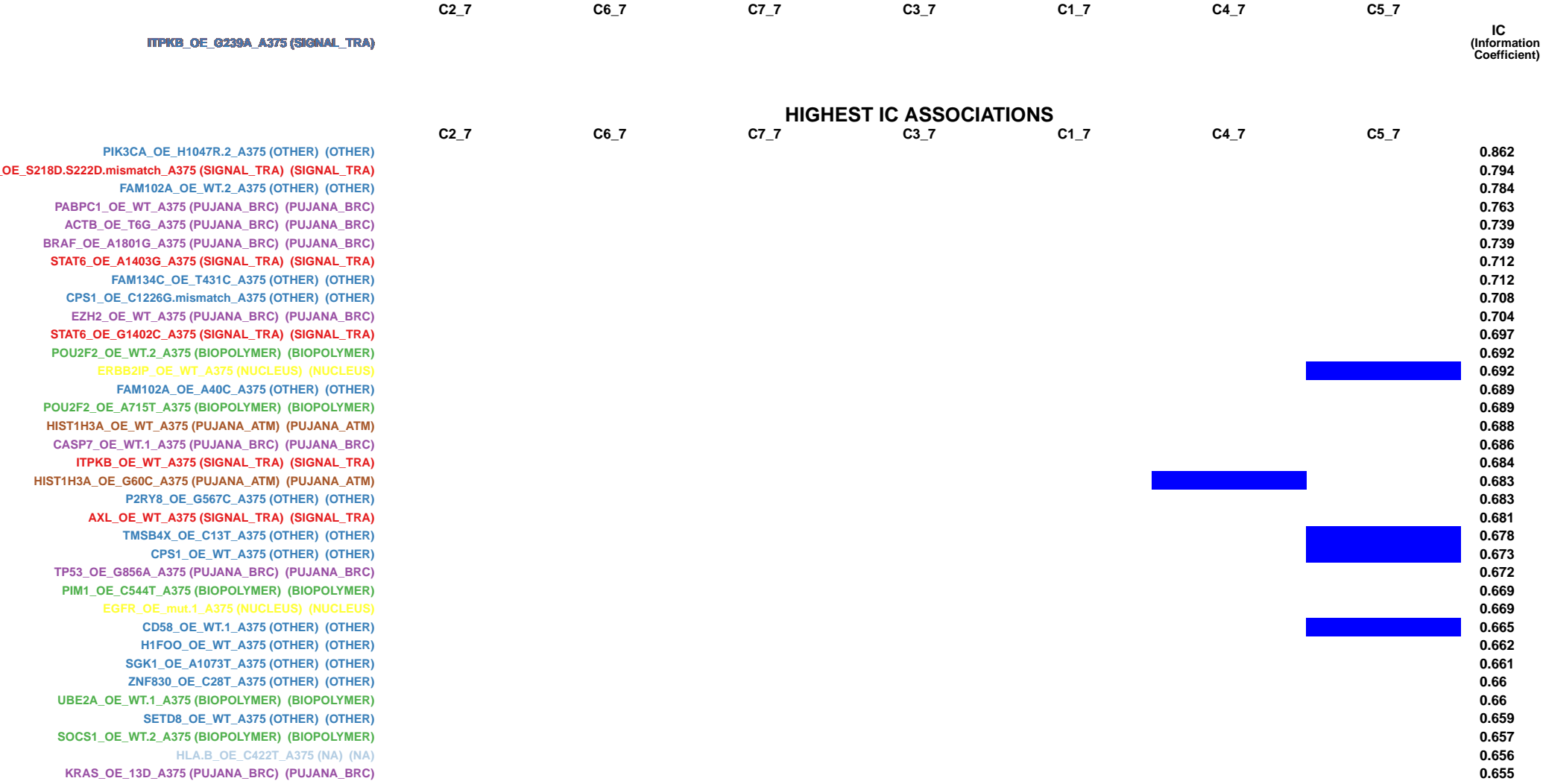
FRS2\_OE\_WT\_A375 (SIGNAL\_TRA)

FRS2\_OE\_WT\_A375 (SIGNAL\_TRA)

IC  
(Information  
Coefficient)

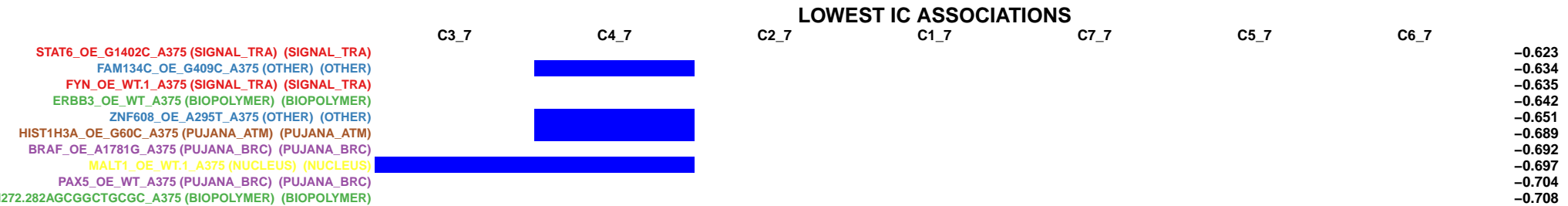


ITPKB\_OE\_G239A\_A375 (SIGNAL\_TRA)



C3_7	C4_7	C2_7	C1_7	C7_7	C5_7	C6_7
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**CARD11\_OE\_Q1876A\_A375 (OTHER)**



FBXO11\_OE\_WT\_A375 (NUCLEUS)

FBXO11\_OE\_WT\_A375 (NUCLEUS)

IC  
(Information  
Coefficient)

HIGHEST IC ASSOCIATIONS

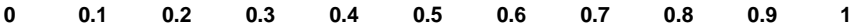


LOWEST IC ASSOCIATIONS



1:1

Legend



Component Amplitude

1:length(leg.set)



C6_7	C1_7	C3_7	C5_7	C7_7	C2_7	C4_7
------	------	------	------	------	------	------

IC  
Information  
Coefficient)

C6\_7      C1\_7      C3\_7      C5\_7      C7\_7      C2\_7      C4\_7

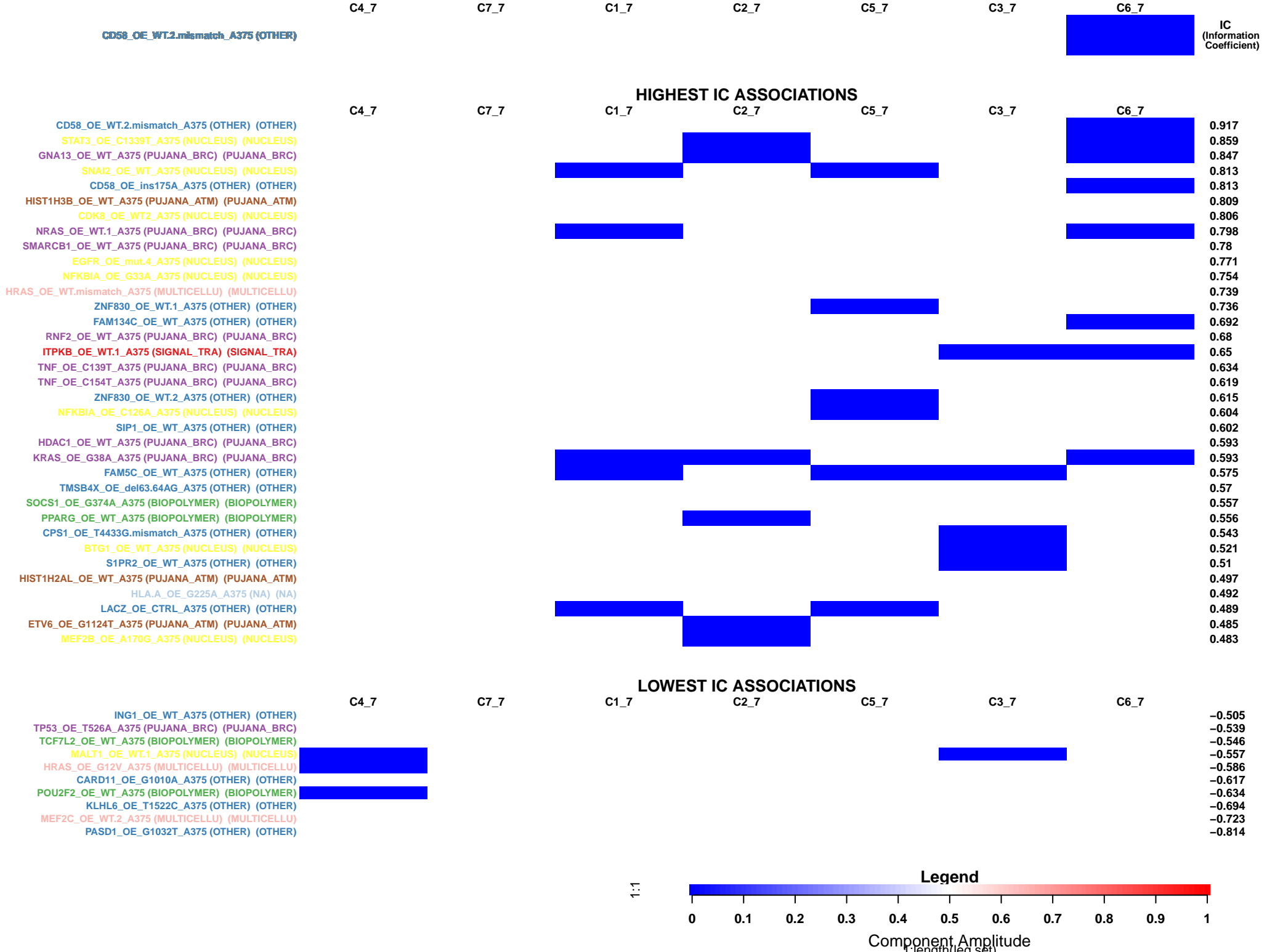


C6_7	C1_7	C3_7	C5_7	C7_7	C2_7	C4_7
------	------	------	------	------	------	------

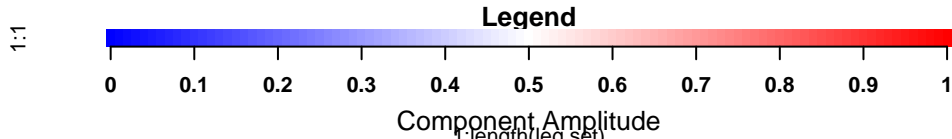
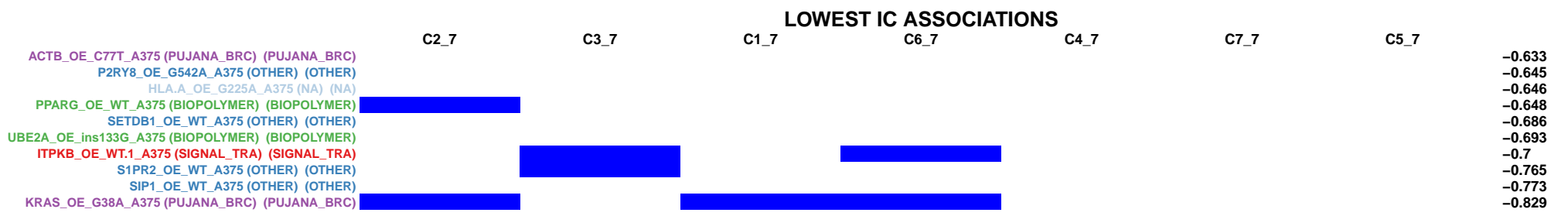
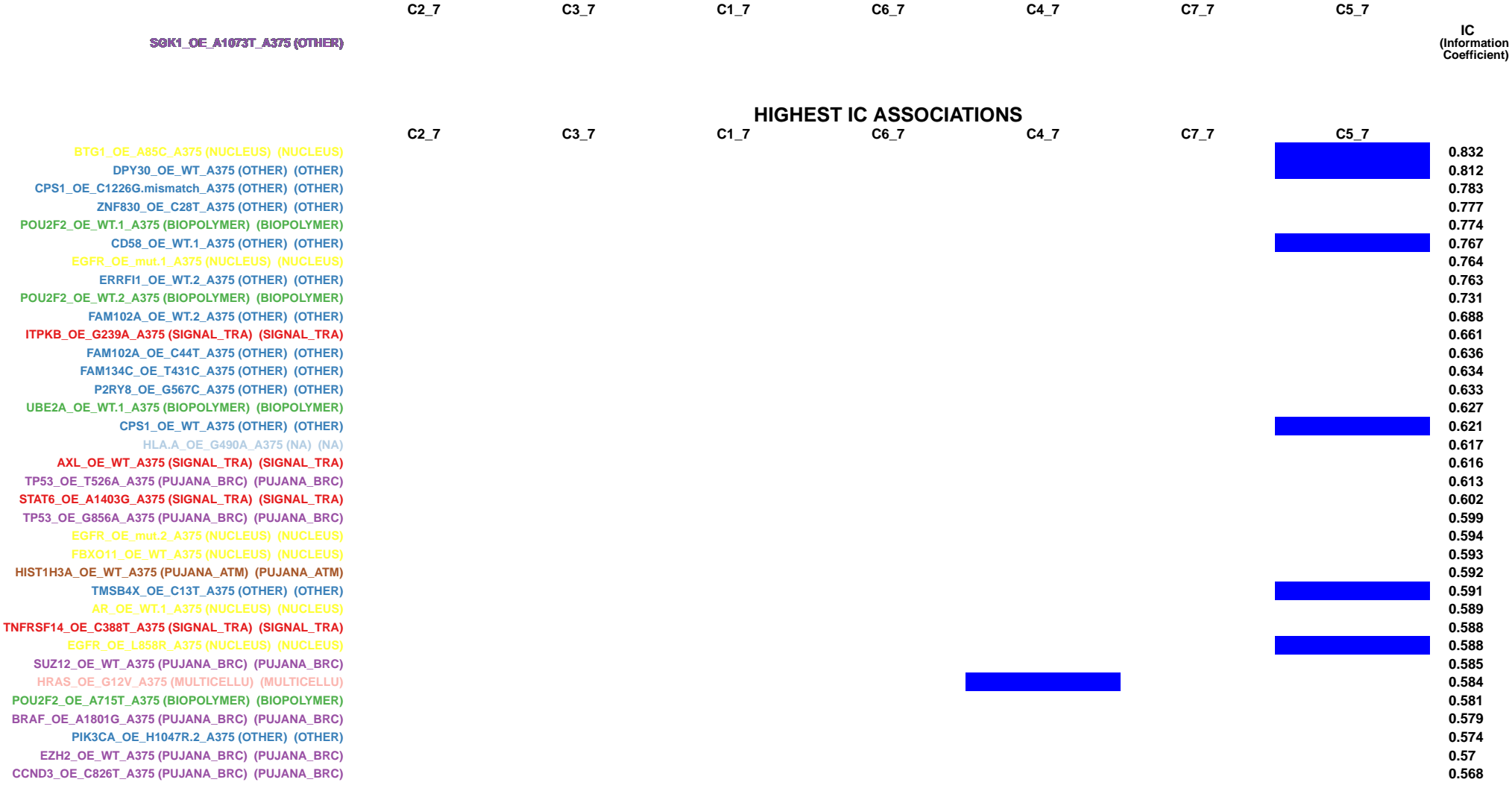


```
1:length(leg.set)
```

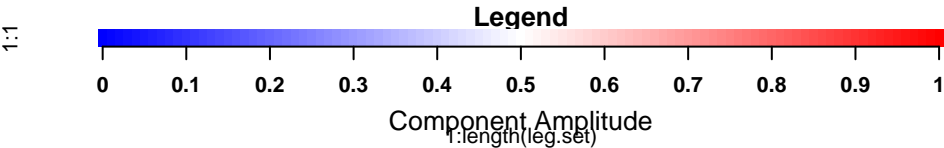
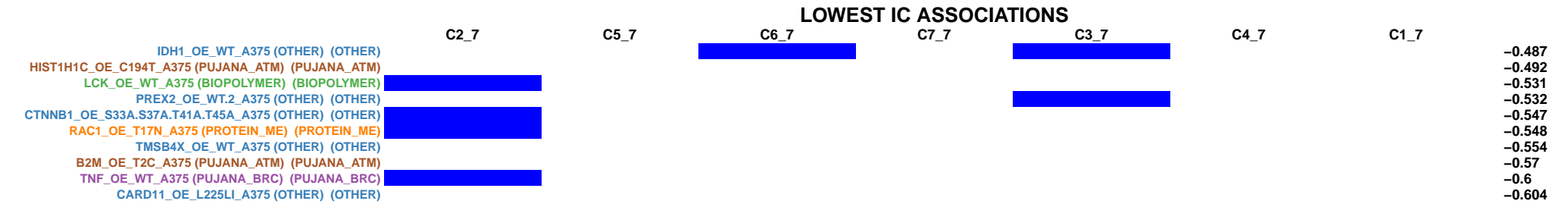
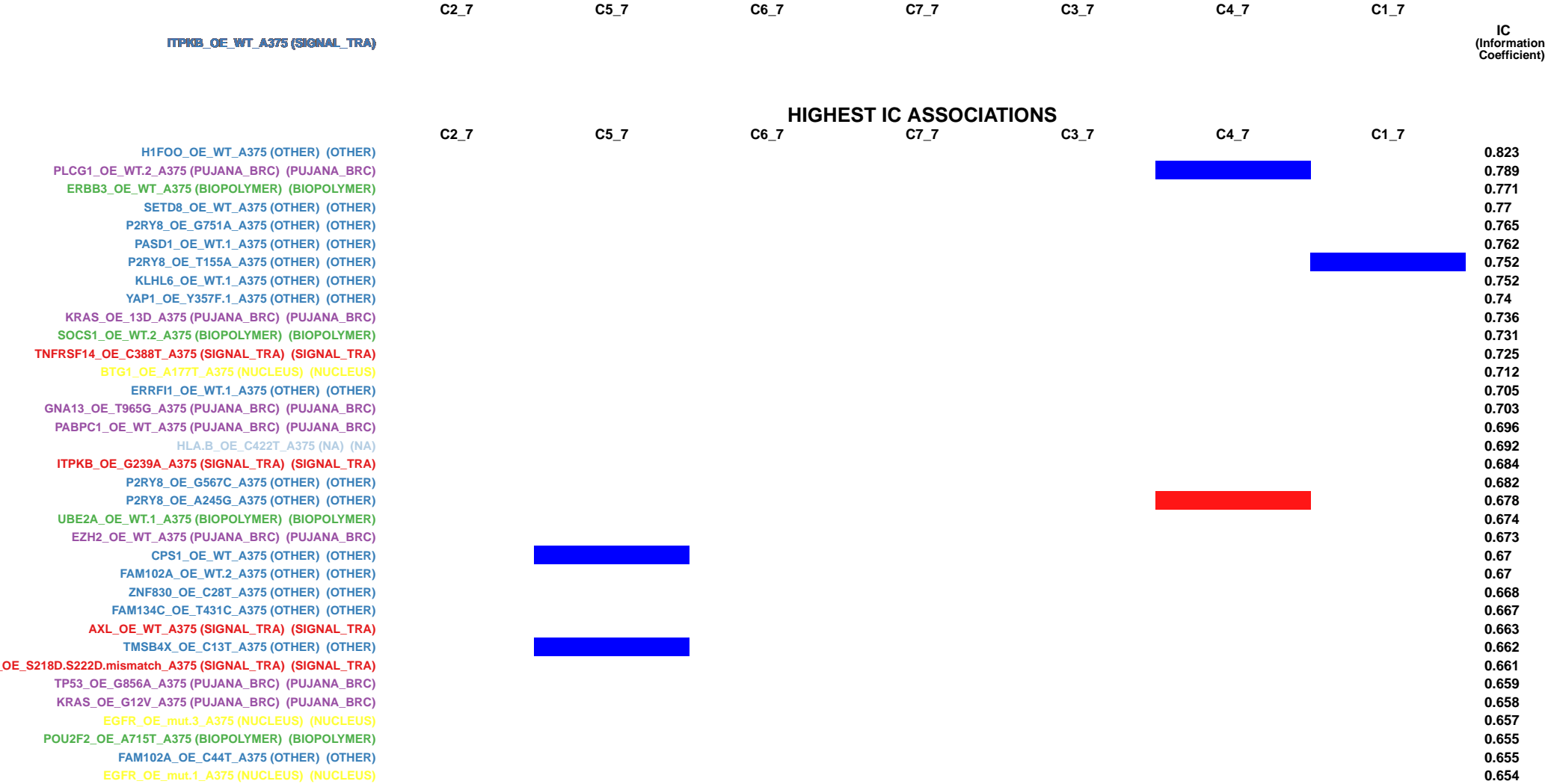
CD58\_OE\_WT.2.mismatch\_A375 (OTHER)



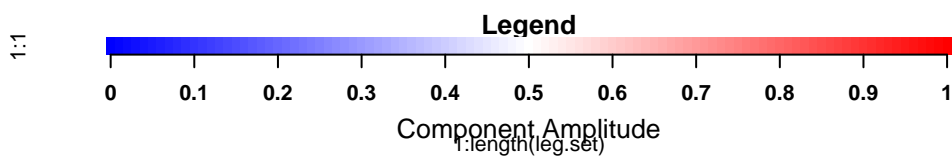
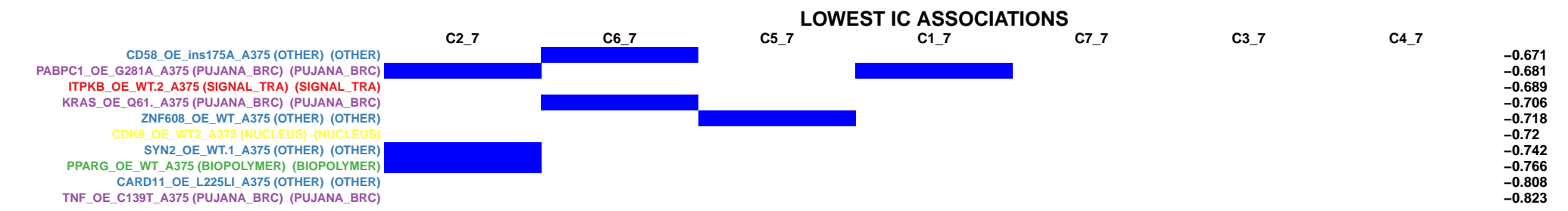
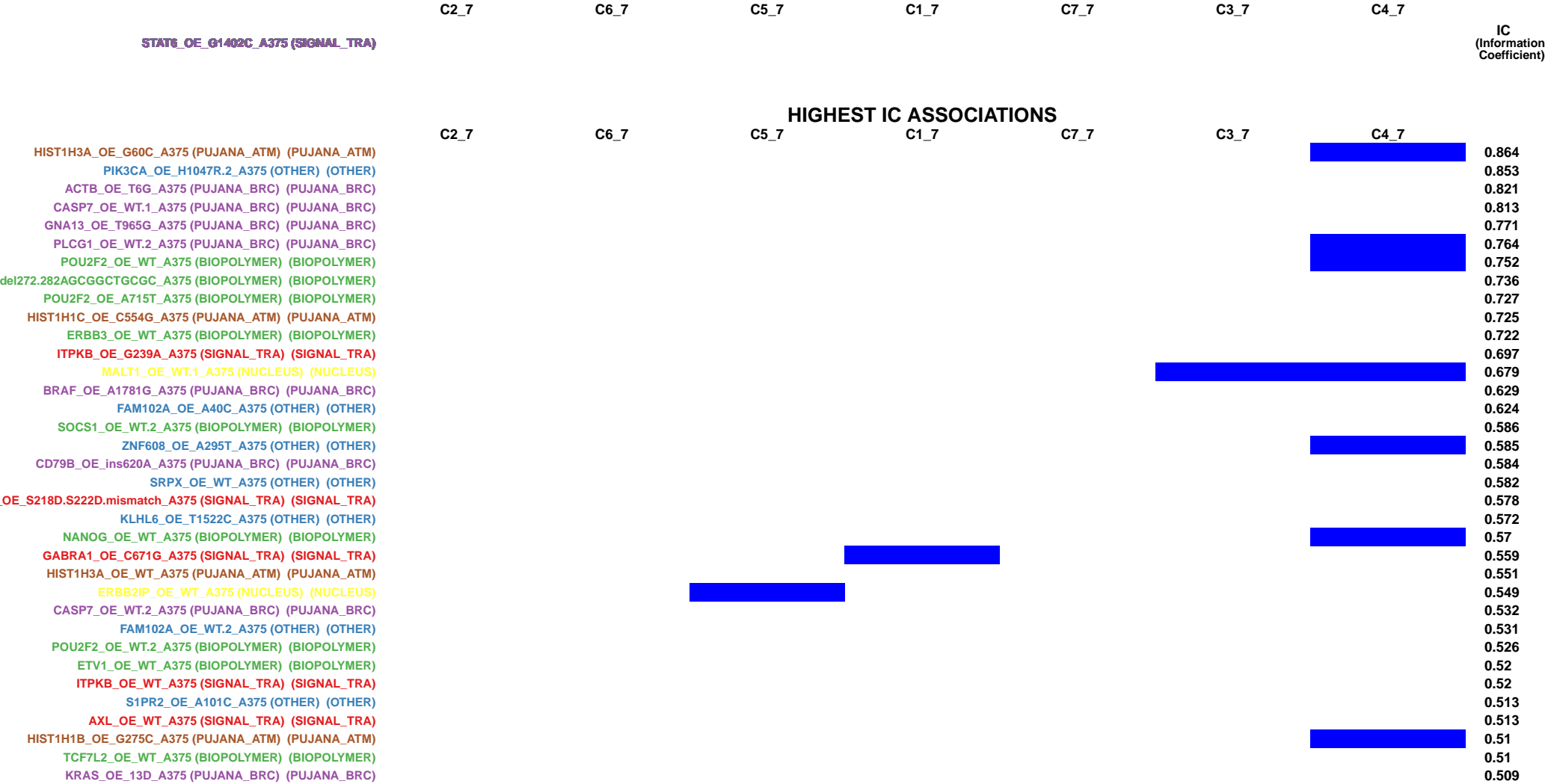
SGK1\_OE\_A1073T\_A375 (OTHER)



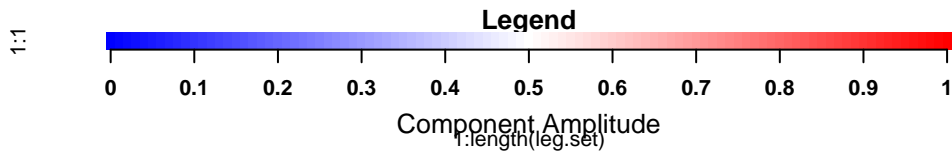
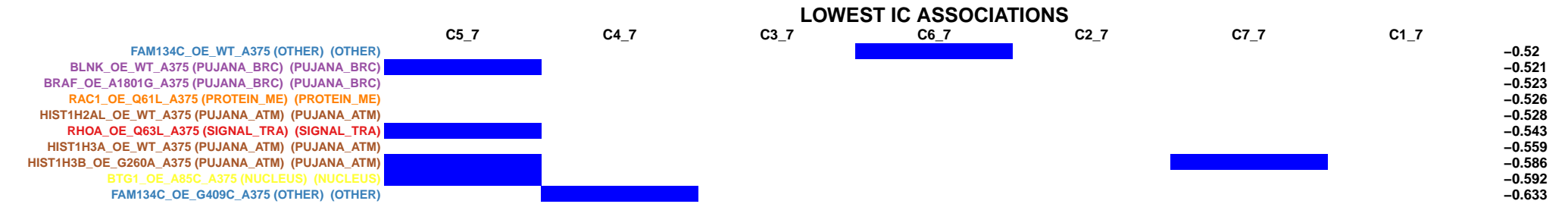
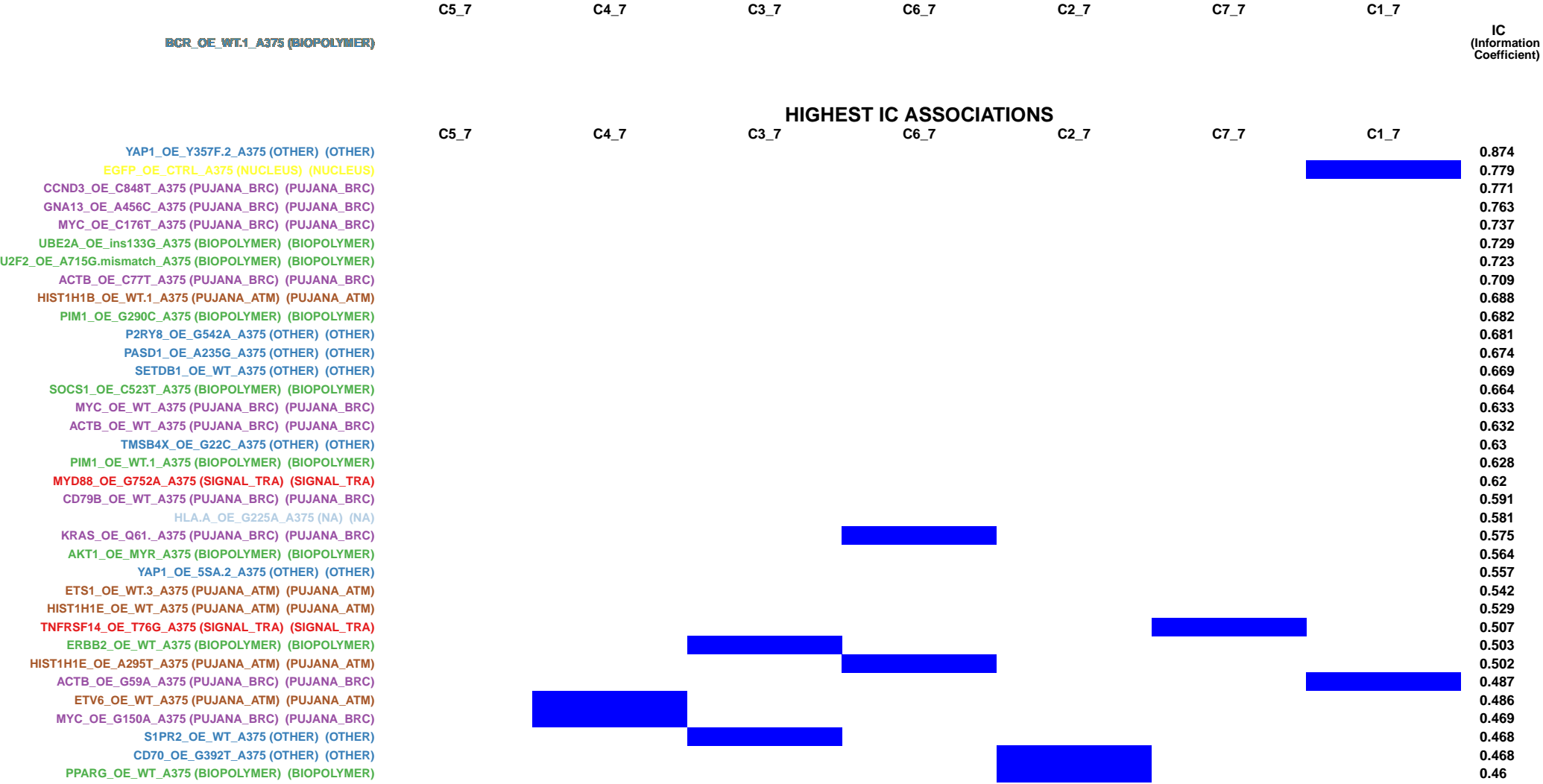
ITPKB\_OE\_WT\_A375 (SIGNAL\_TRA)



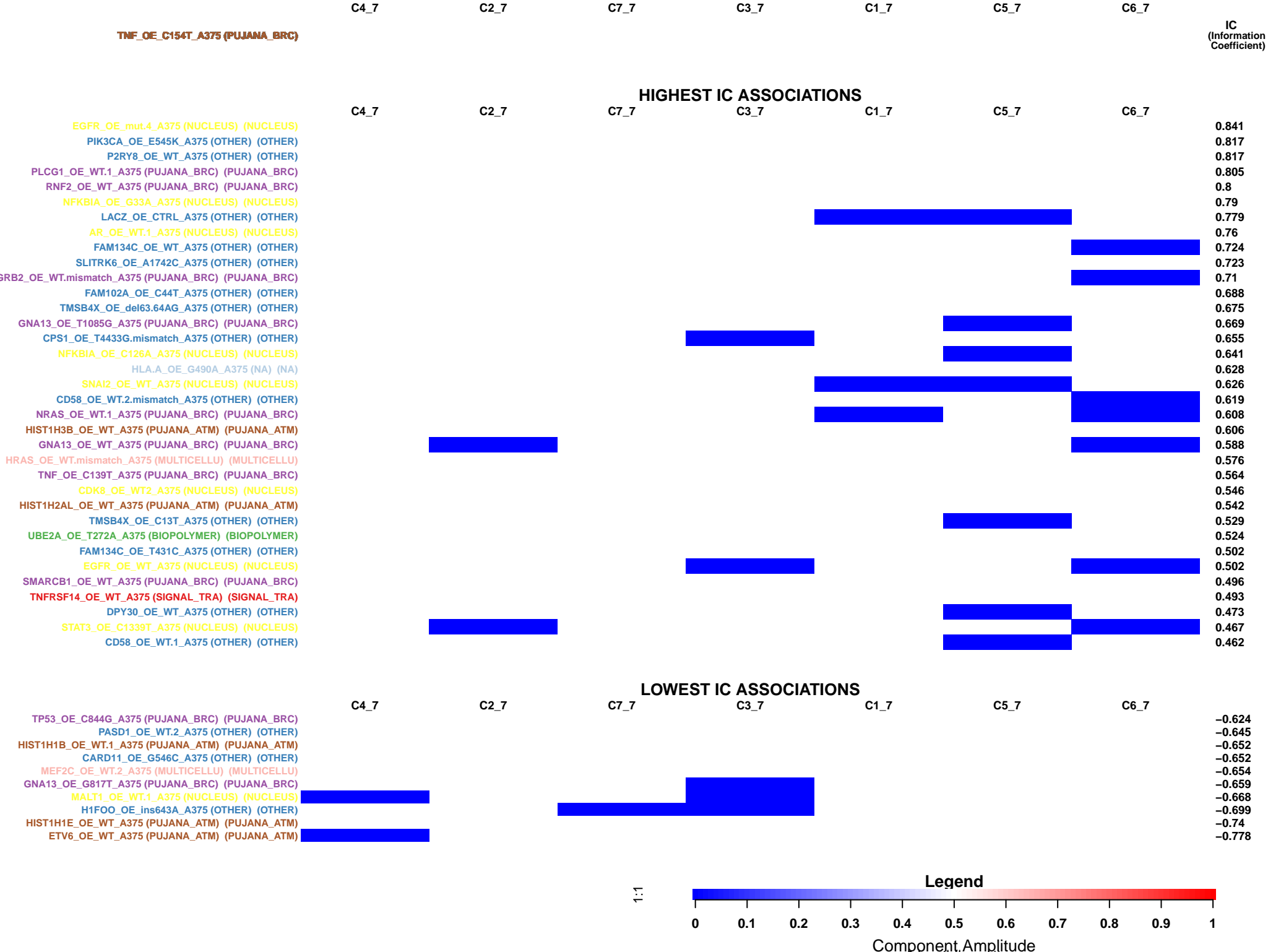
STAT6\_OE\_G1402C\_A375 (SIGNAL\_TRA)



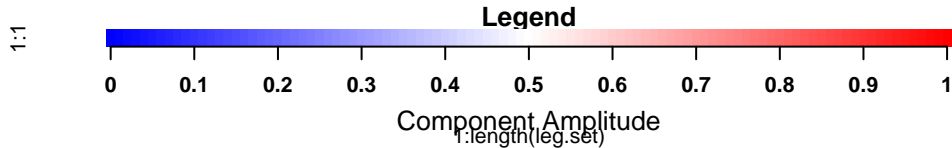
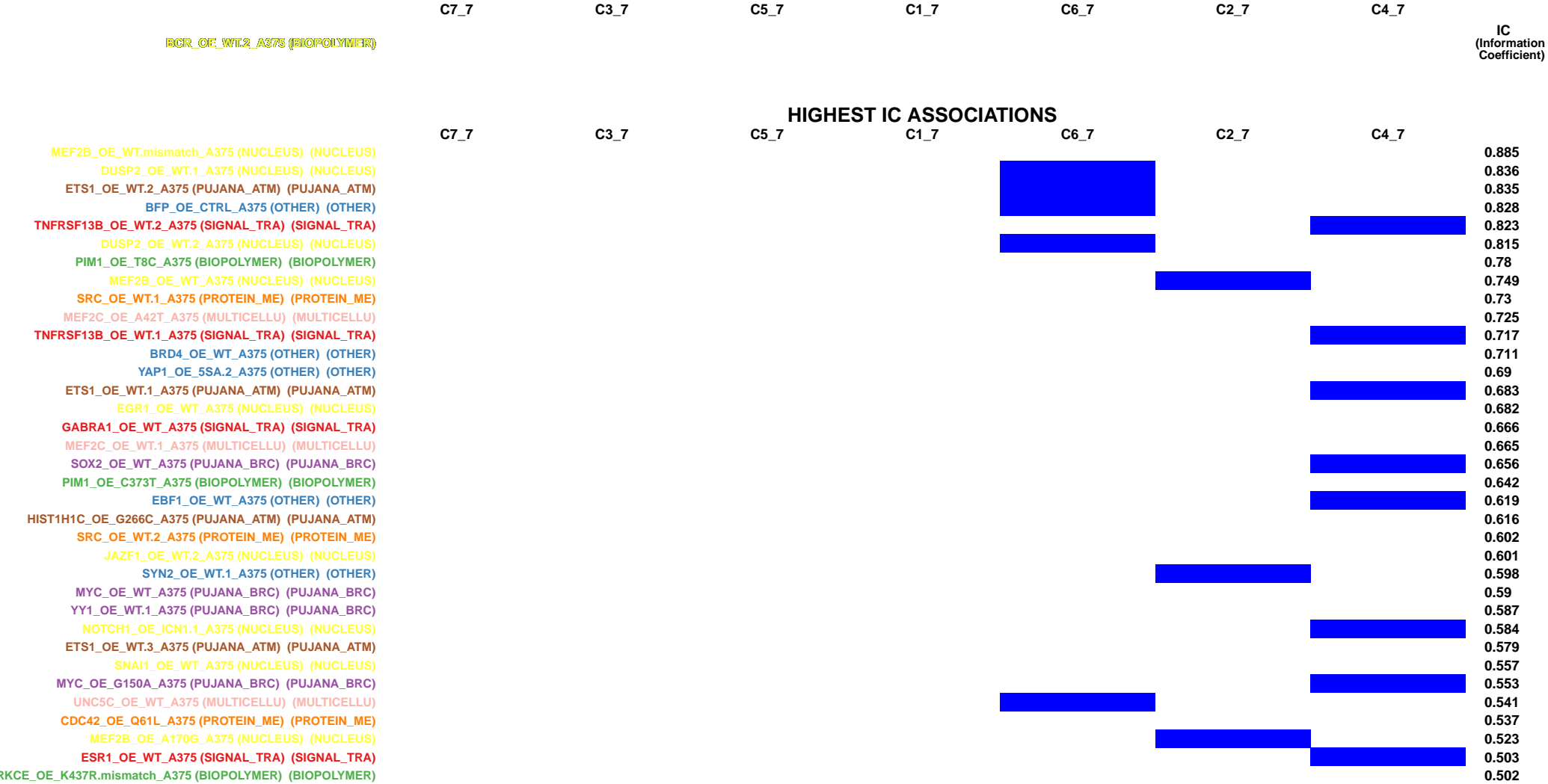
BCR\_OE\_WT.1\_A375 (BIOPOLYMER)



TNF\_OE\_C154T\_A375 (PUJANA\_BRC)

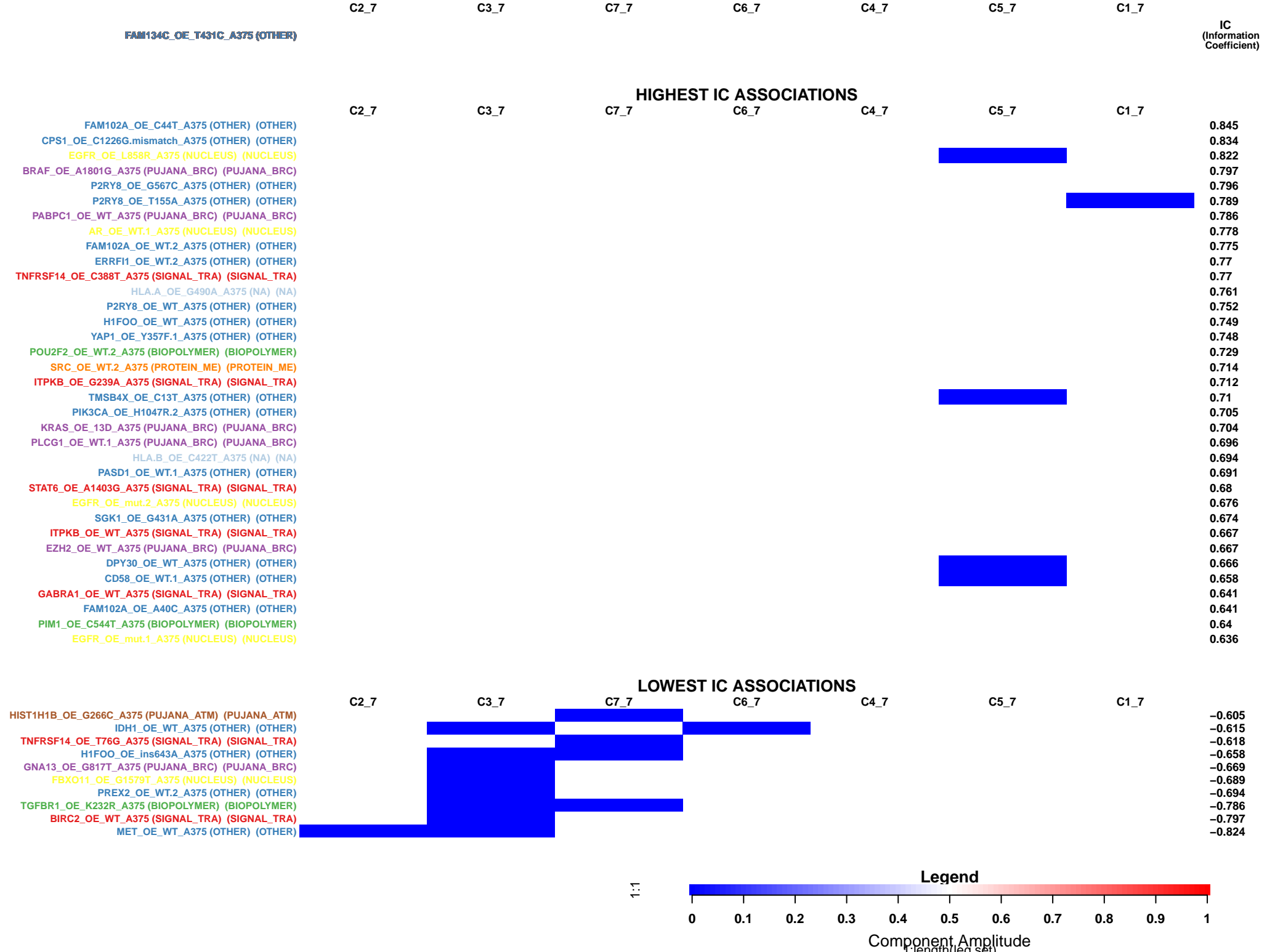


BCR\_OE\_WT.2\_A375 (BIOPOLYMER)

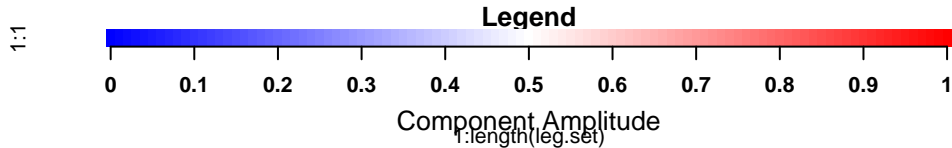
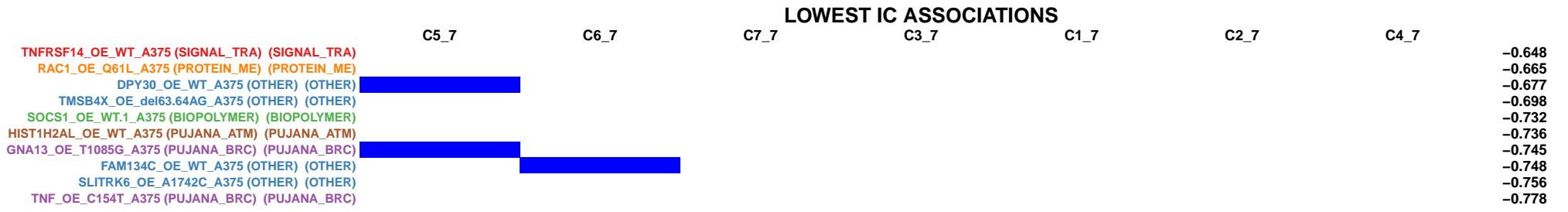
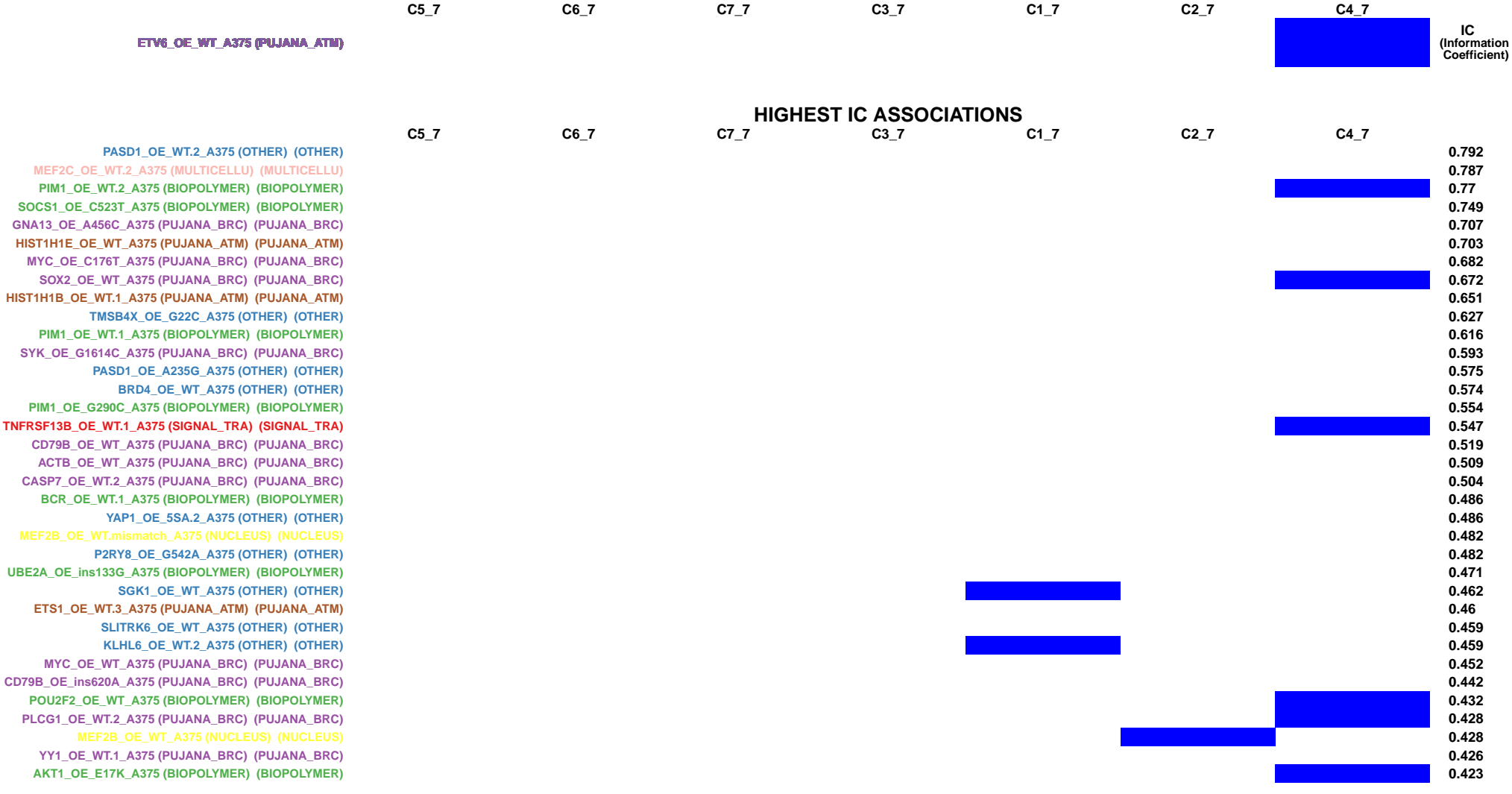




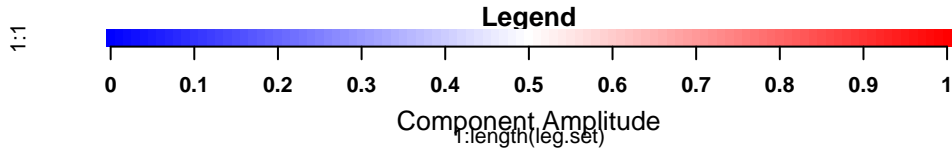
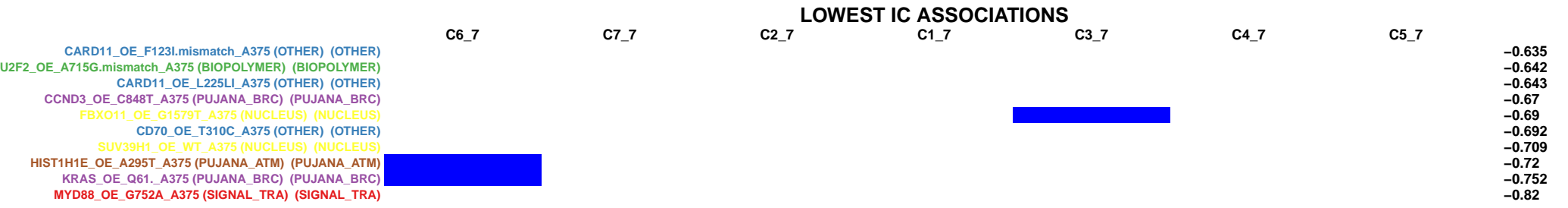
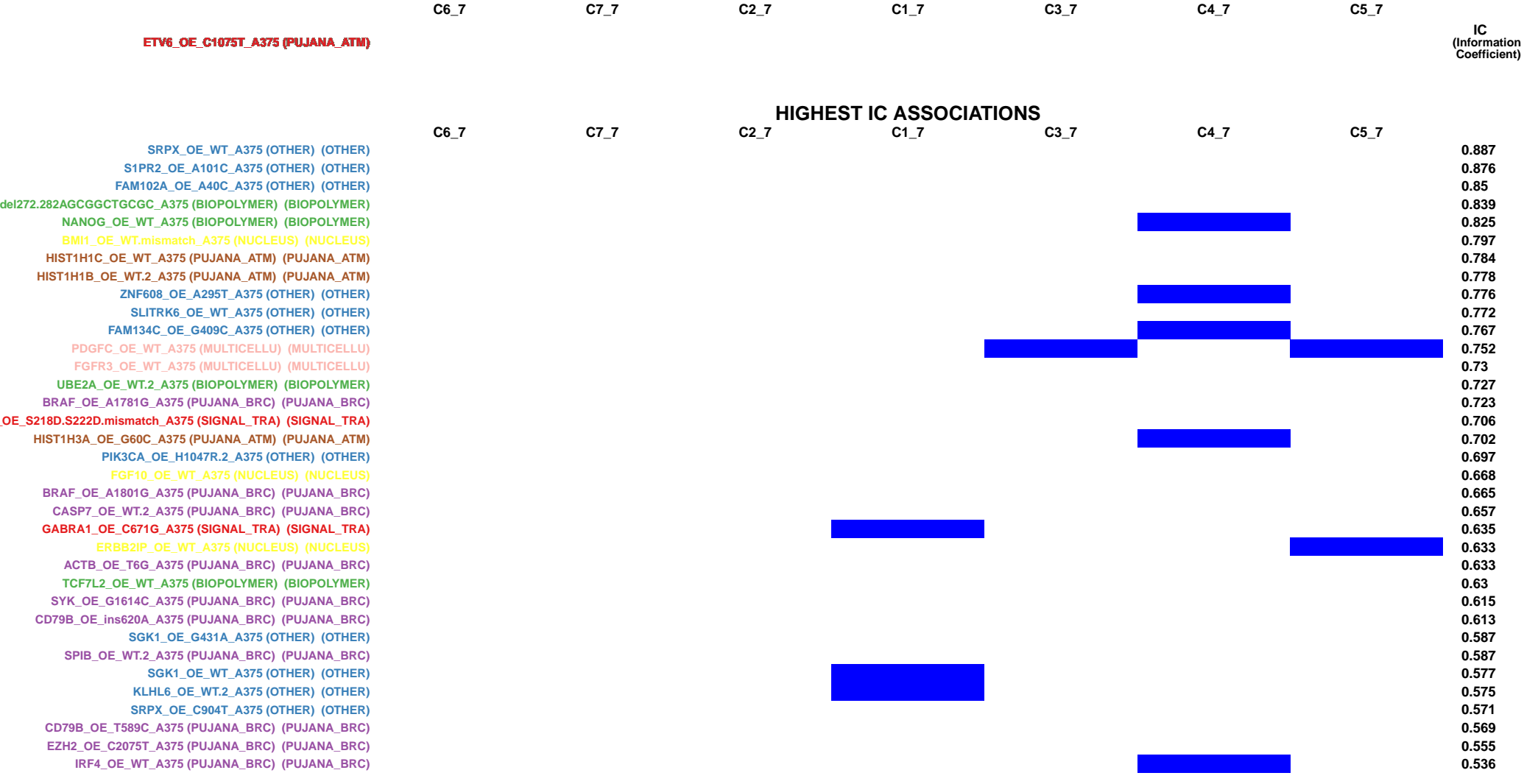
FAM134C\_OE\_T431C\_A375 (OTHER)



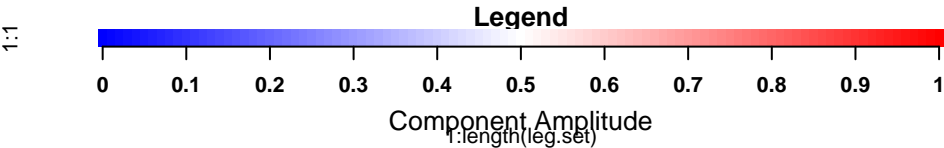
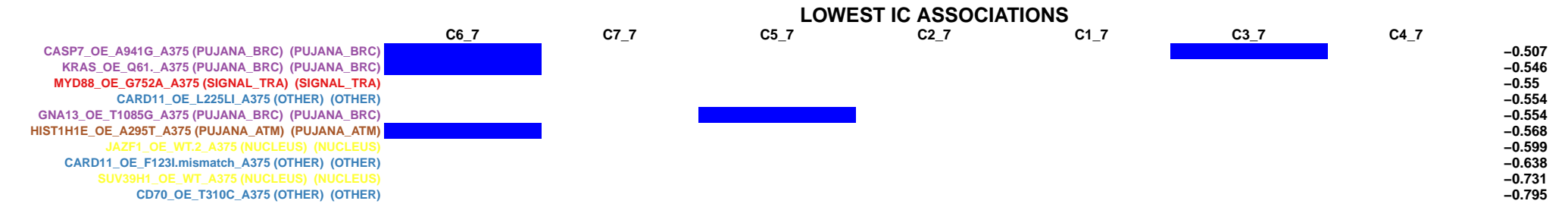
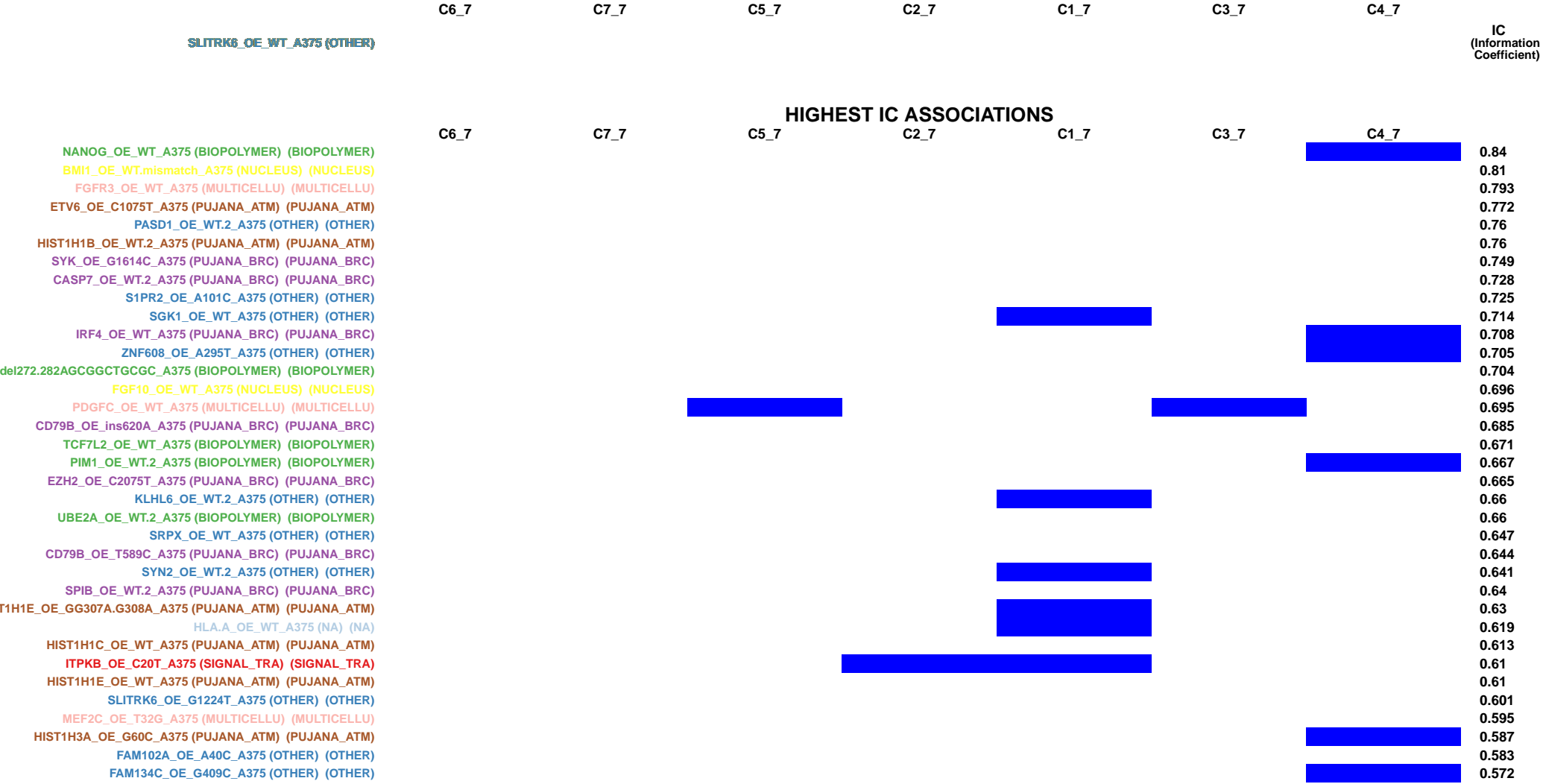
ETV6\_OE\_WT\_A375 (PUJANA\_ATM)



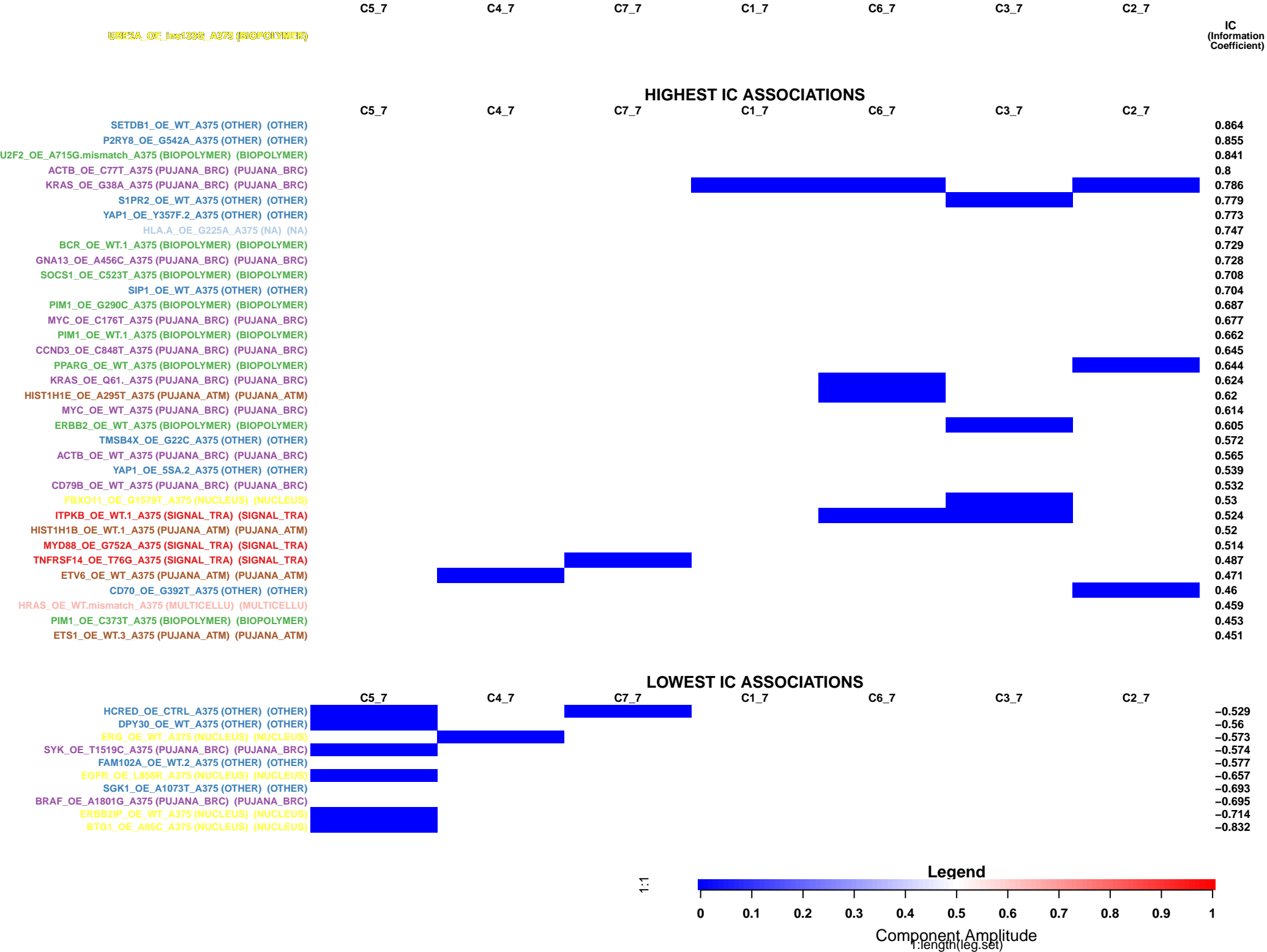
ETV6\_OE\_C1075T\_A375 (PUJANA\_ATM)



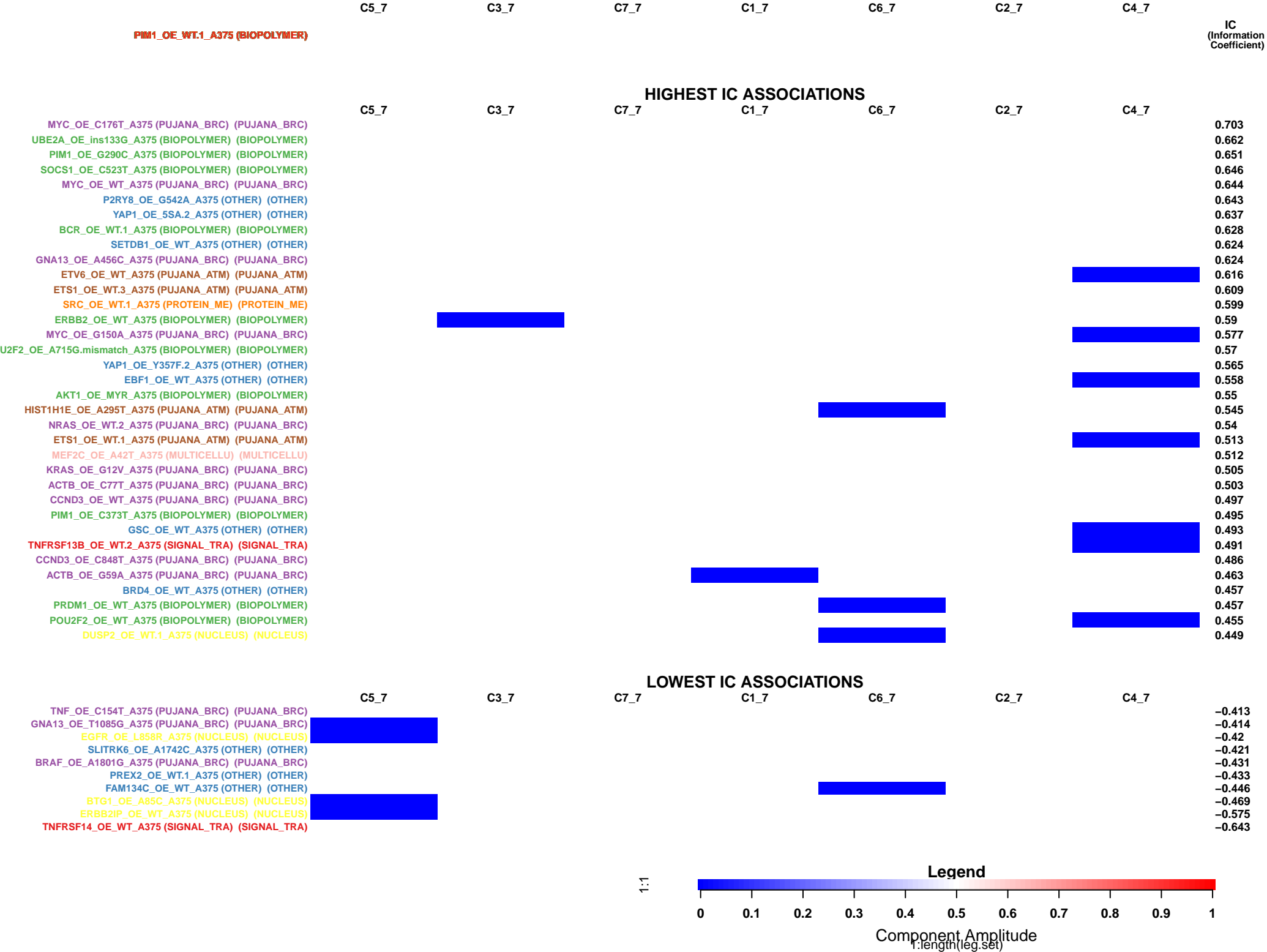
SLITRK6\_OE\_WT\_A375 (OTHER)



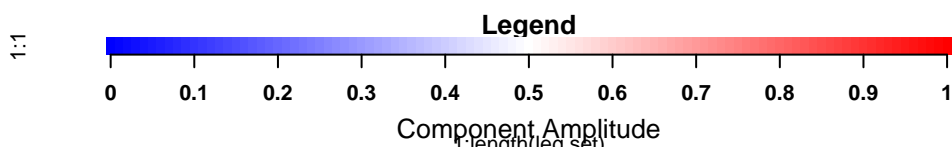
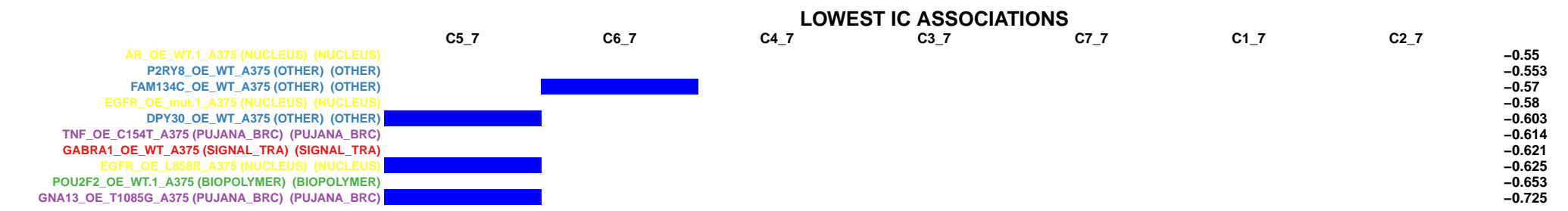
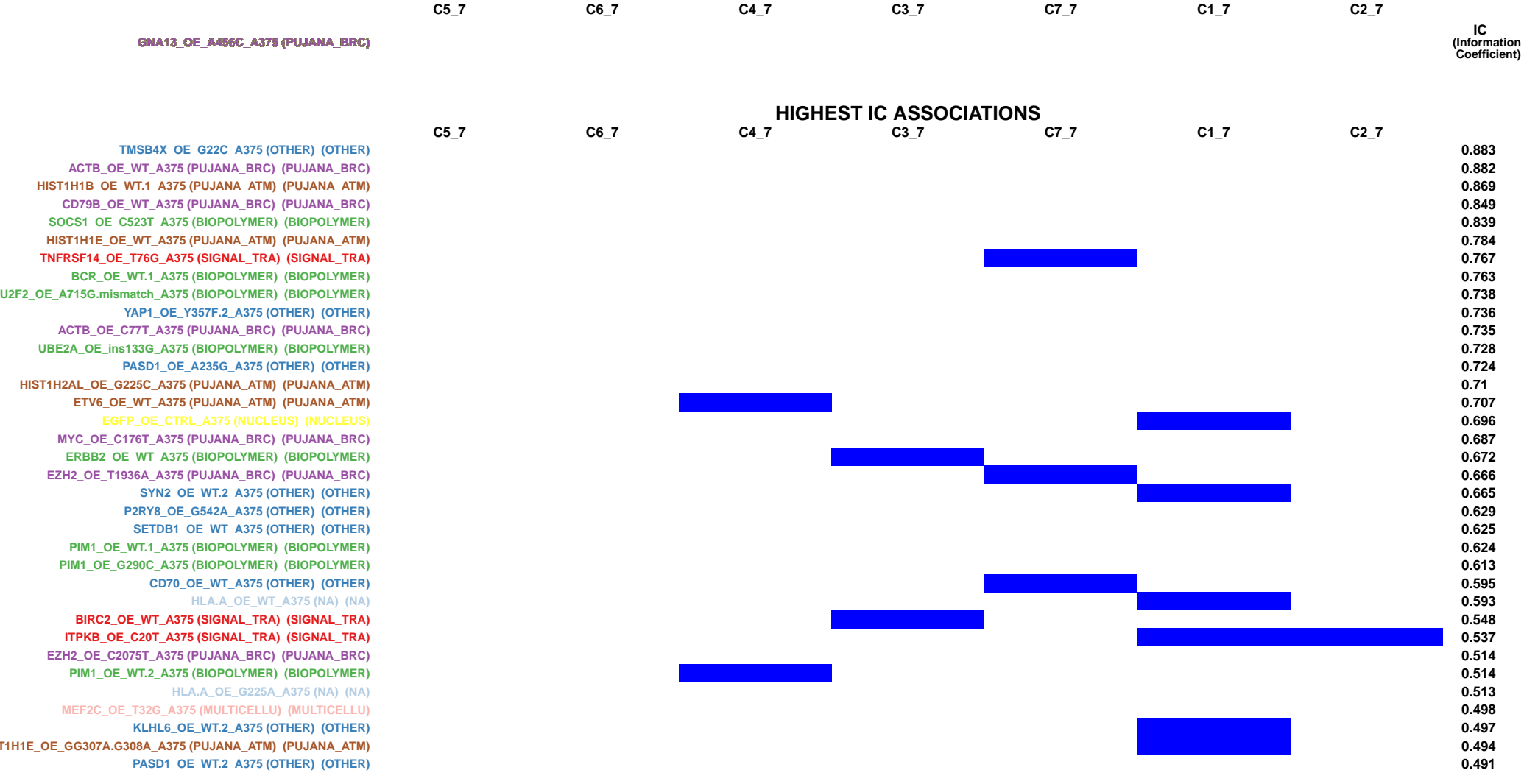
# UBE2A\_OE\_ins133G\_A375 (BIOPOLYMER)



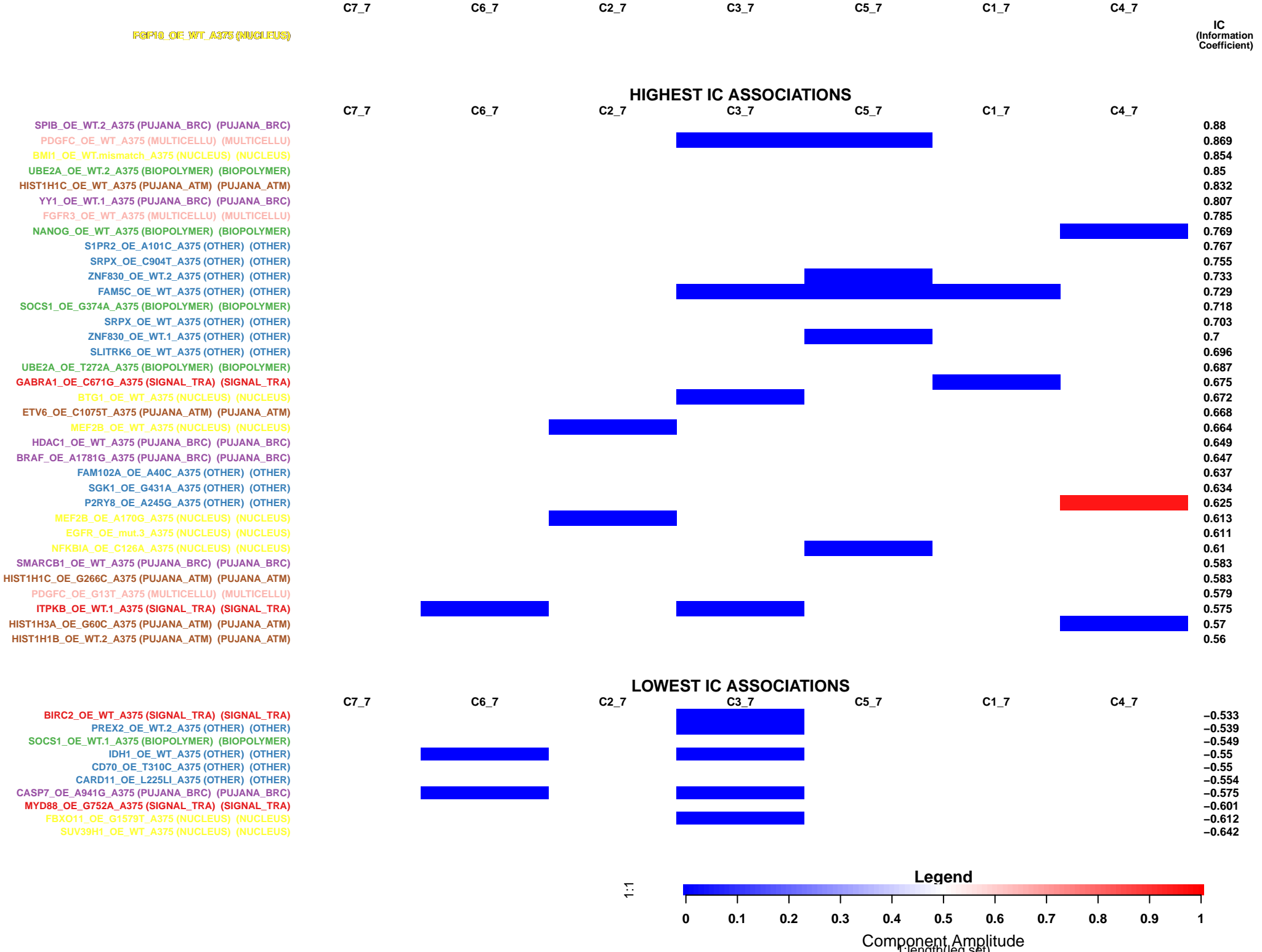
PIM1\_OE\_WT.1\_A375 (BIOPOLYMER)



GNA13\_OE\_A456C\_A375 (PUJANA\_BRC)

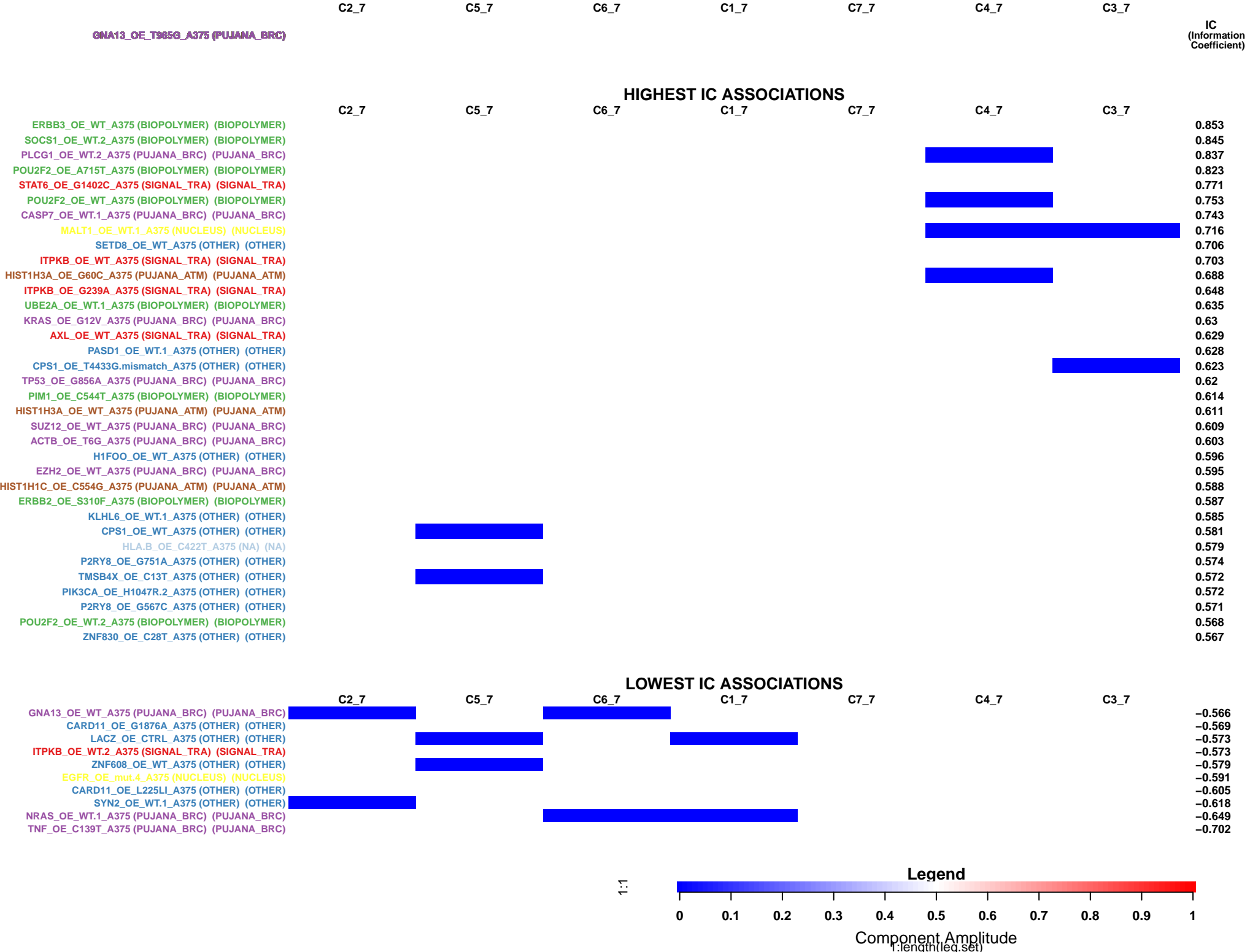


FGF10\_OE\_WT\_A375 (NUCLEUS)

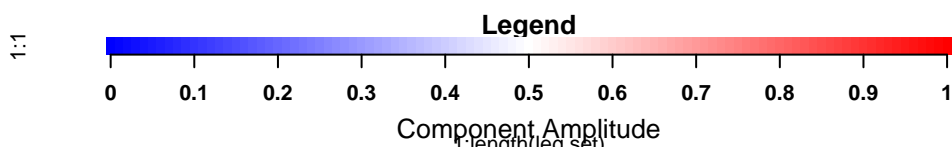
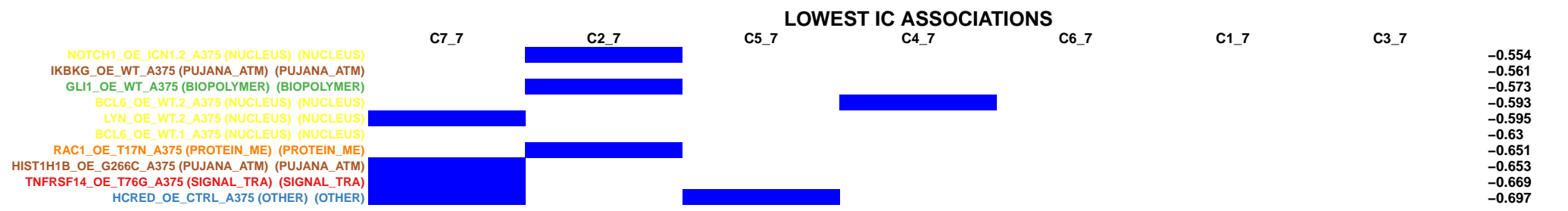
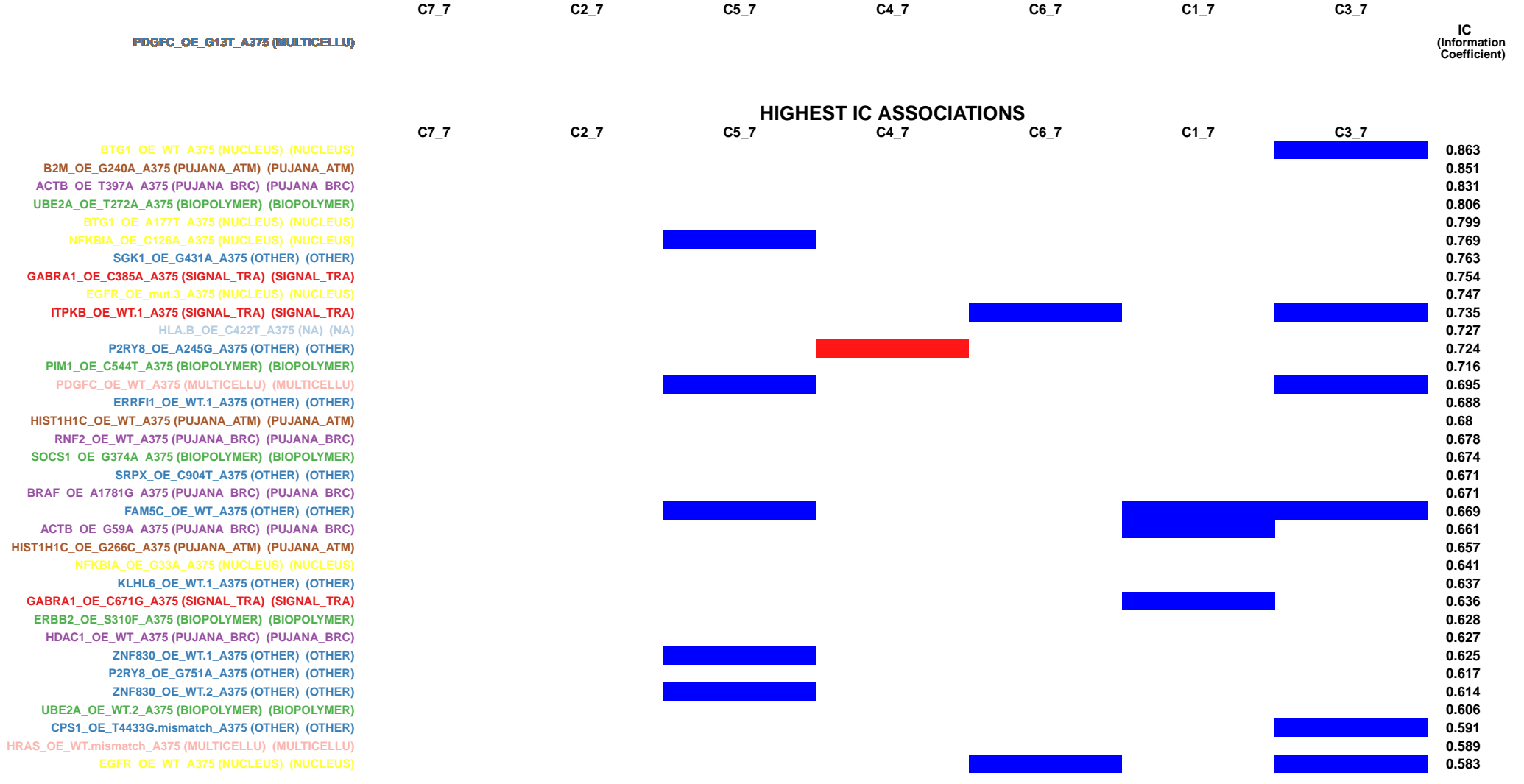




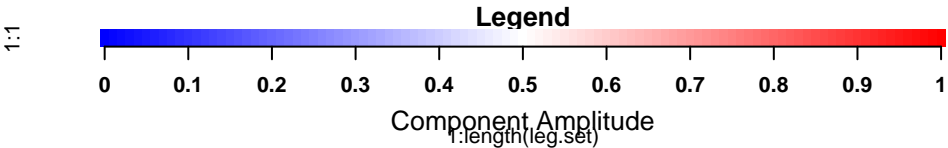
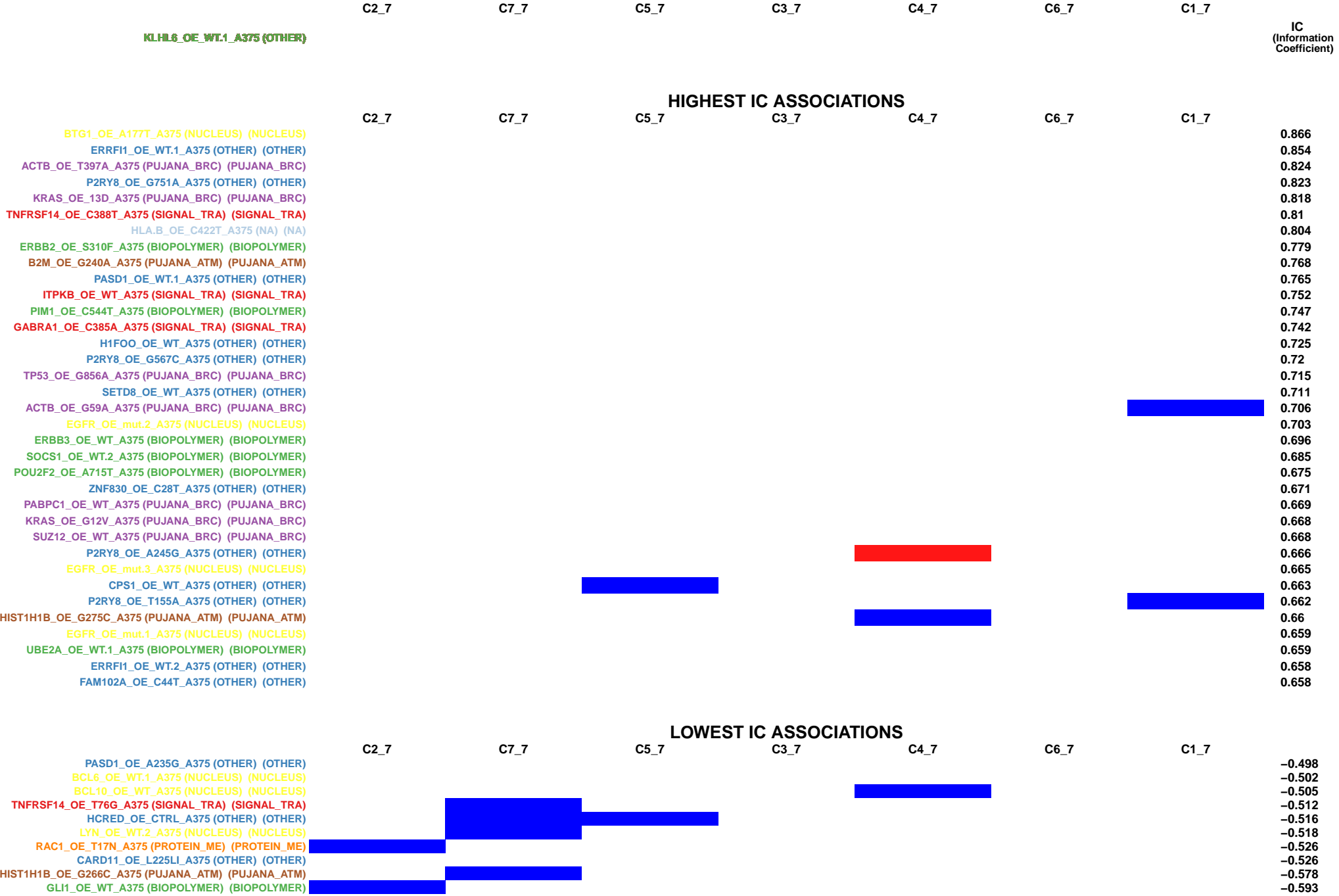
GNA13\_OE\_T965G\_A375 (PUJANA\_BRC)



PDGFC\_OE\_G13T\_A375 (MULTICELLU)

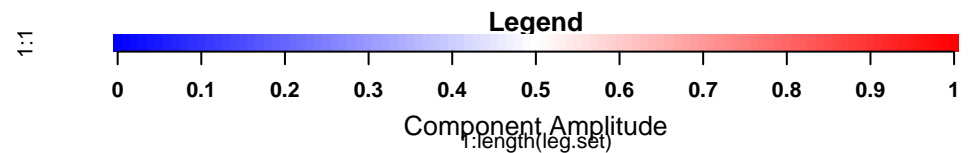


KLHL6\_OE\_WT.1\_A375 (OTHER)

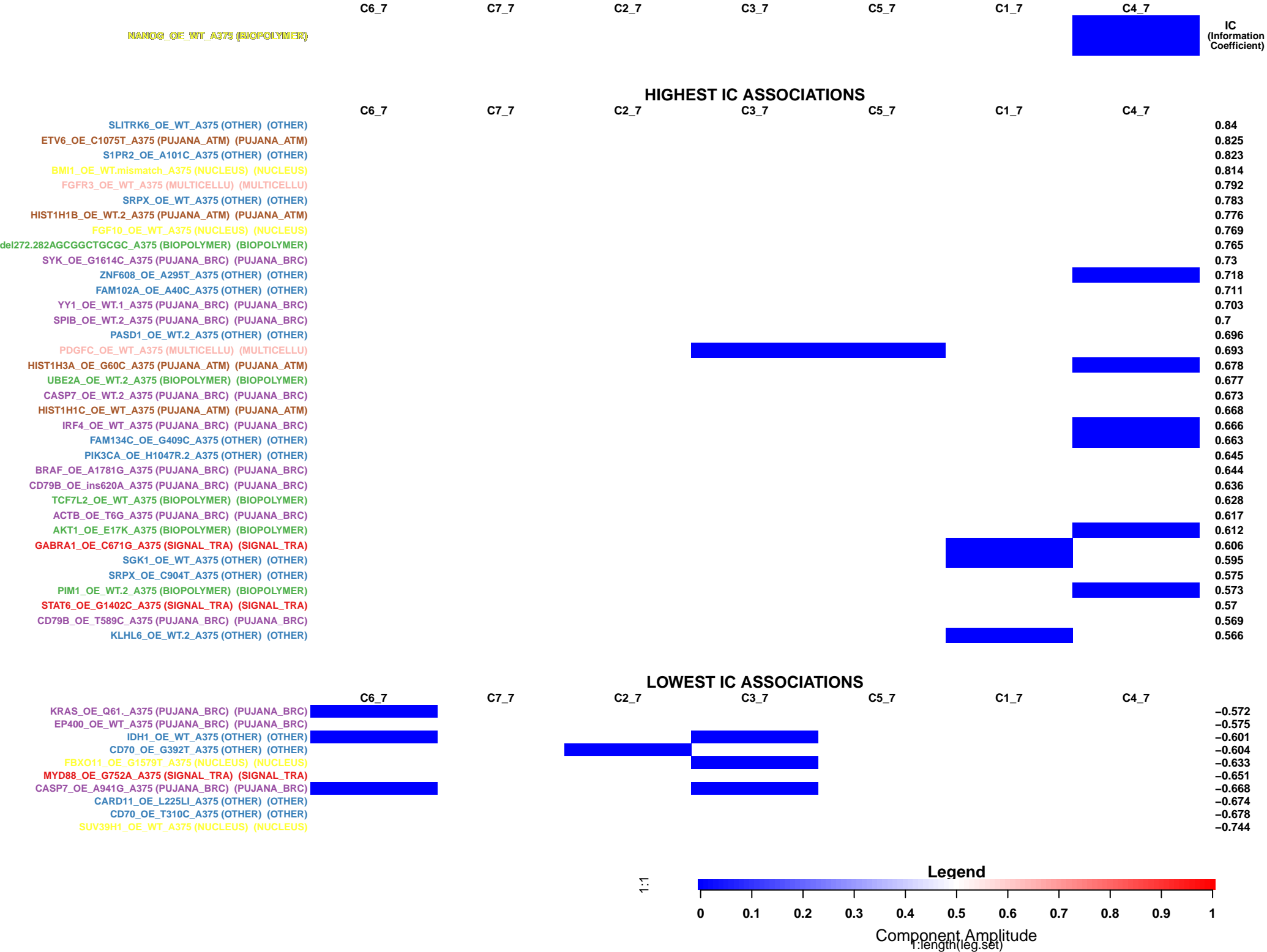


[illegible]

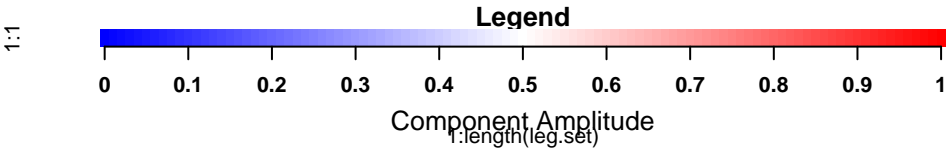
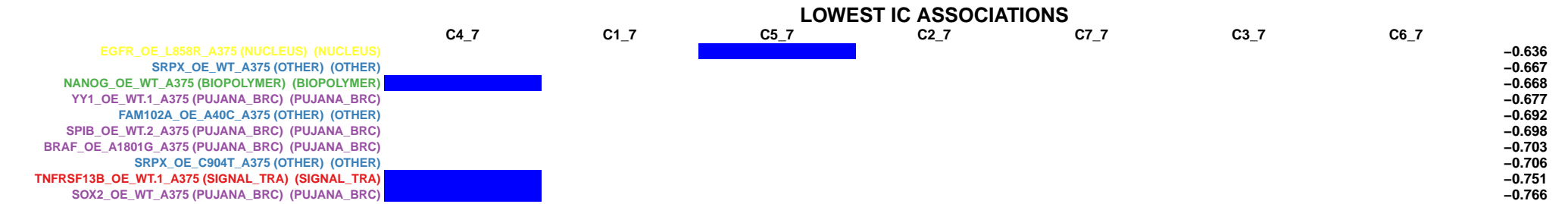
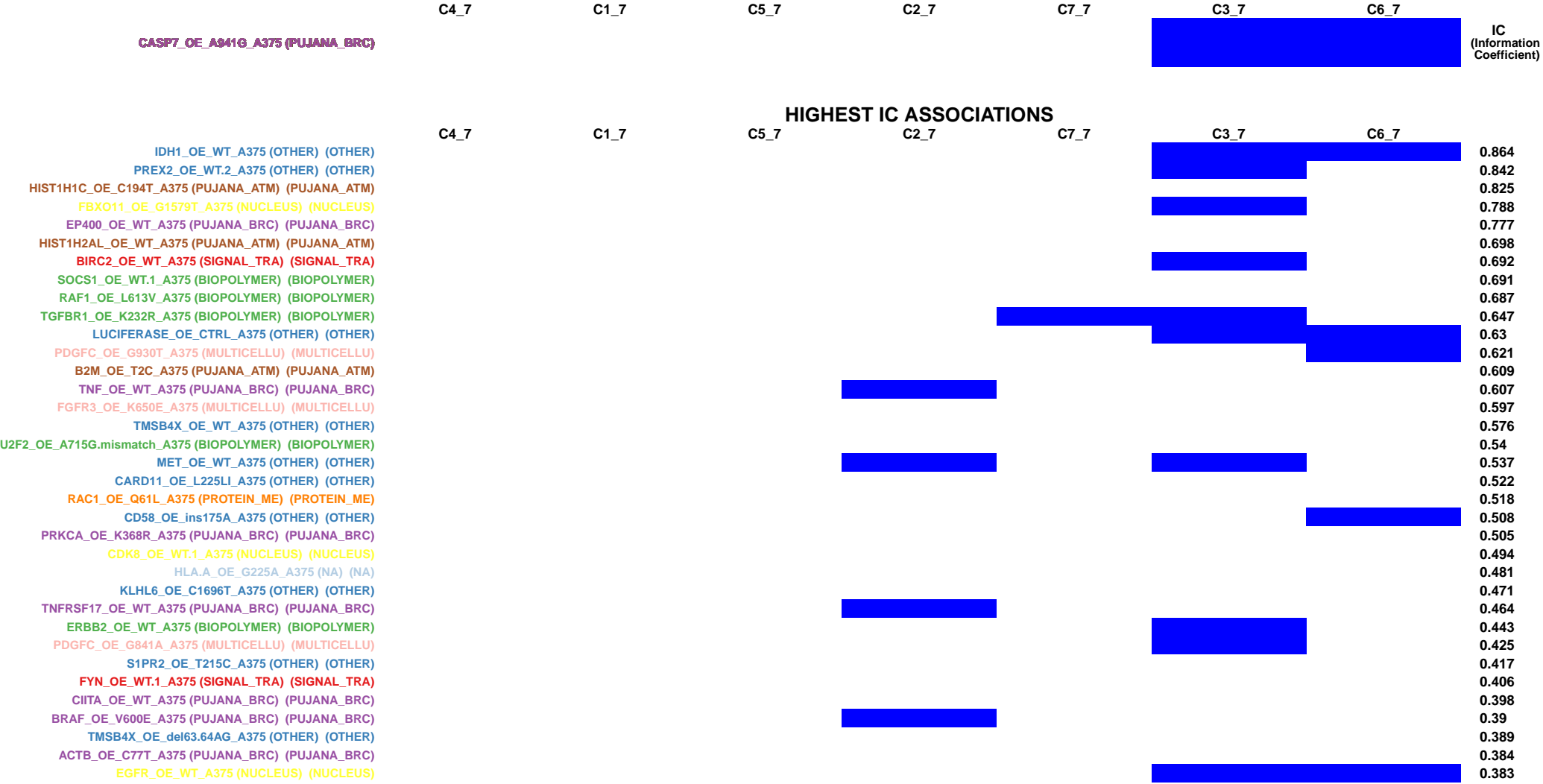
C2_7	C7_7	C1_7	C6_7	C5_7	C3_7	C4_7
						-0.514
						-0.516
						-0.533
						-0.534
						-0.543
						-0.564
						-0.608
						-0.64
						-0.744
						-0.794



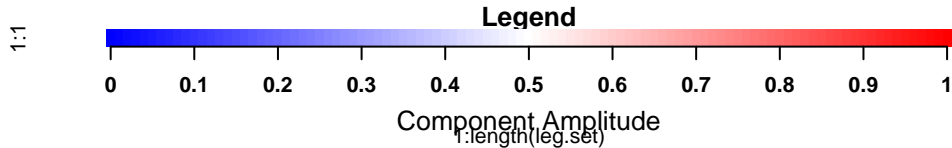
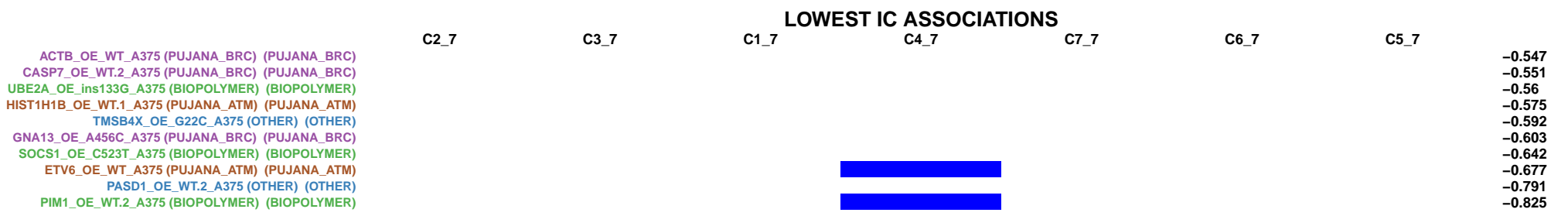
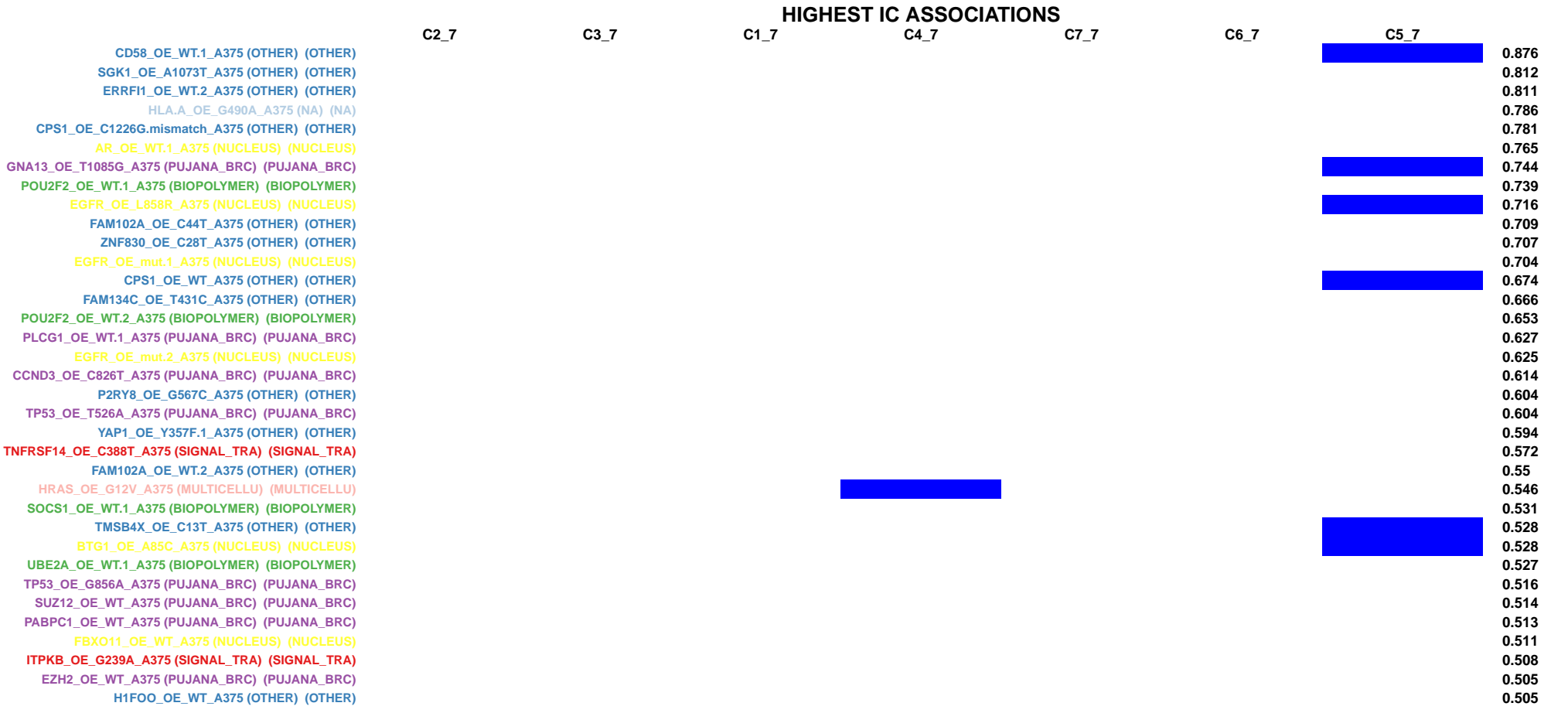
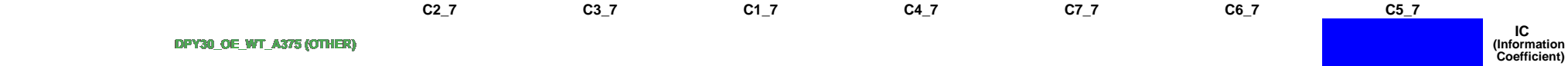
NANOG\_OE\_WT\_A375 (BIOPOLYMER)



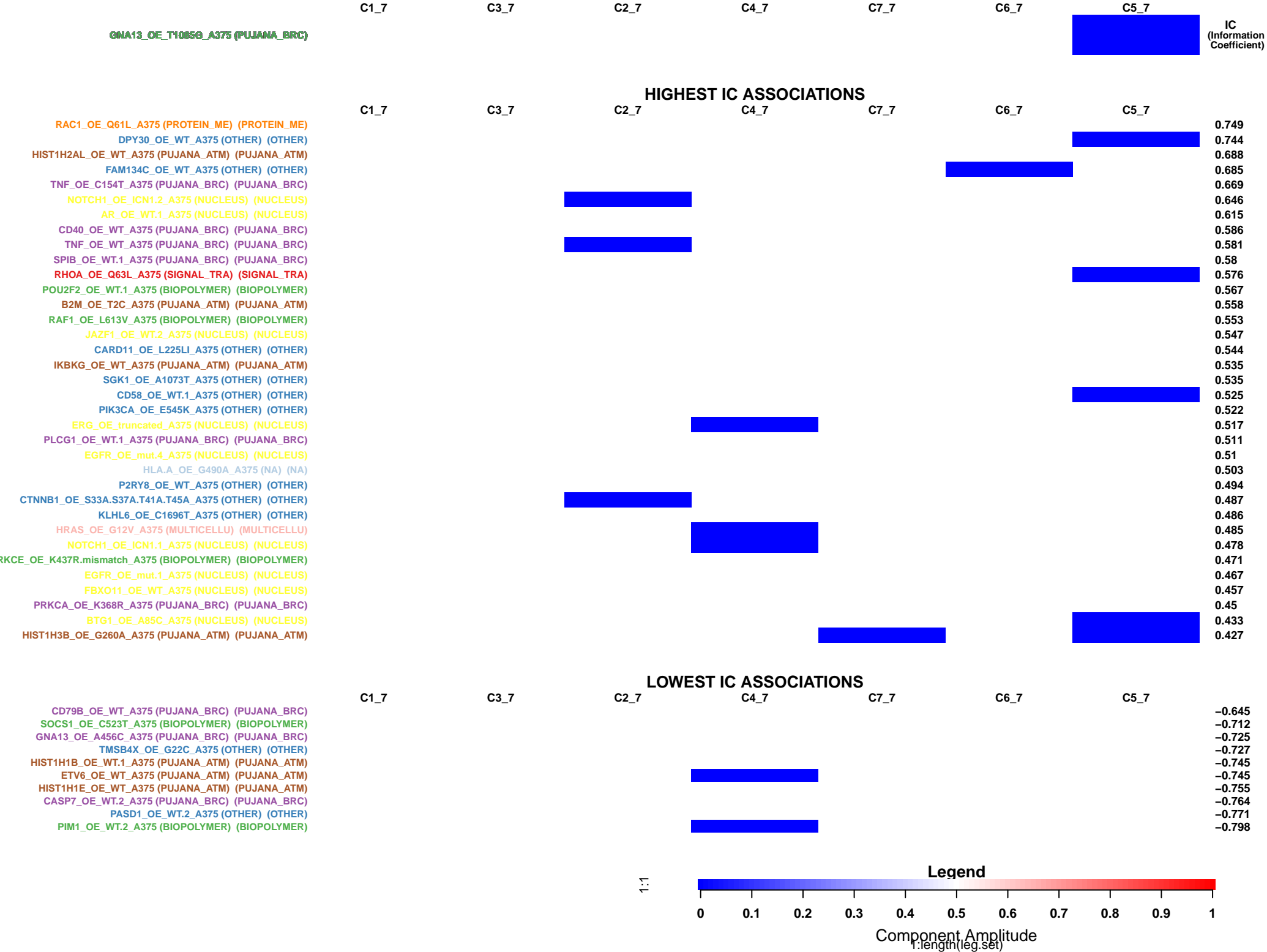
CASP7\_OE\_A941G\_A375 (PUJANA\_BRC)



DPY30\_OE\_WT\_A375 (OTHER)



GNA13\_OE\_T1085G\_A375 (PUJANA\_BRC)





SETD8\_OE\_WT\_A375 (OTHER)

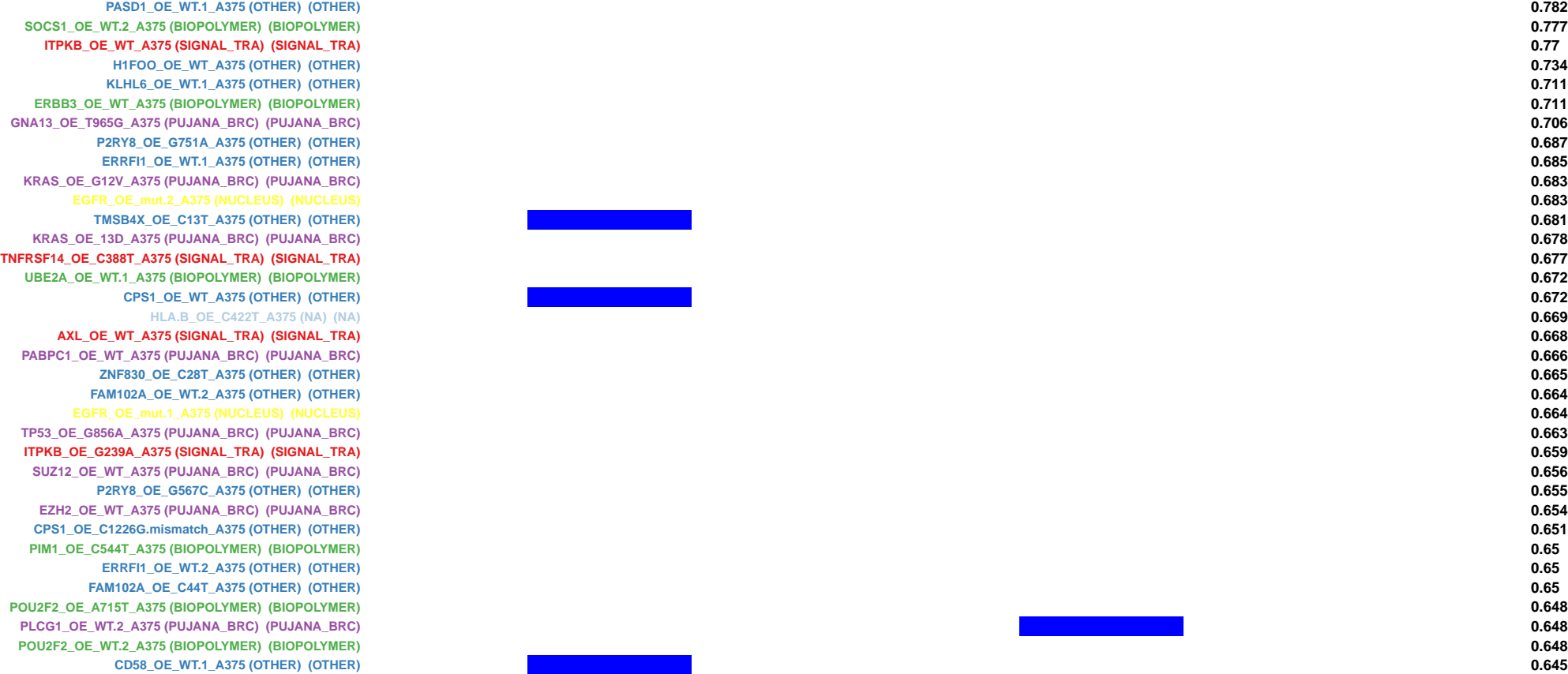
C2\_7C5\_7C3\_7C6\_7C4\_7C7\_7C1\_7

SETD8\_OE\_WT\_A375 (OTHER)

IC  
(Information  
Coefficient)

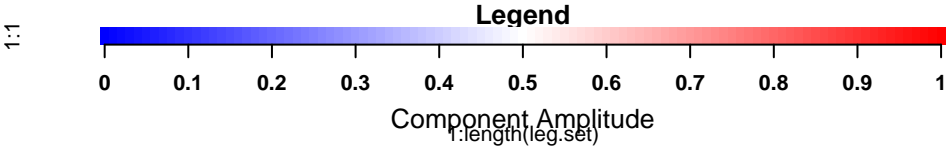
HIGHEST IC ASSOCIATIONS

C2\_7C5\_7C3\_7C6\_7C4\_7C7\_7C1\_7

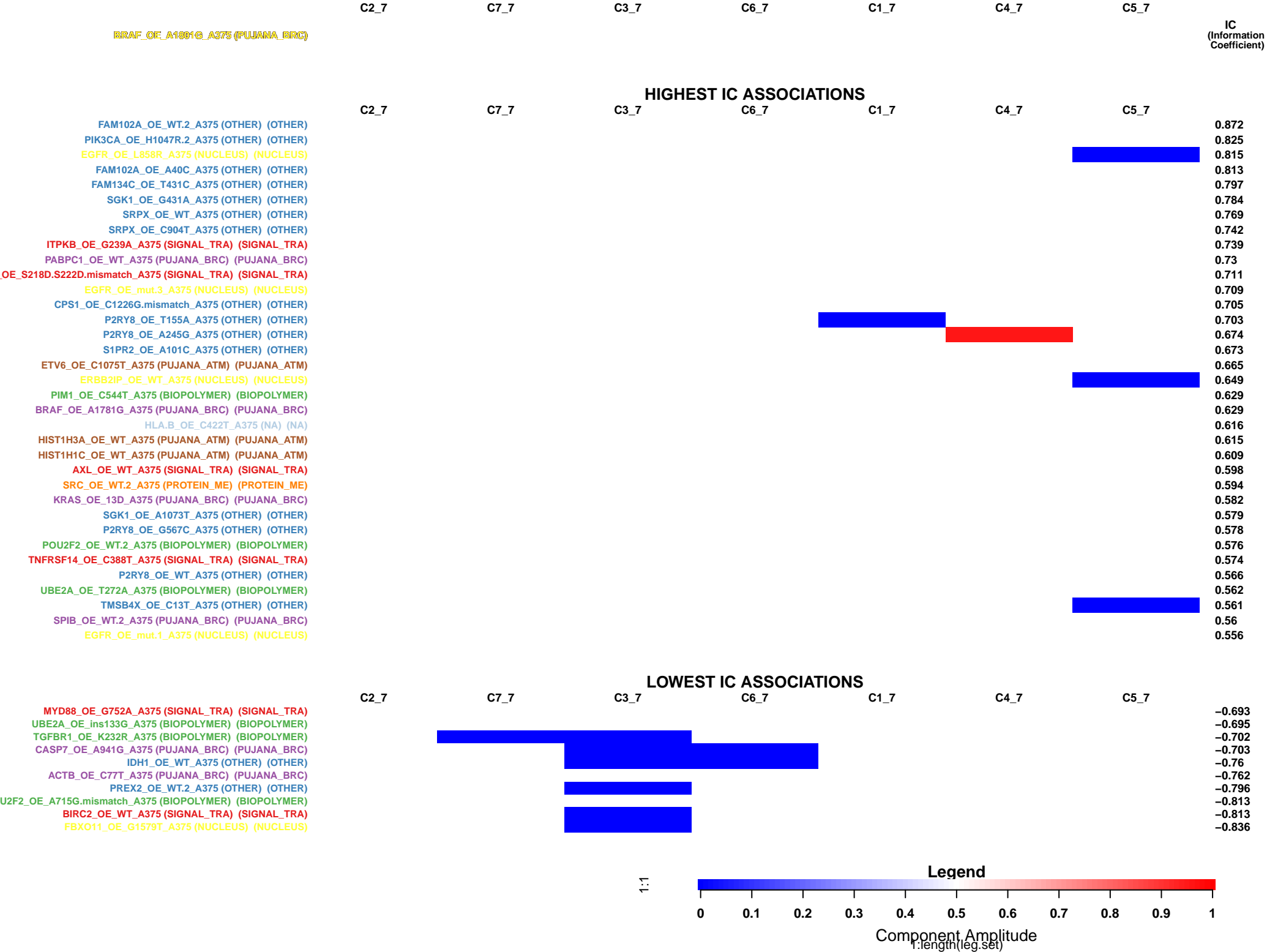


LOWEST IC ASSOCIATIONS

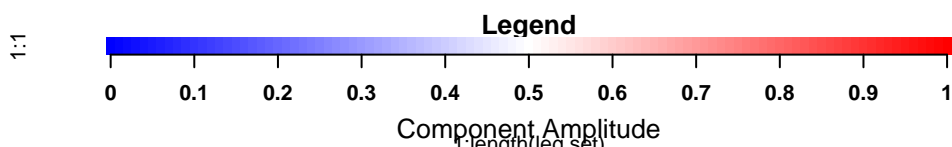
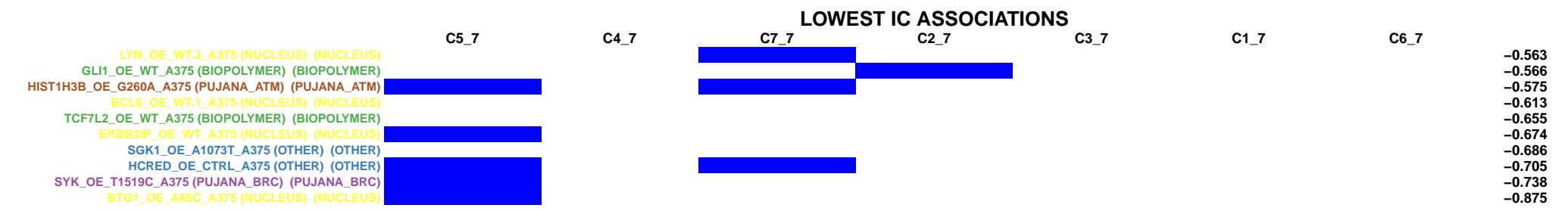
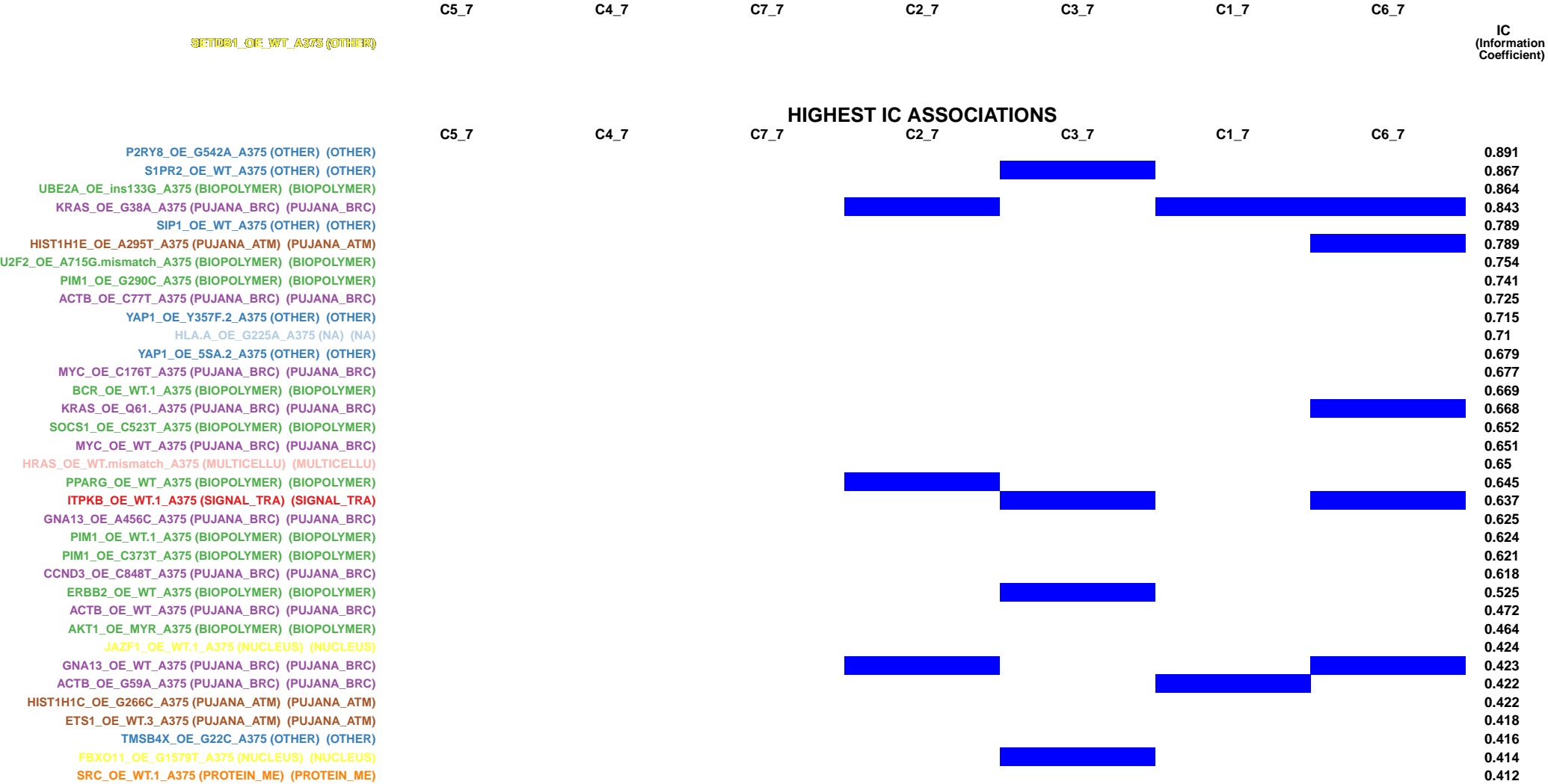
C2\_7C5\_7C3\_7C6\_7C4\_7C7\_7C1\_7



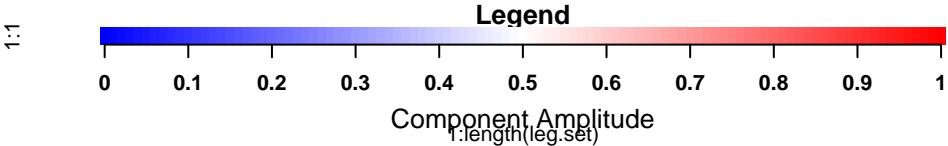
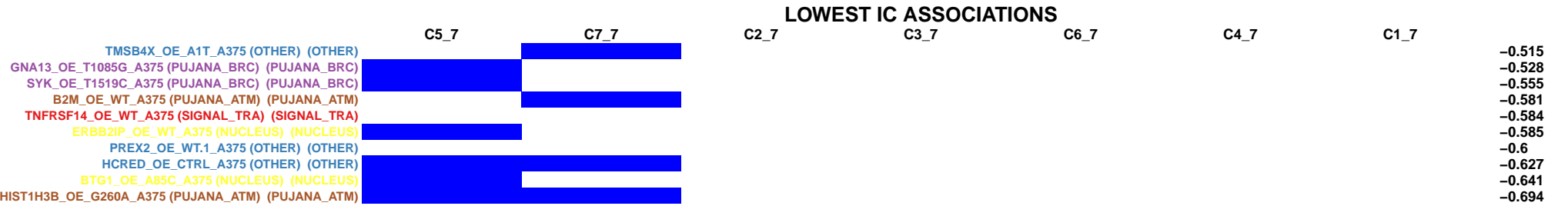
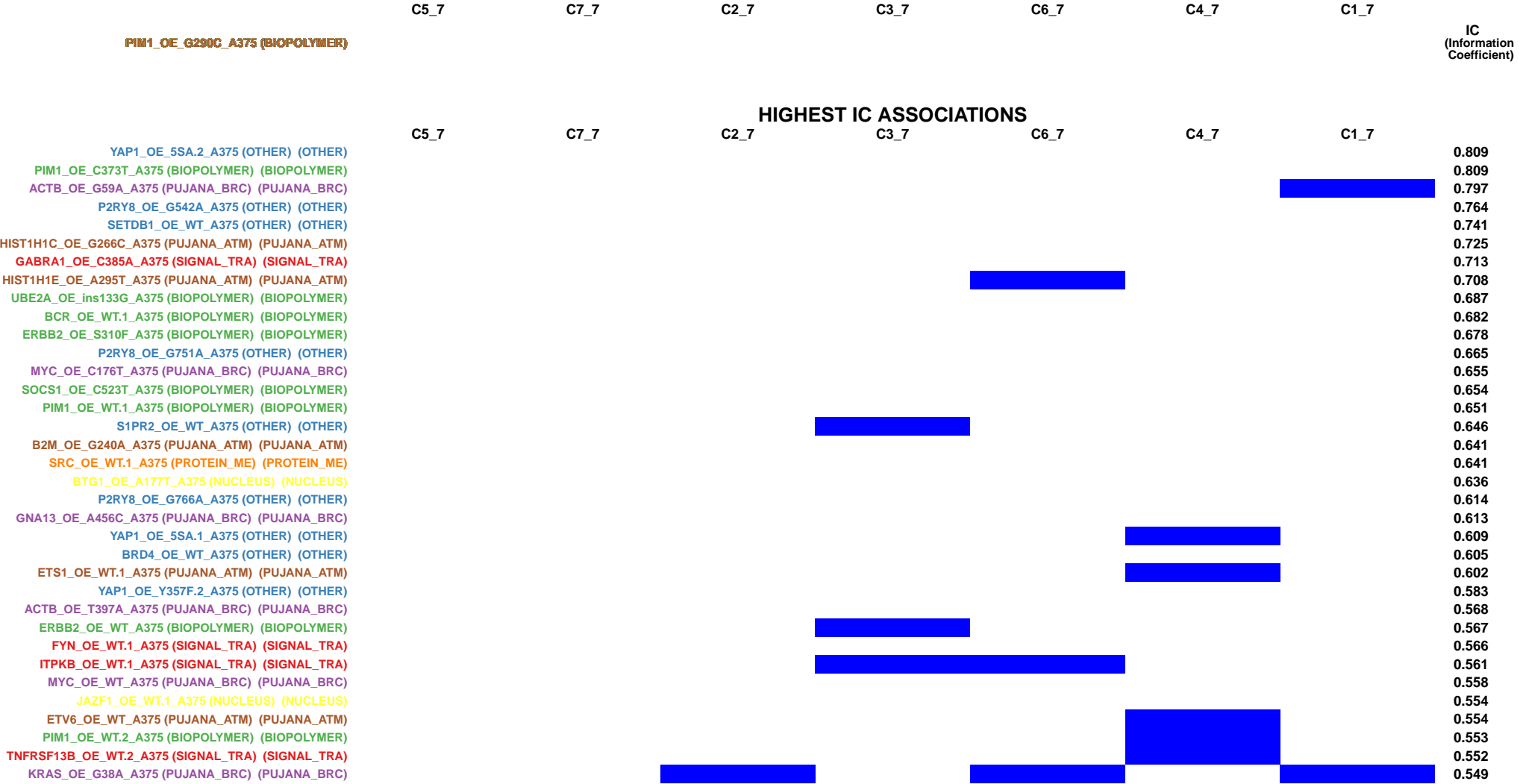
BRAF\_OE\_A1801G\_A375 (PUJANA\_BRC)

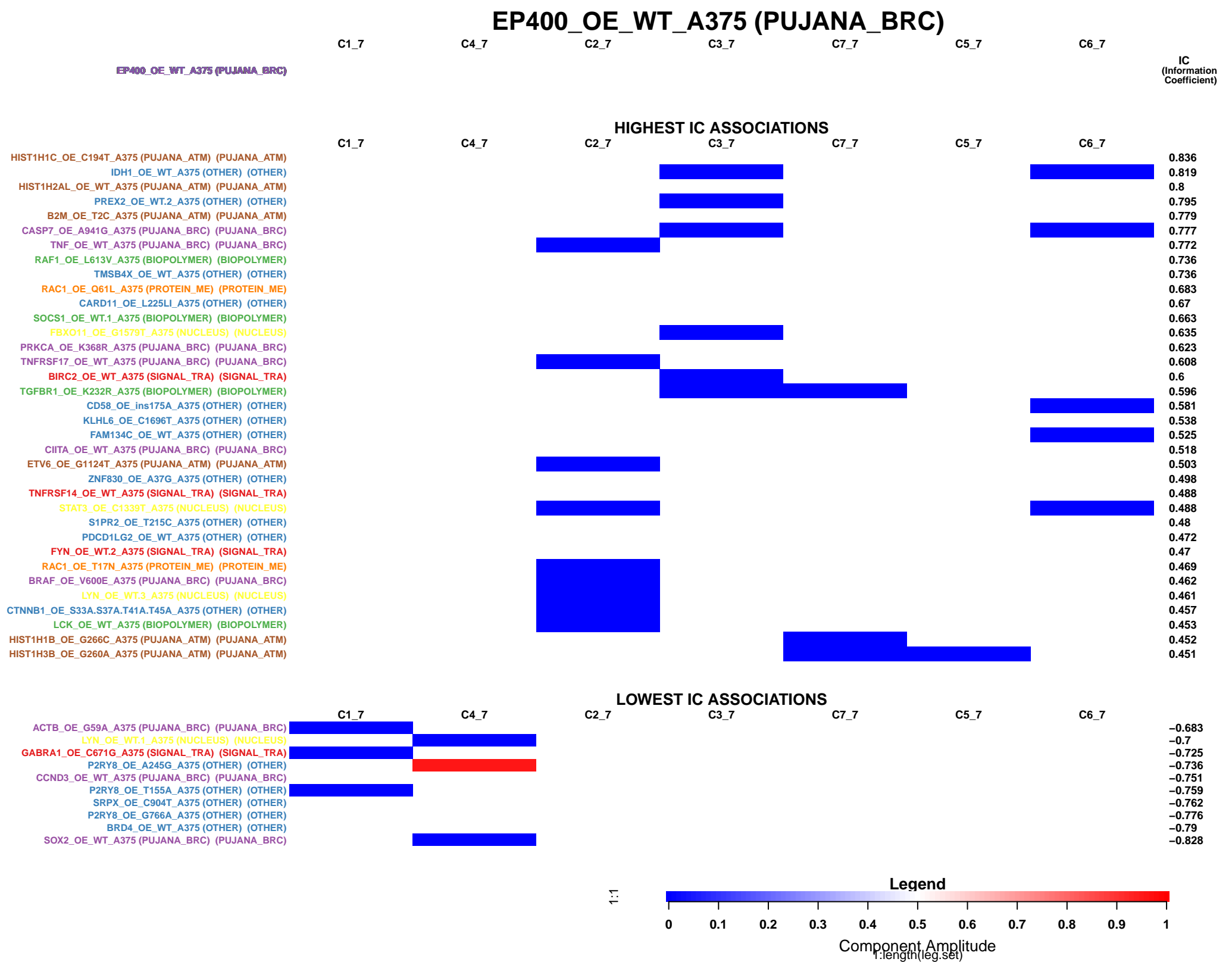


SETDB1\_OE\_WT\_A375 (OTHER)

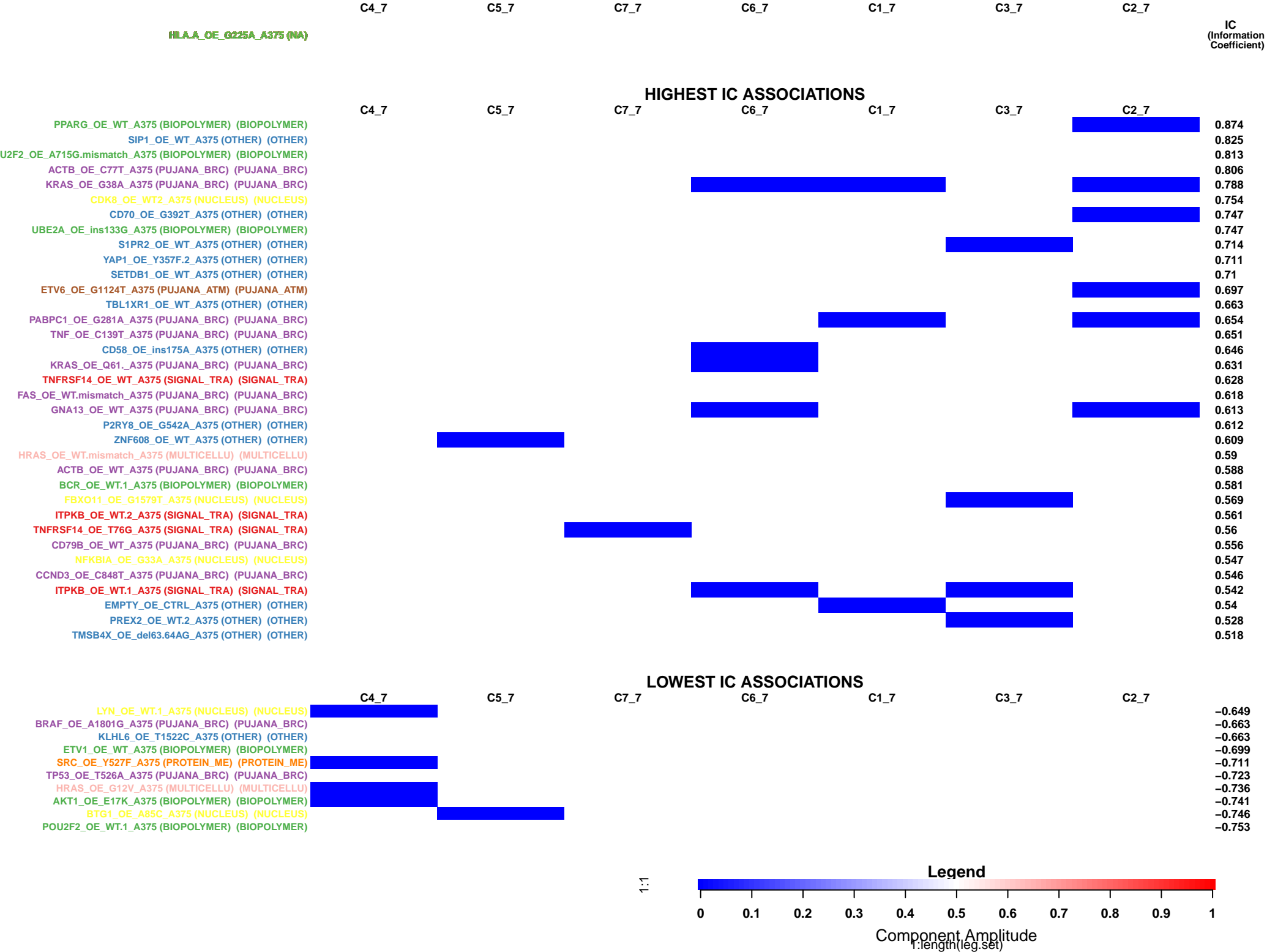


PIM1\_OE\_G290C\_A375 (BIOPOLYMER)





HLA.A\_OE\_G225A\_A375 (NA)

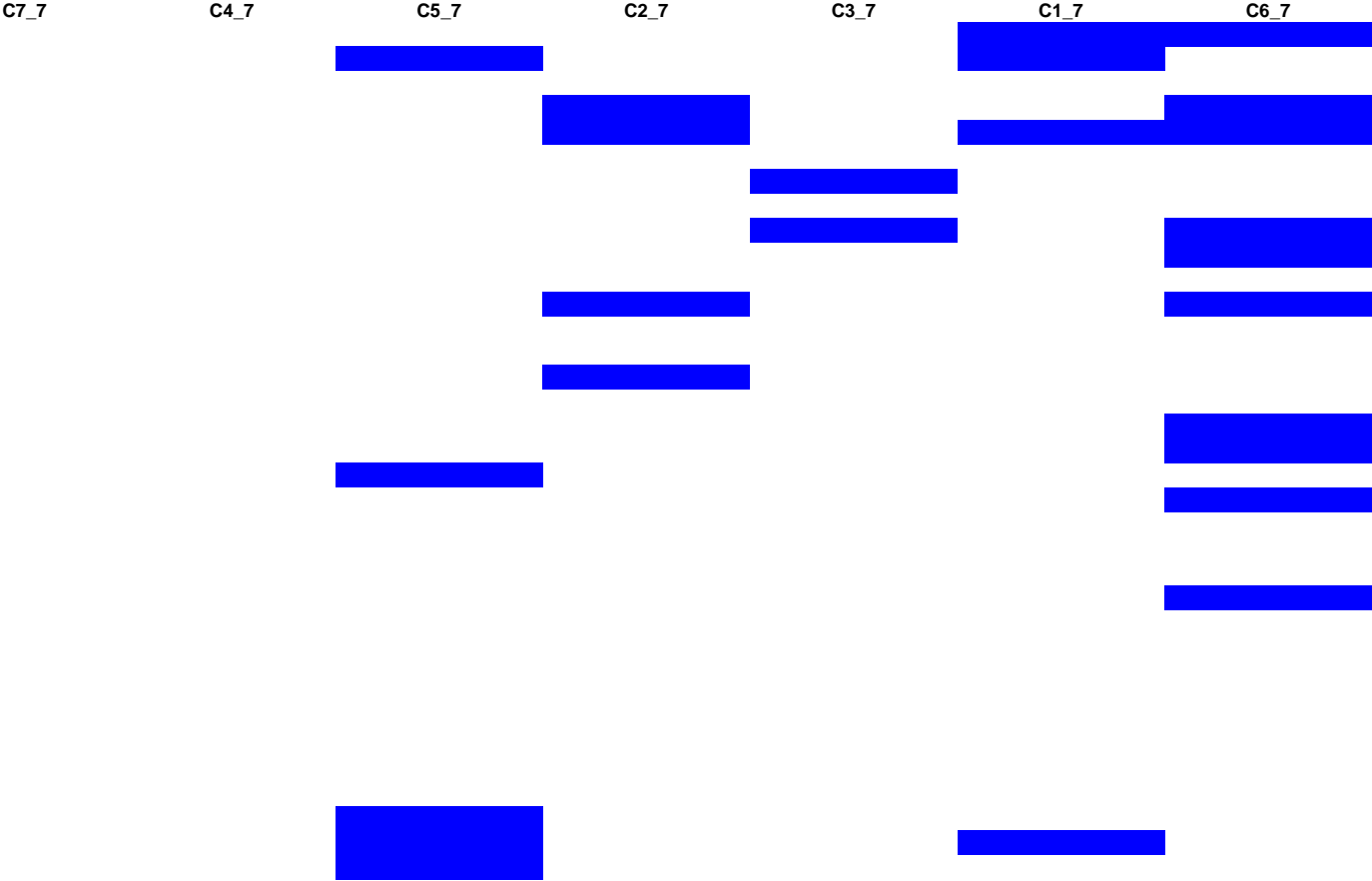


HRAS\_OE\_WT.mismatch\_A375 (MULTICELLU)

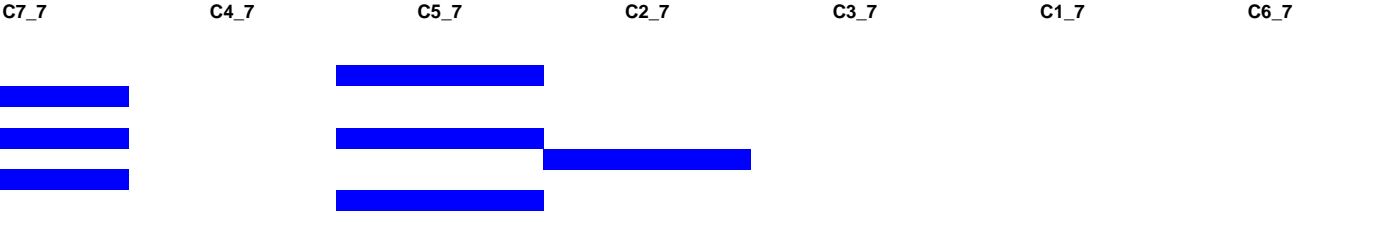
C7\_7C4\_7C5\_7C2\_7C3\_7C1\_7C6\_7

IC  
(Information  
Coefficient)

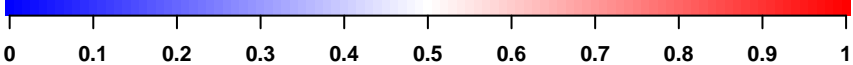
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS



Legend

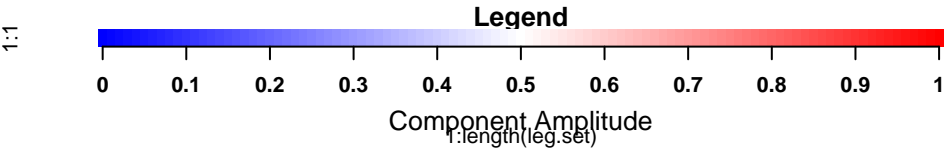
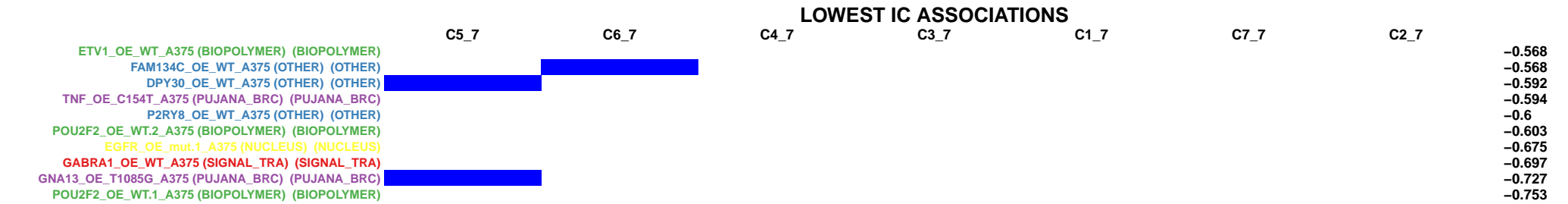
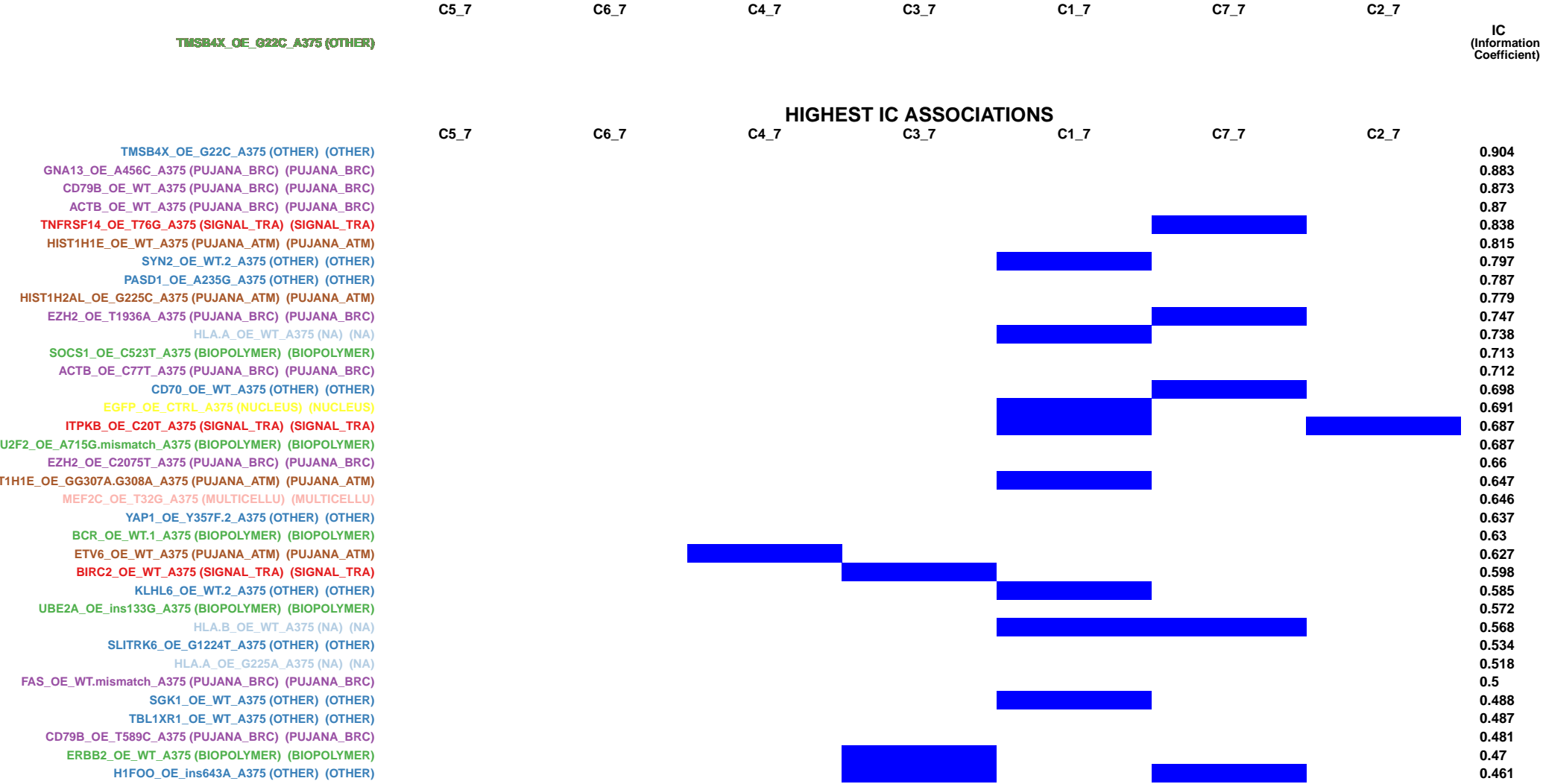


Component Amplitude

1:1

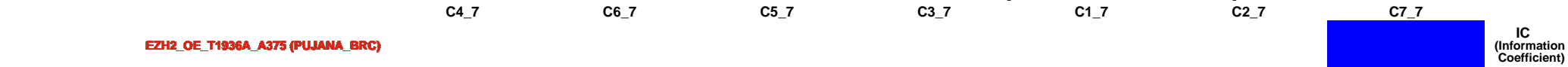
1:length(leg.set)

TMSB4X\_OE\_G22C\_A375 (OTHER)

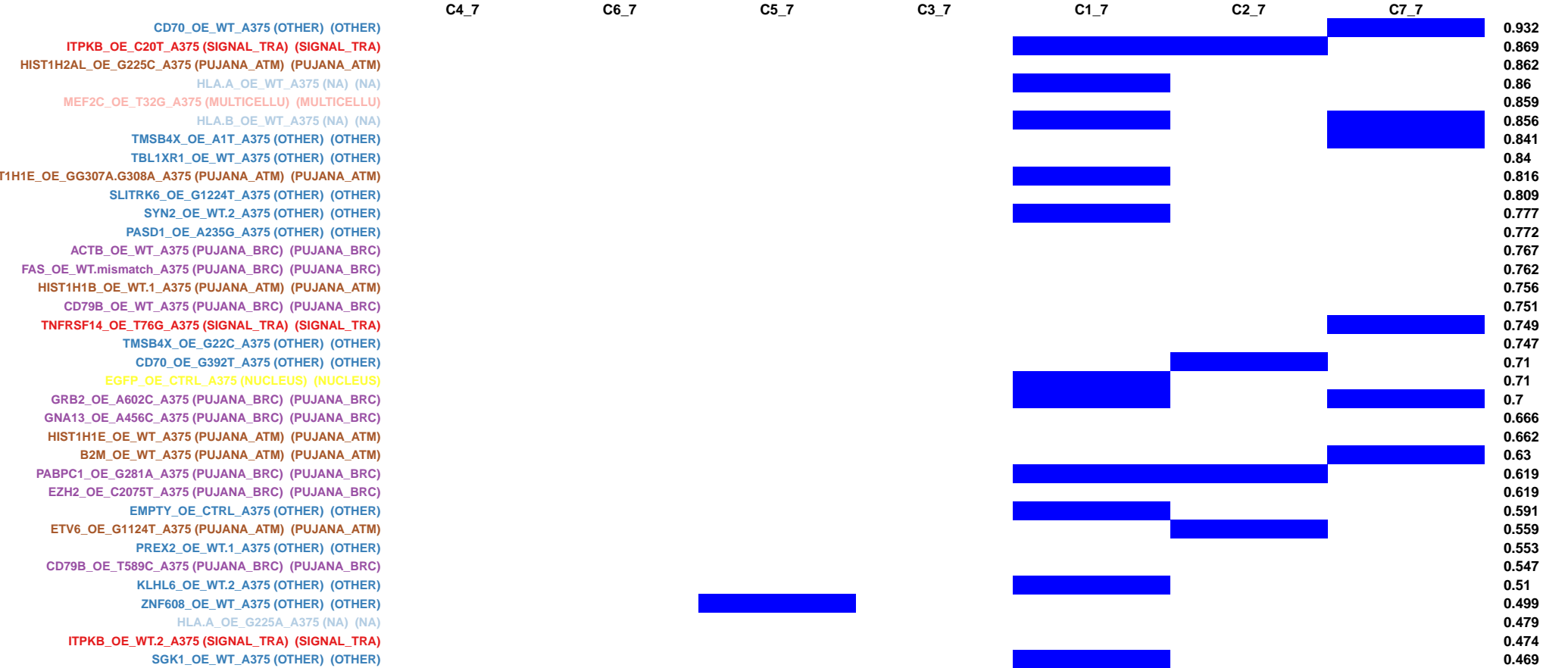




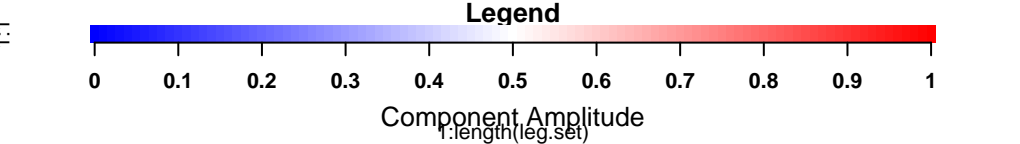
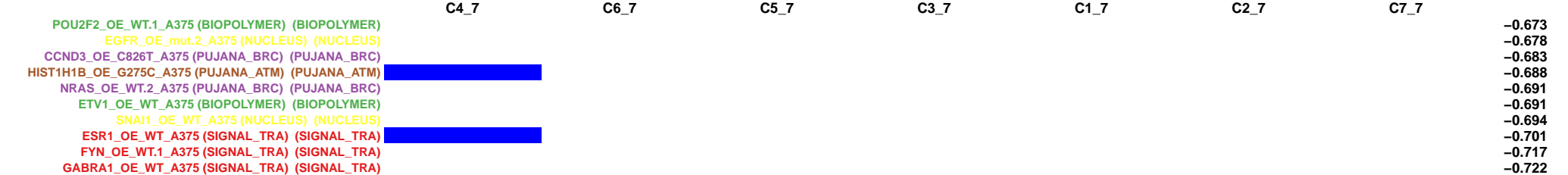
EZH2\_OE\_T1936A\_A375 (PUJANA\_BRC)



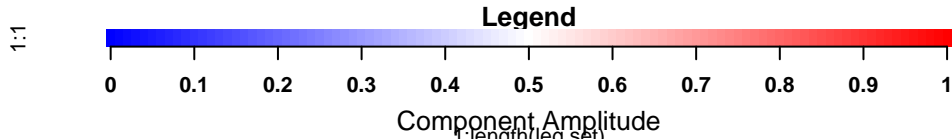
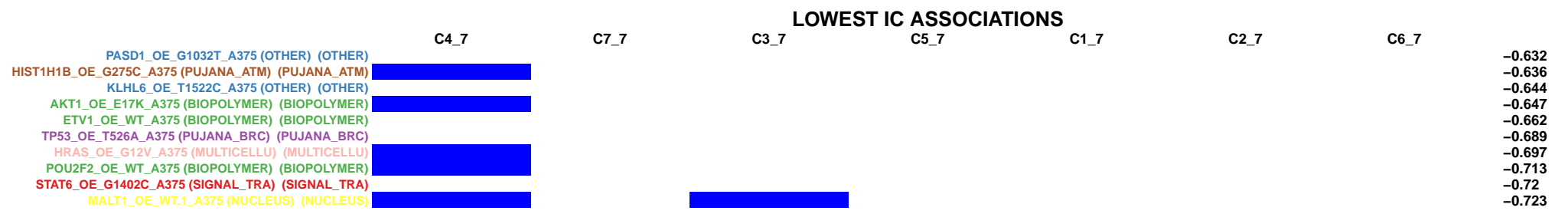
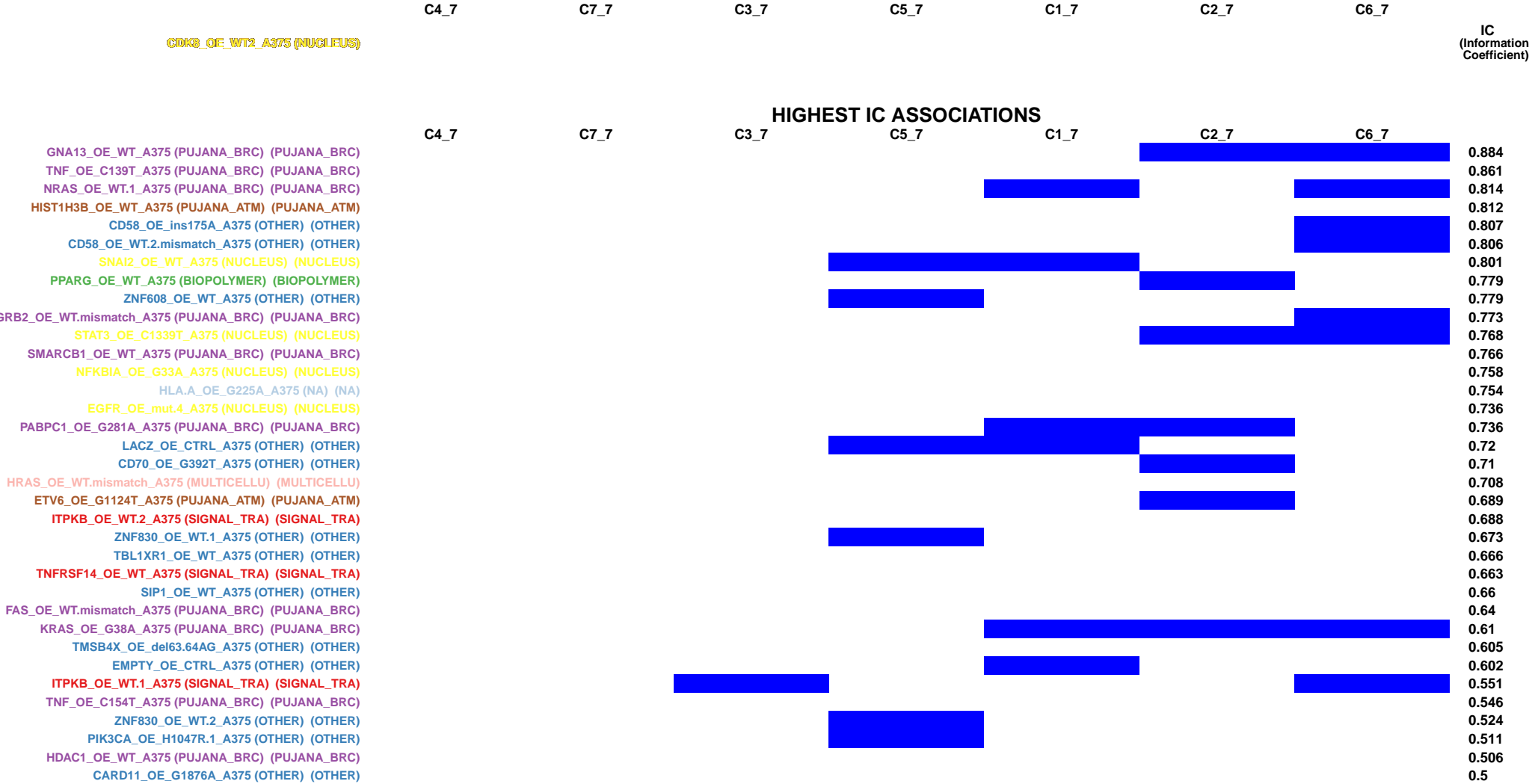
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS



CDK8\_OE\_WT2\_A375 (NUCLEUS)



FAM102A\_OE\_C44T\_A375 (OTHER)

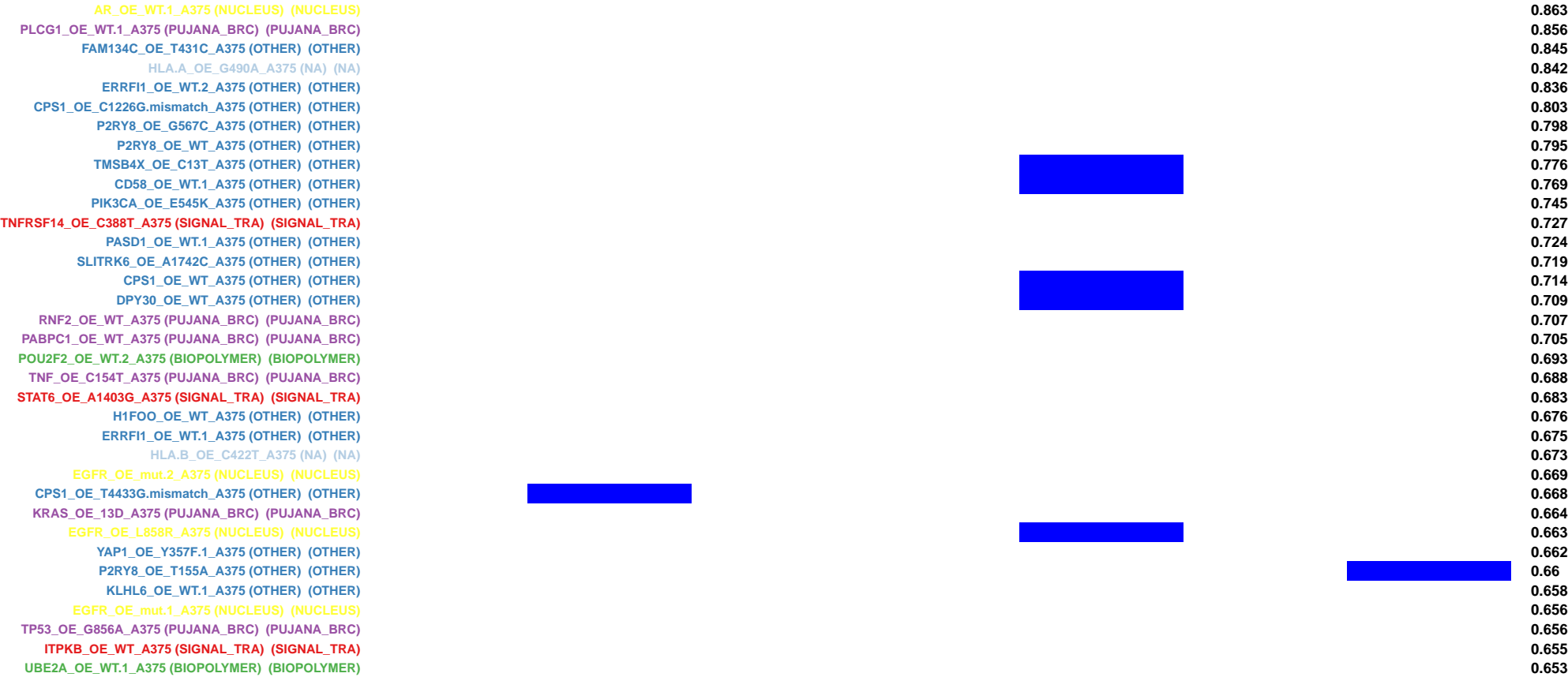
FAM102A\_OE\_C44T\_A375 (OTHER)

IC  
(Information  
Coefficient)

C2\_7C3\_7C4\_7C7\_7C5\_7C6\_7C1\_7

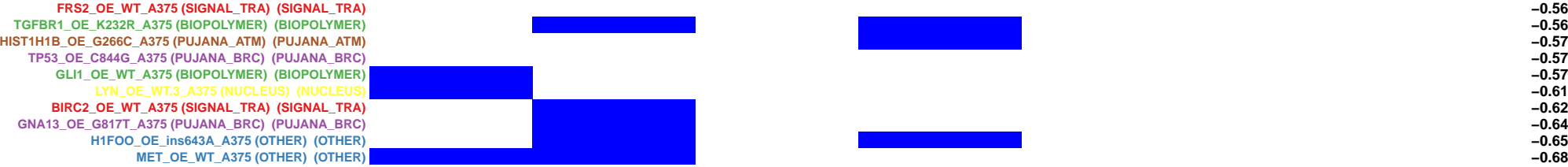
HIGHEST IC ASSOCIATIONS

C2\_7C3\_7C4\_7C7\_7C5\_7C6\_7C1\_7

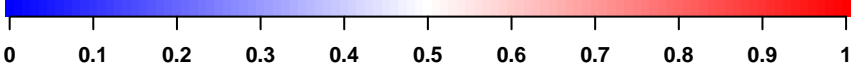


LOWEST IC ASSOCIATIONS

C2\_7C3\_7C4\_7C7\_7C5\_7C6\_7C1\_7



Legend

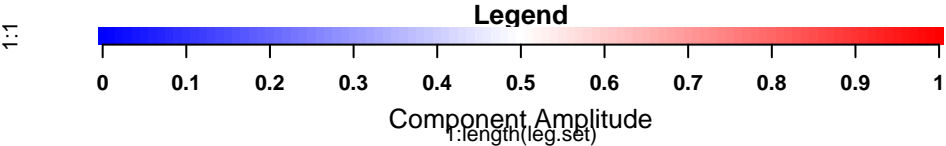
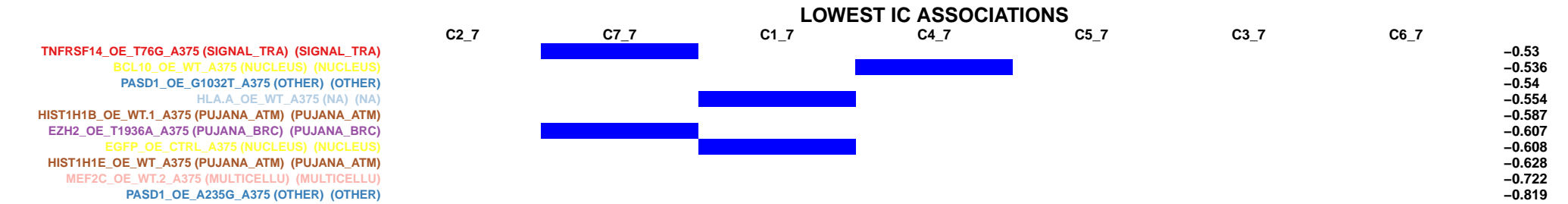
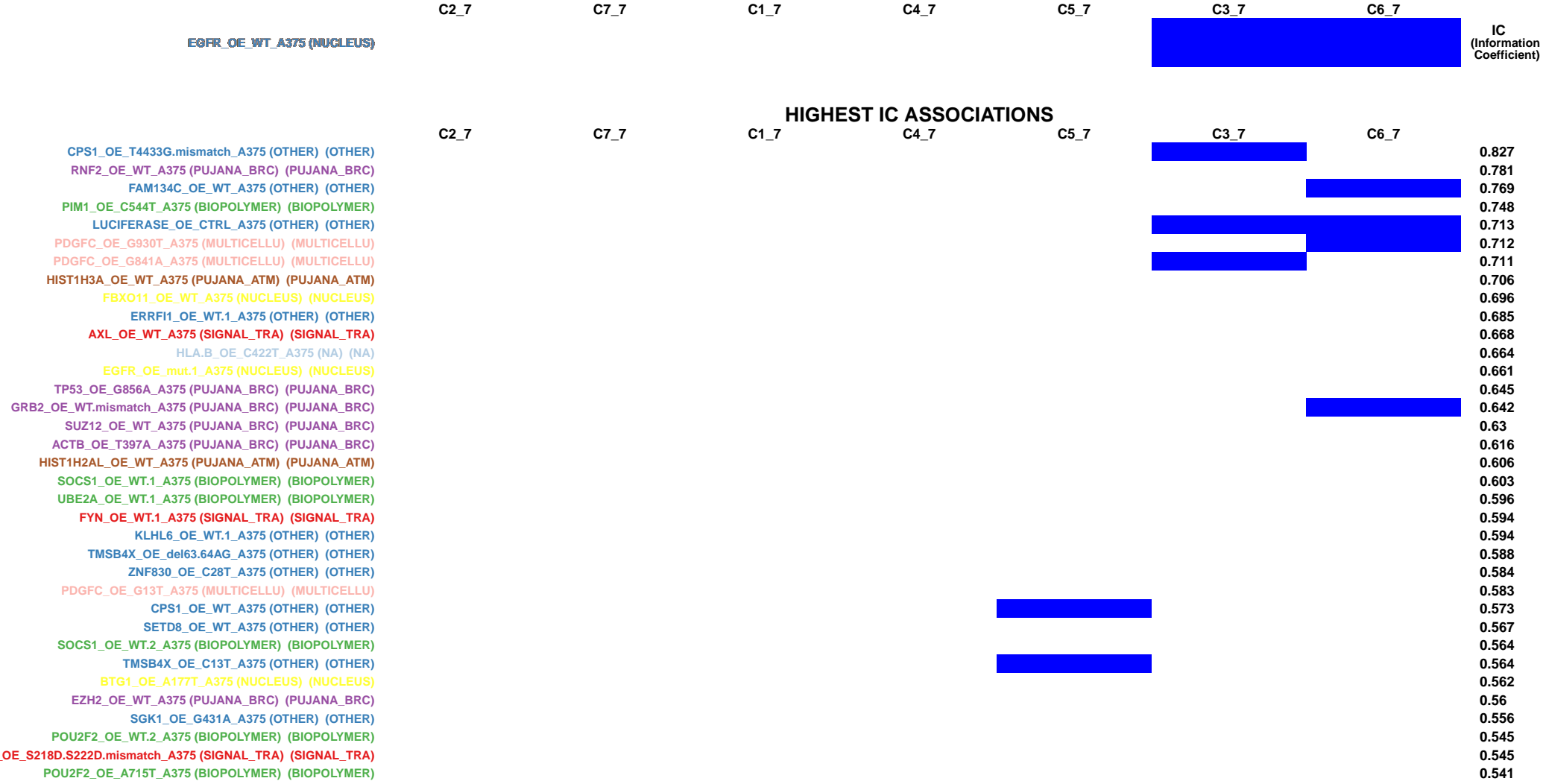


Component Amplitude

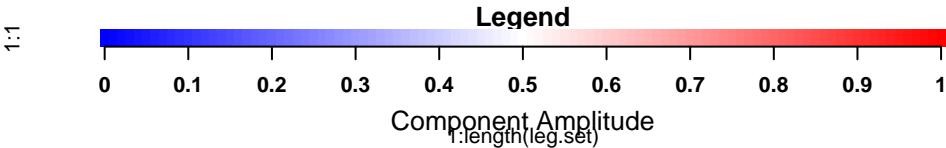
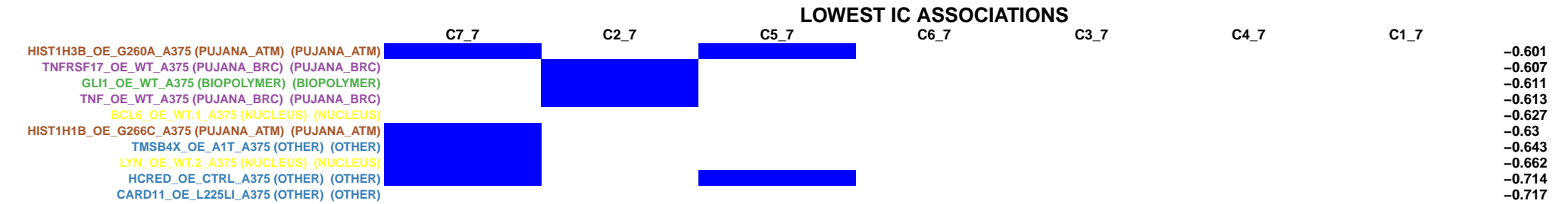
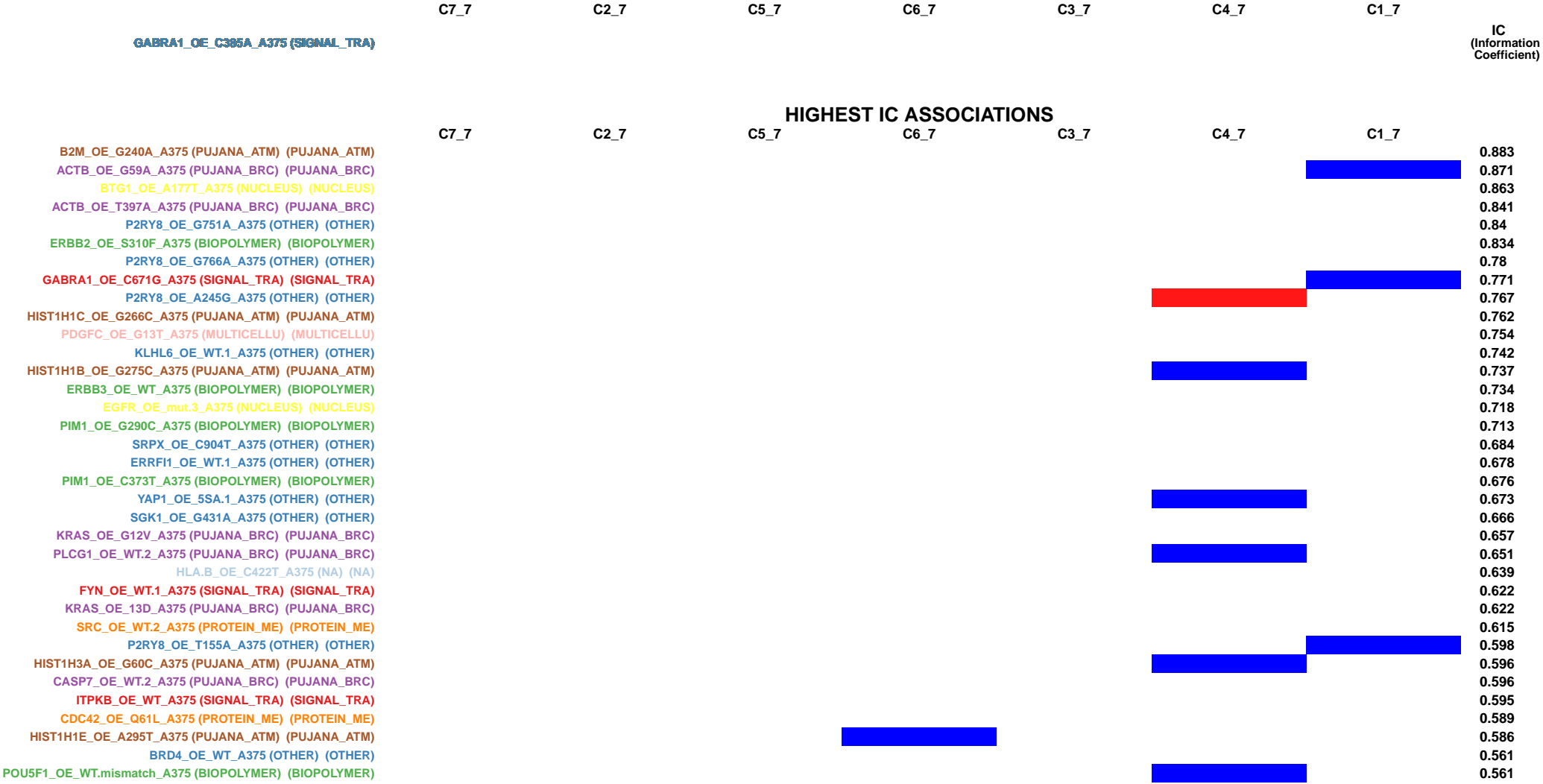
1:1

1:length(leg.set)

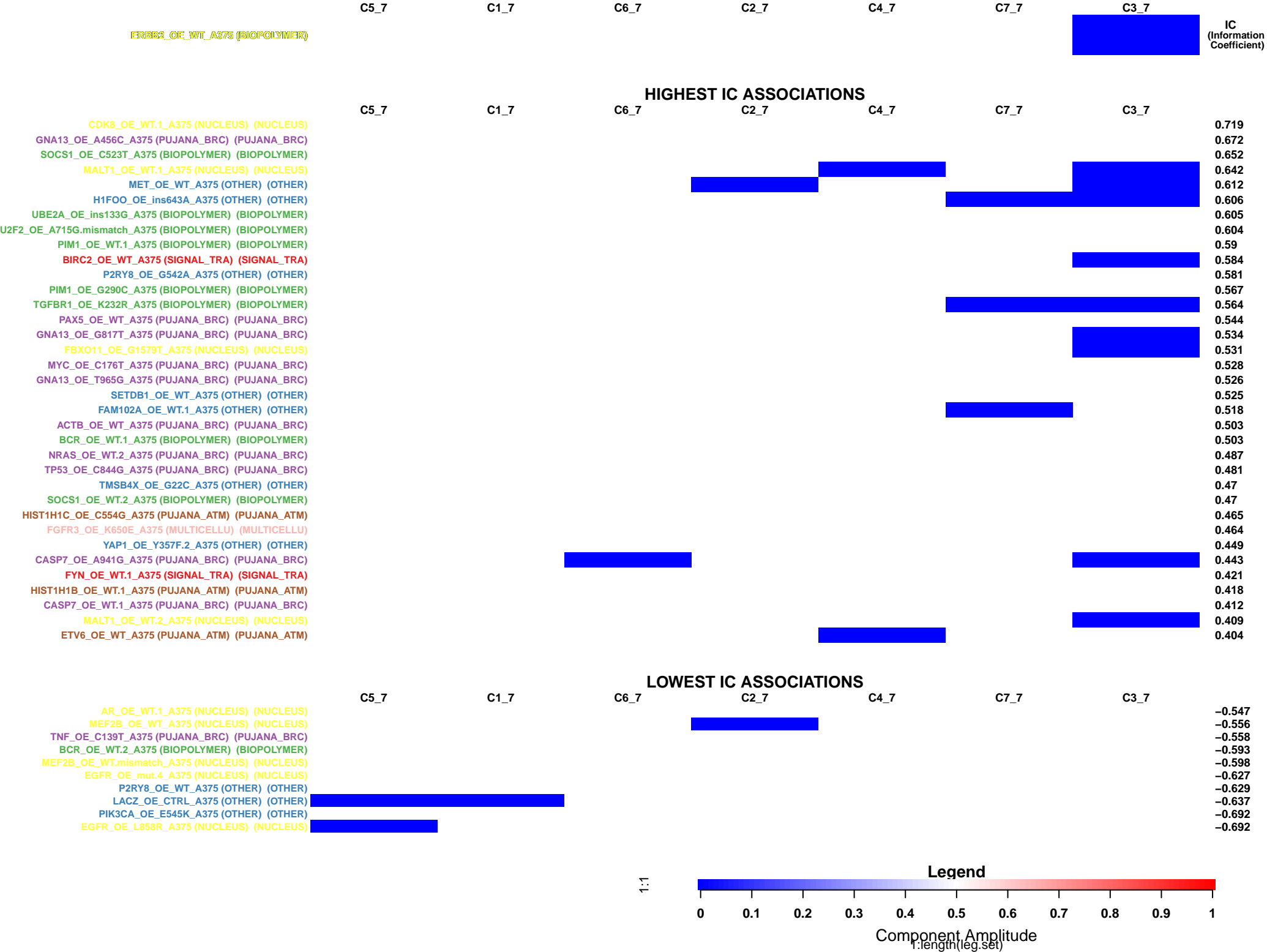
EGFR\_OE\_WT\_A375 (NUCLEUS)



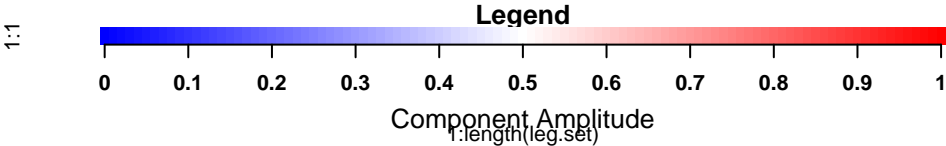
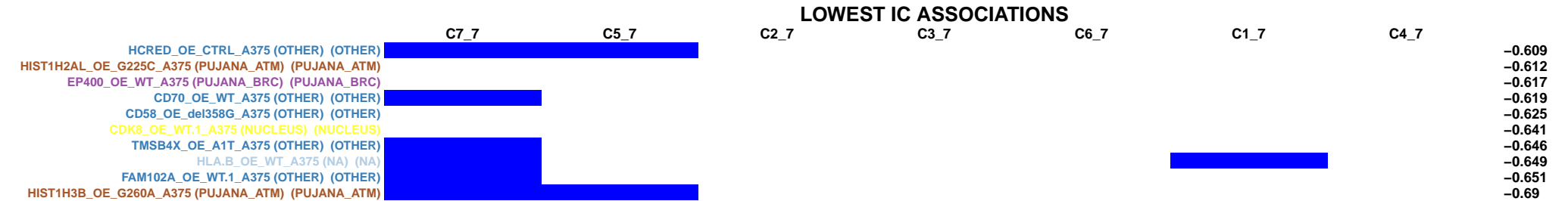
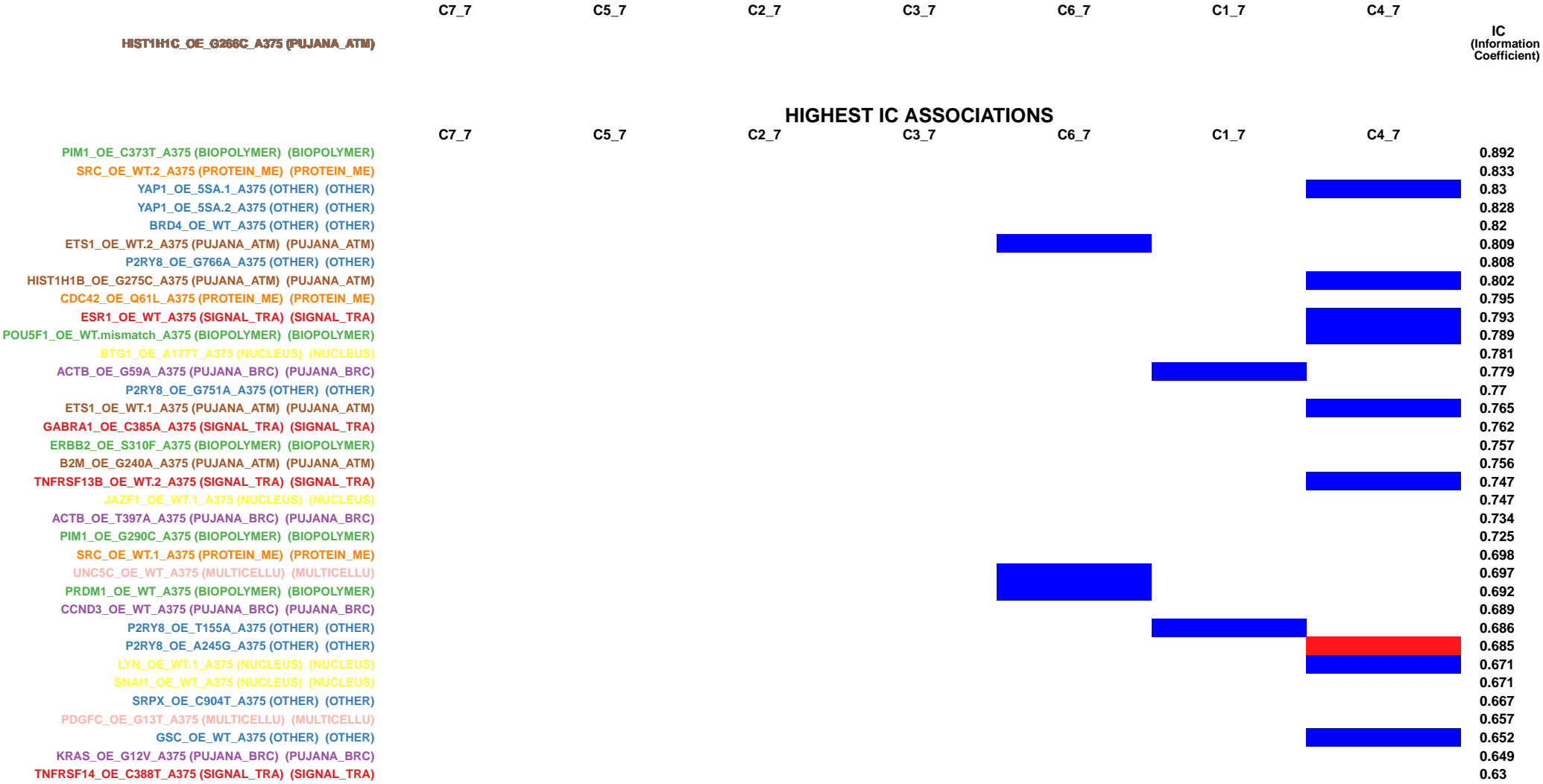
GABRA1\_OE\_C385A\_A375 (SIGNAL\_TRA)



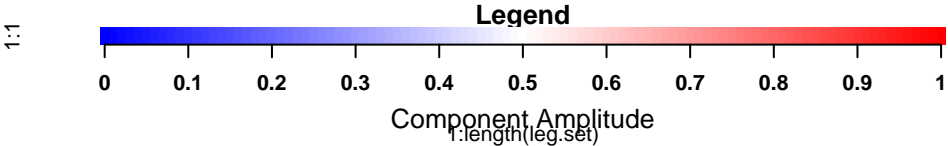
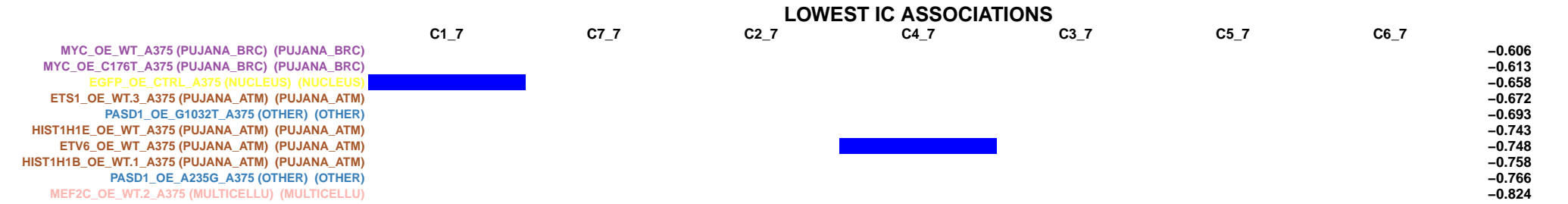
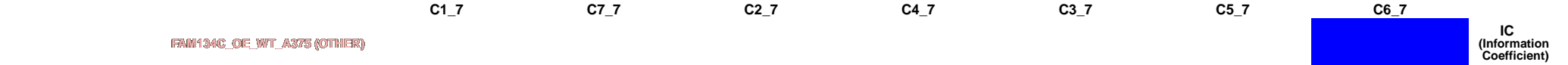
ERBB2\_OE\_WT\_A375 (BIOPOLYMER)



HIST1H1C\_OE\_G266C\_A375 (PUJANA\_ATM)

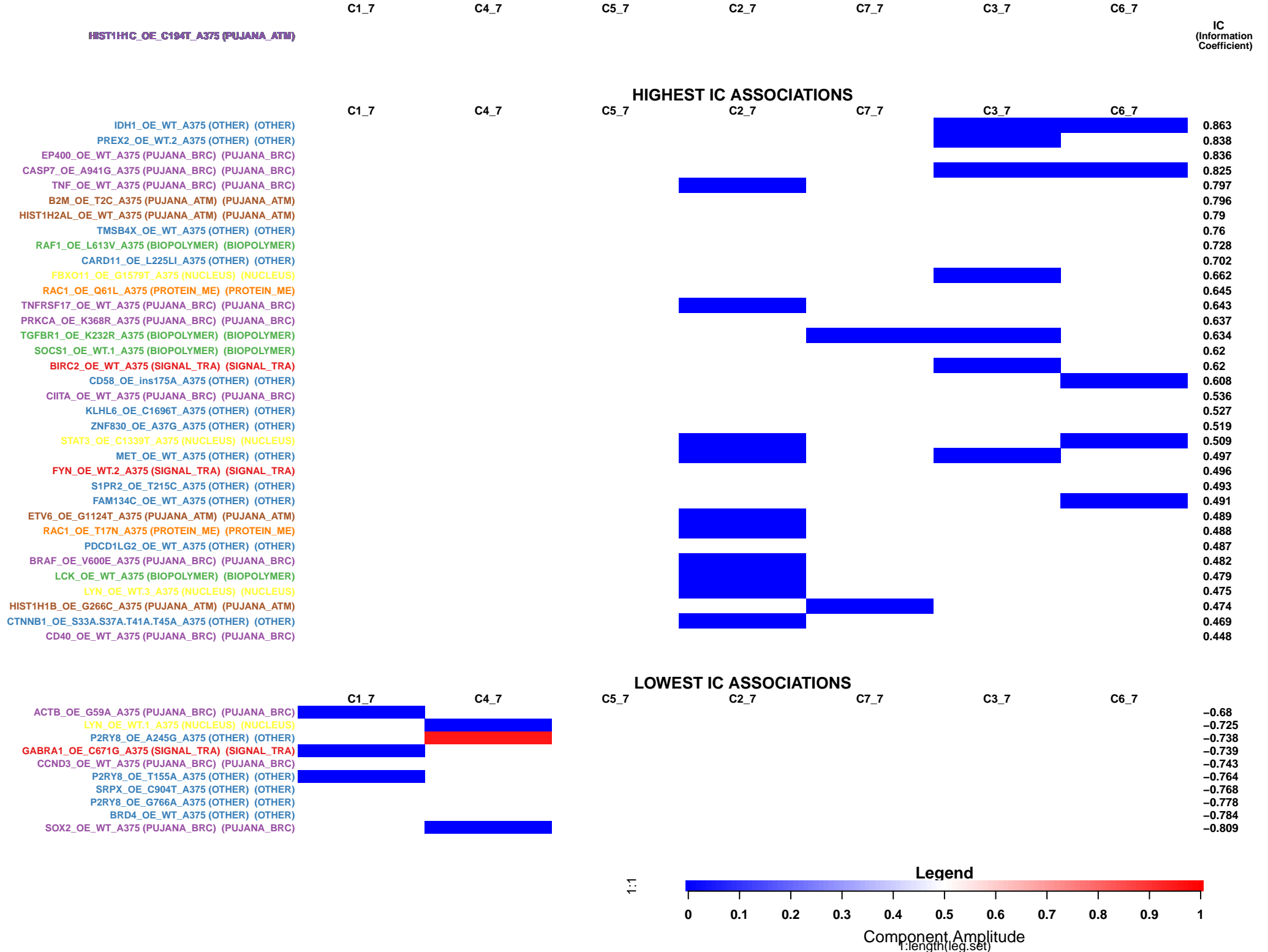


FAM134C\_OE\_WT\_A375 (OTHER)

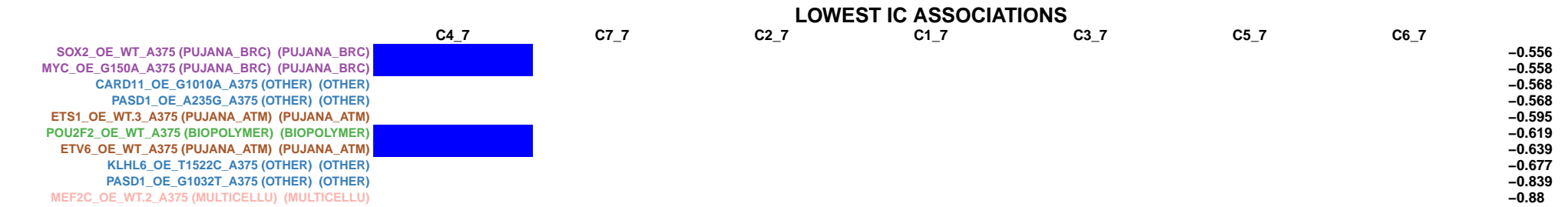




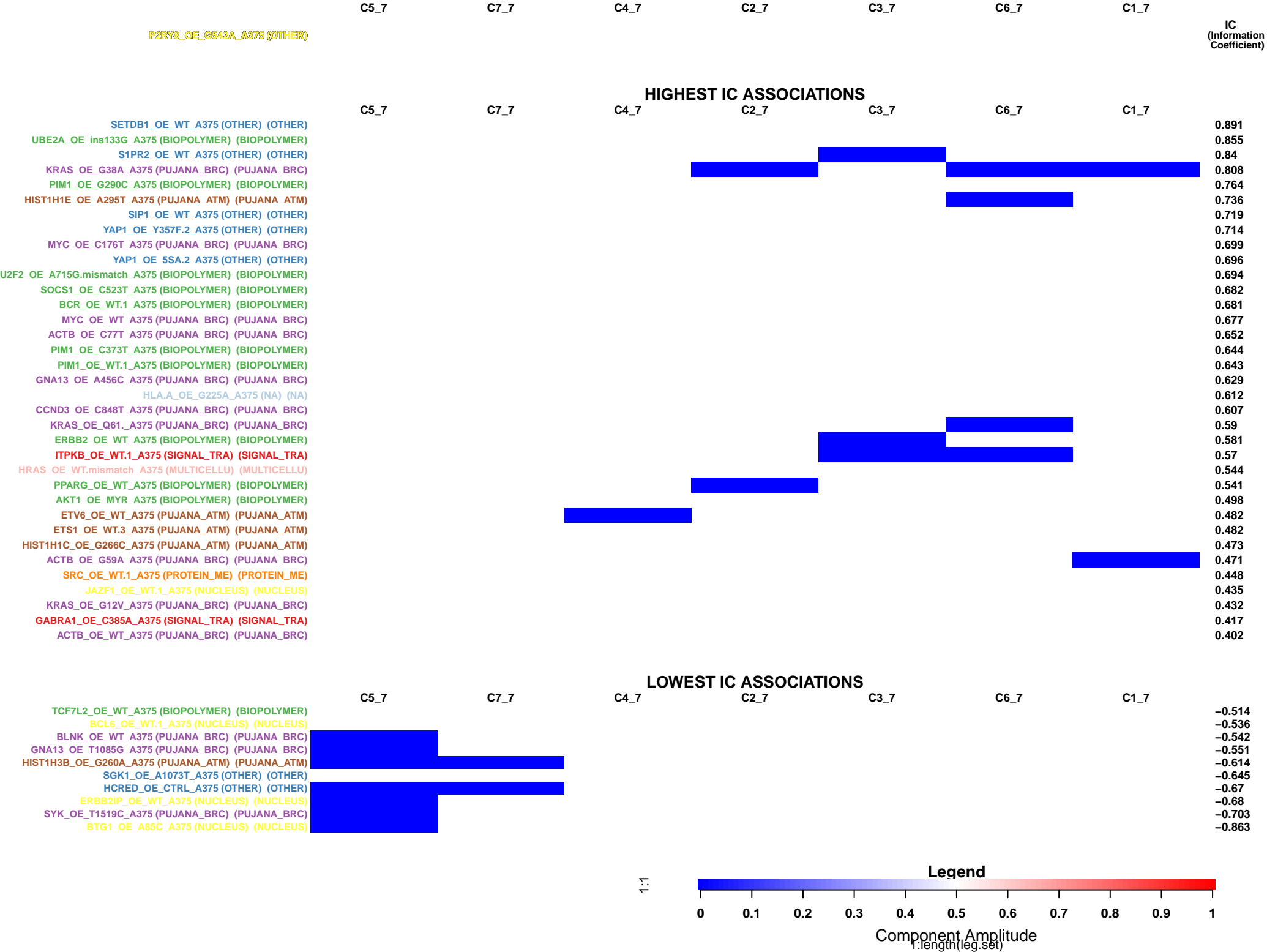
HIST1H1C\_OE\_C194T\_A375 (PUJANA\_ATM)



C4_7	C7_7	C2_7	C1_7	C3_7	C5_7	C6_7	IC (Information Coefficient)
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P2RY8\_OE\_G542A\_A375 (OTHER)



C2_7	C3_7	C7_7	C4_7	C5_7	C1_7	C6_7
------	------	------	------	------	------	------

IC  
Information  
Coefficient)

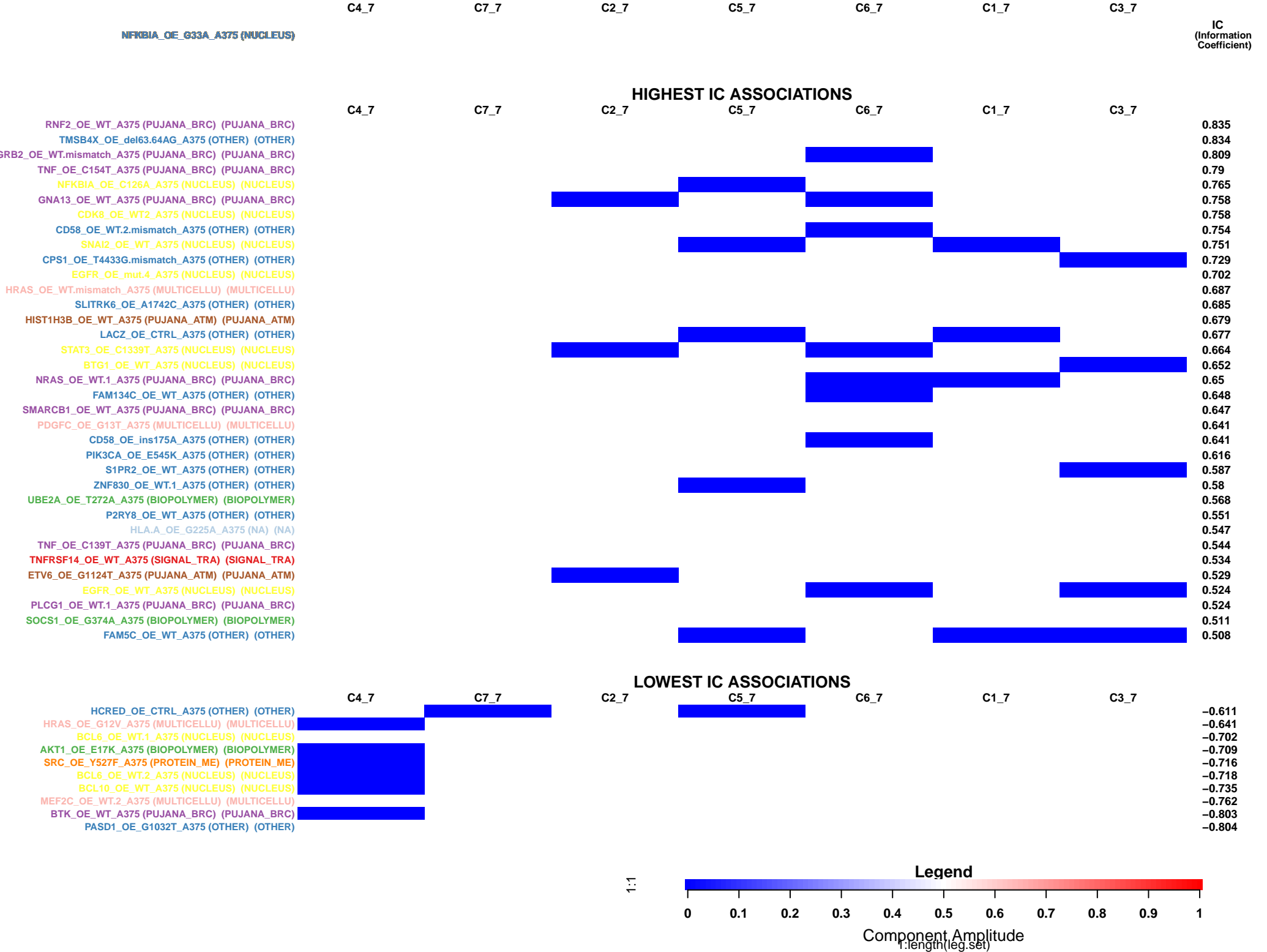
C2_7	C3_7	C7_7	C4_7	C5_7	C1_7	C6_7
------	------	------	------	------	------	------



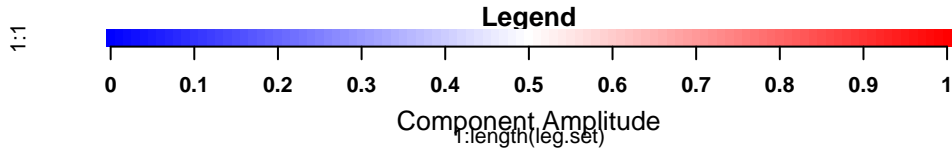
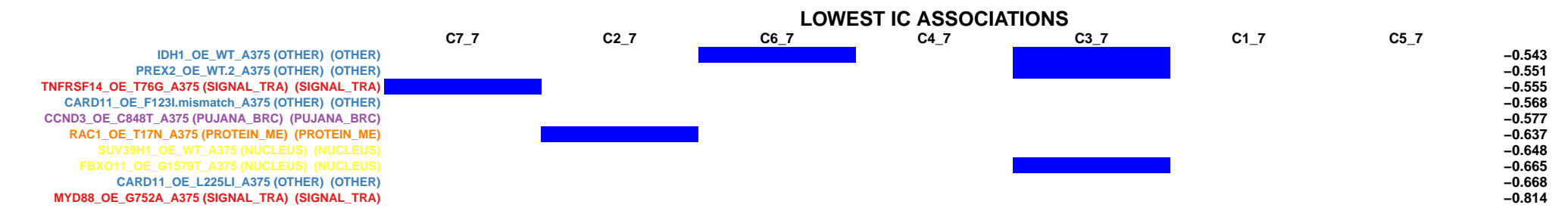
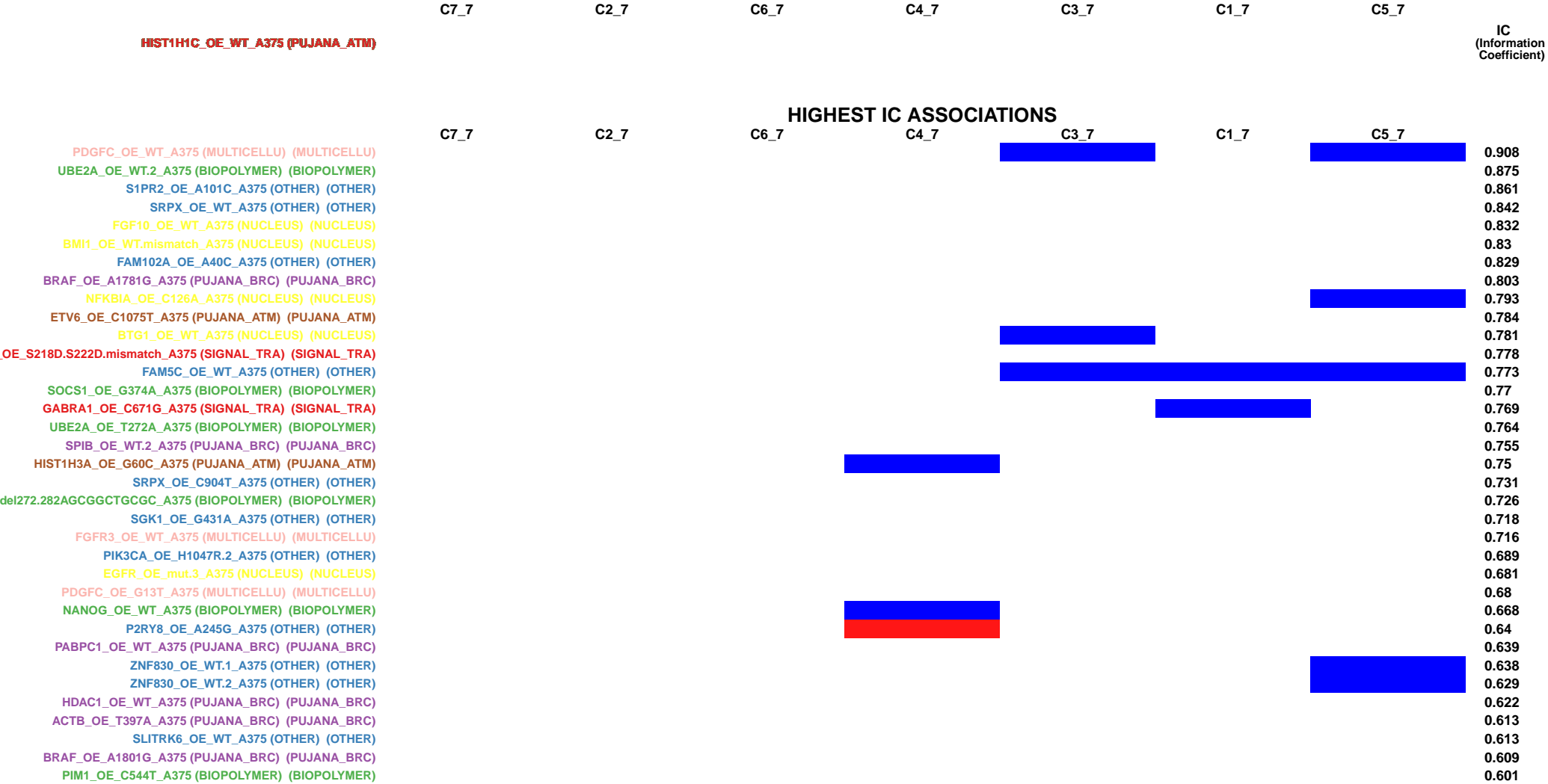
Legend

length (leng.sət)

NFKBIA\_OE\_G33A\_A375 (NUCLEUS)



HIST1H1C\_OE\_WT\_A375 (PUJANA\_ATM)

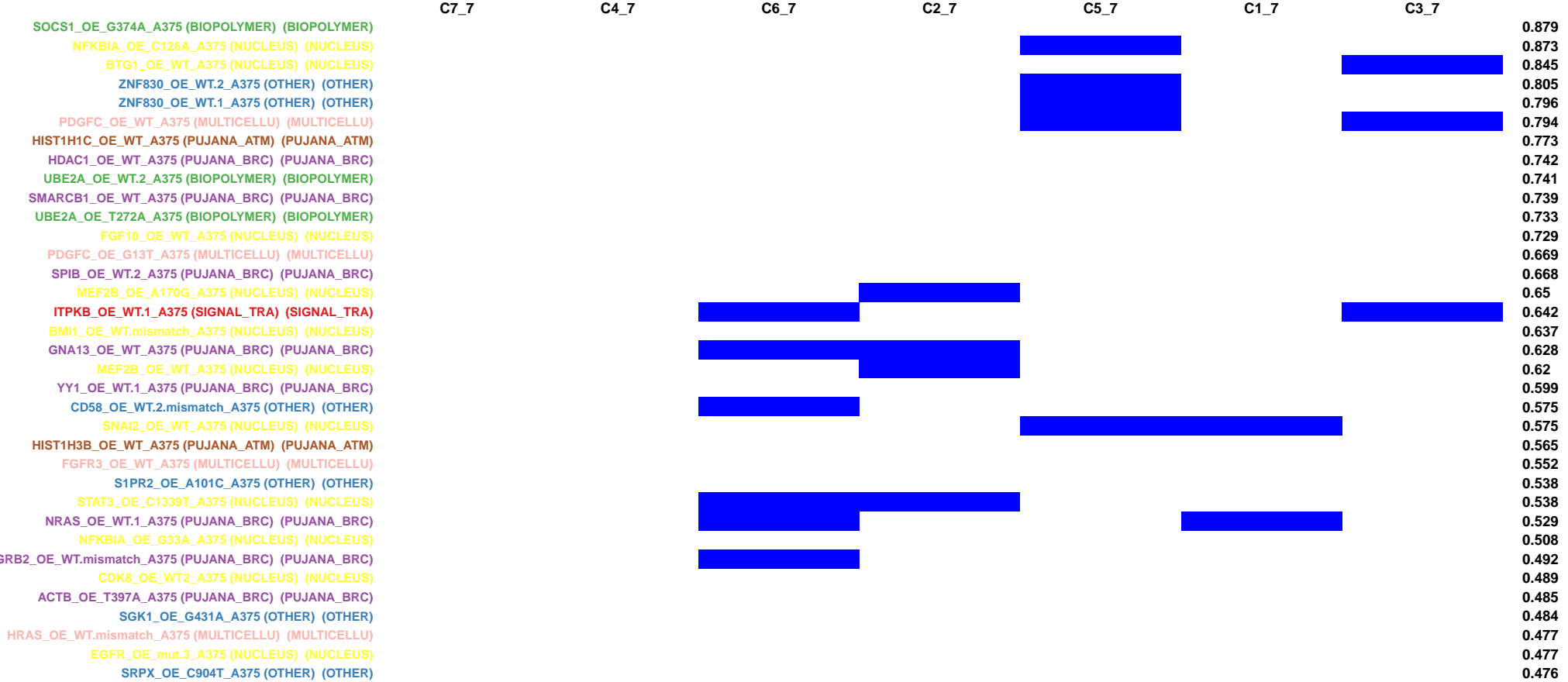


FAM5C\_OE\_WT\_A375 (OTHER)

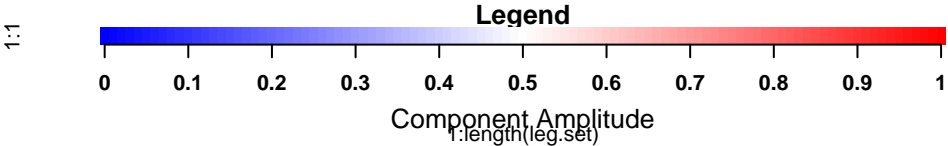
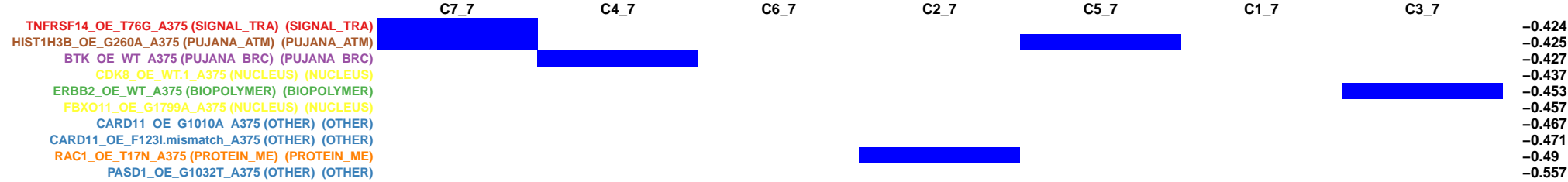
FAM5C\_OE\_WT\_A375 (OTHER)

IC  
(Information  
Coefficient)

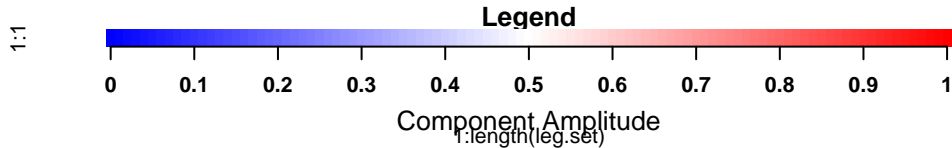
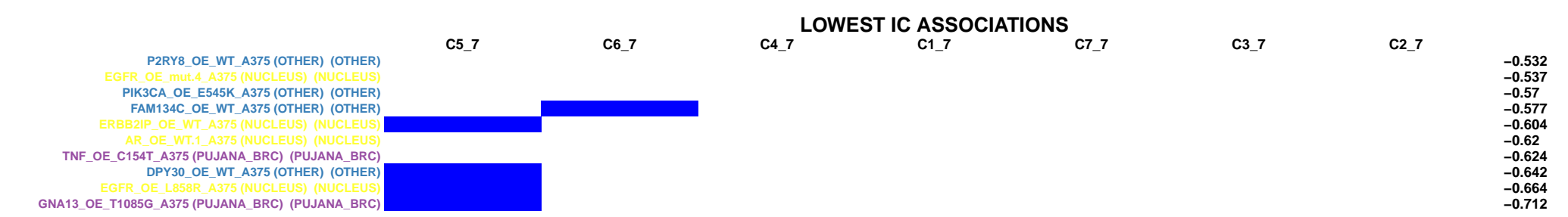
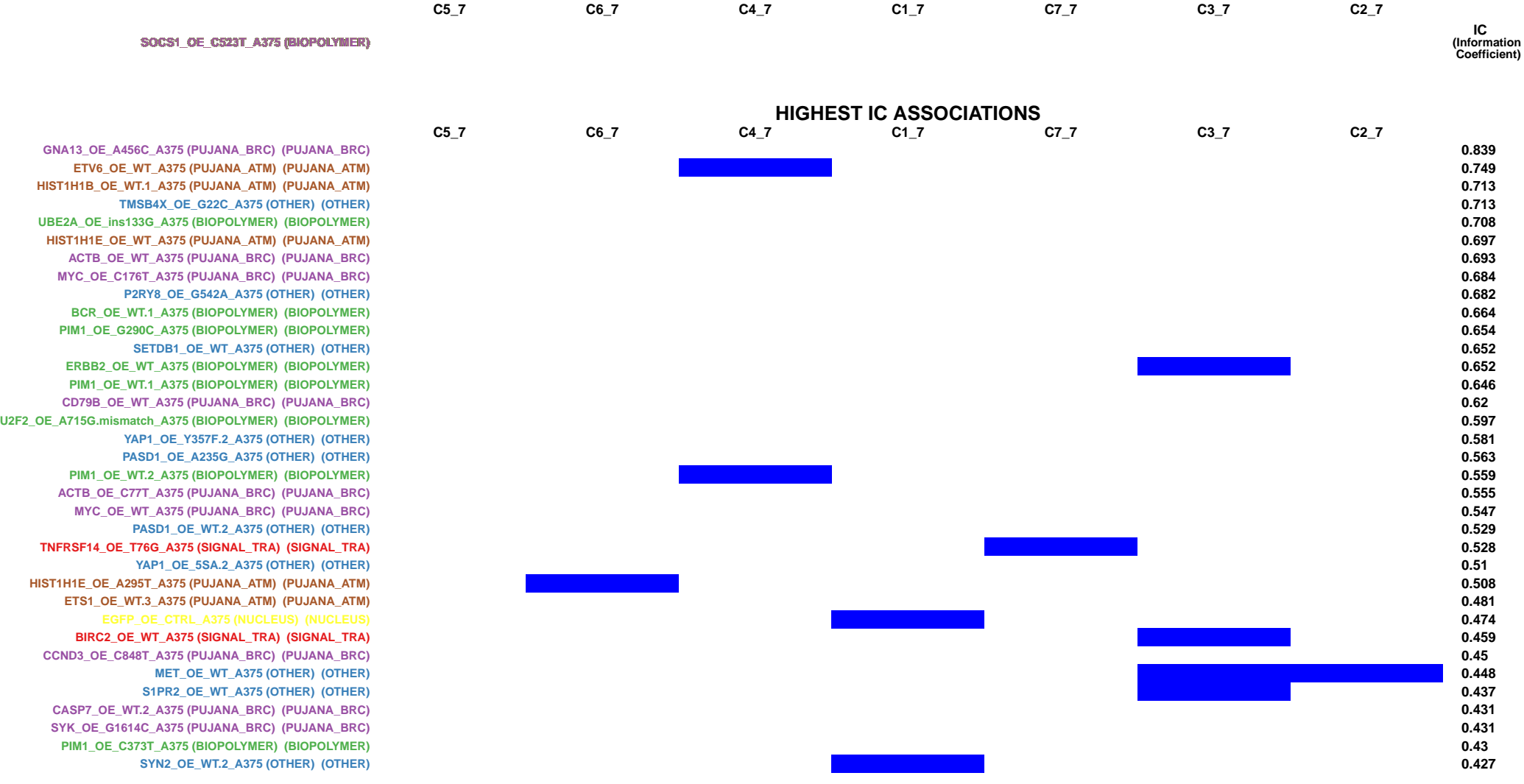
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS



SOCS1\_OE\_C523T\_A375 (BIOPOLYMER)



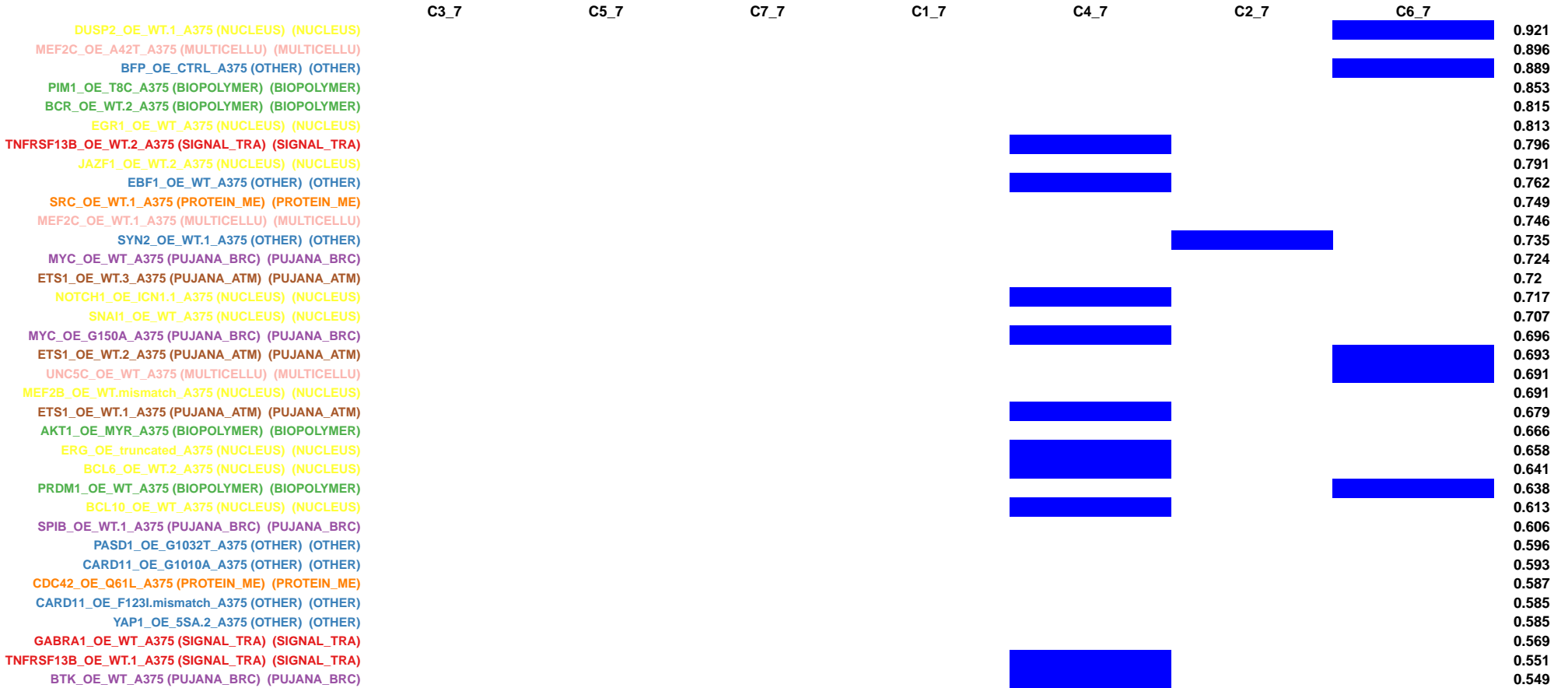


# DUSP2\_OE\_WT.2\_A375 (NUCLEUS)

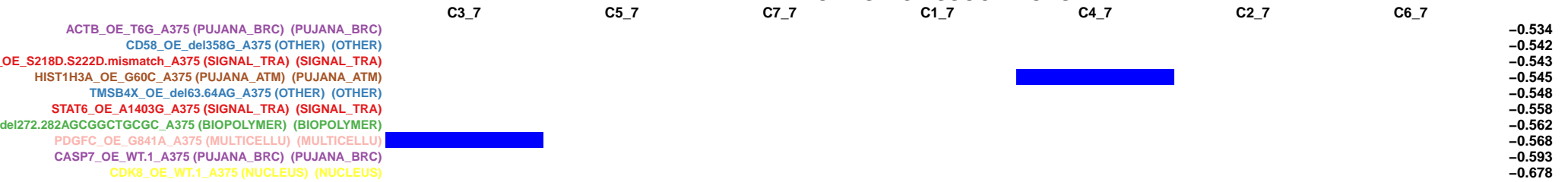
DUSP2\_OE\_WT.2\_A375 (NUCLEUS)

IC  
(Information  
Coefficient)

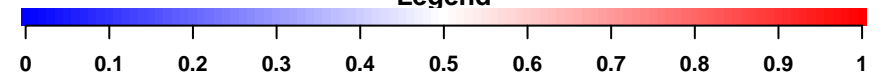
## HIGHEST IC ASSOCIATIONS



## LOWEST IC ASSOCIATIONS



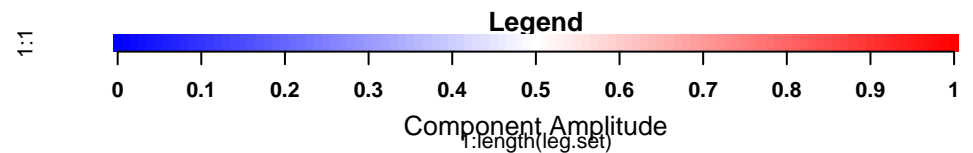
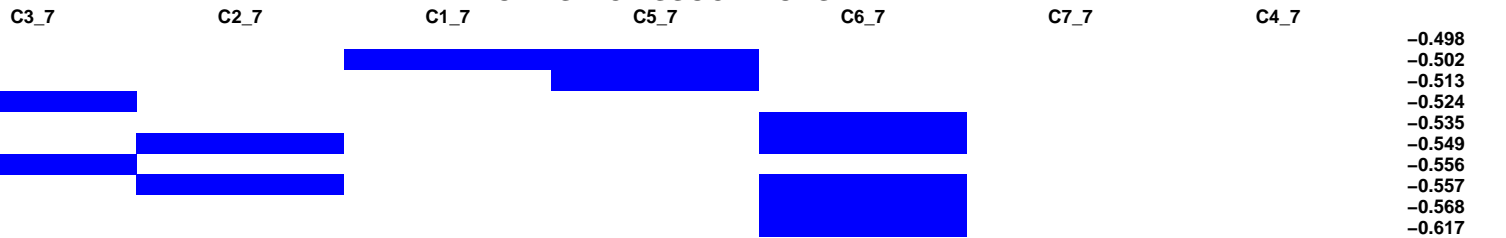
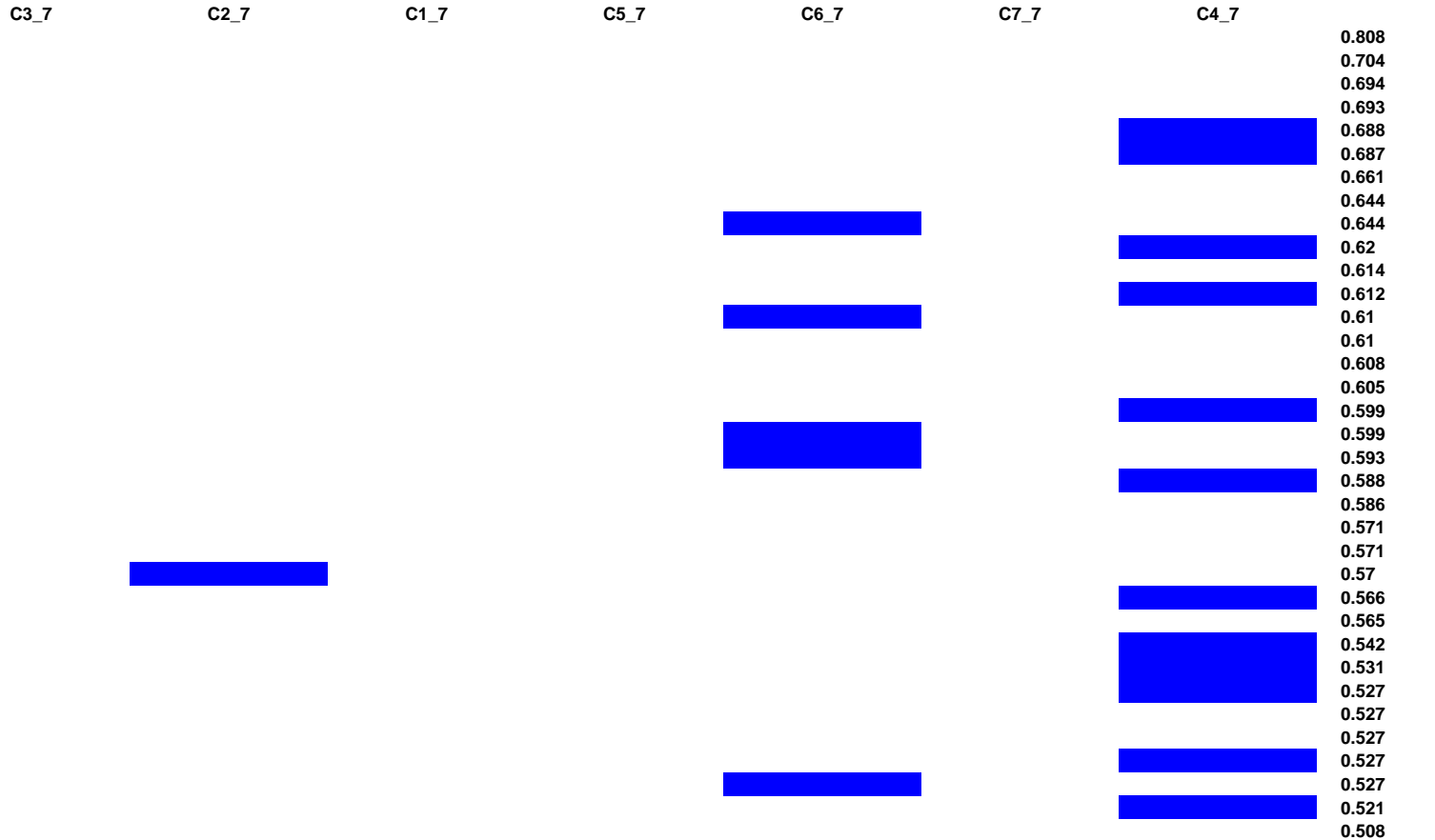
## Legend



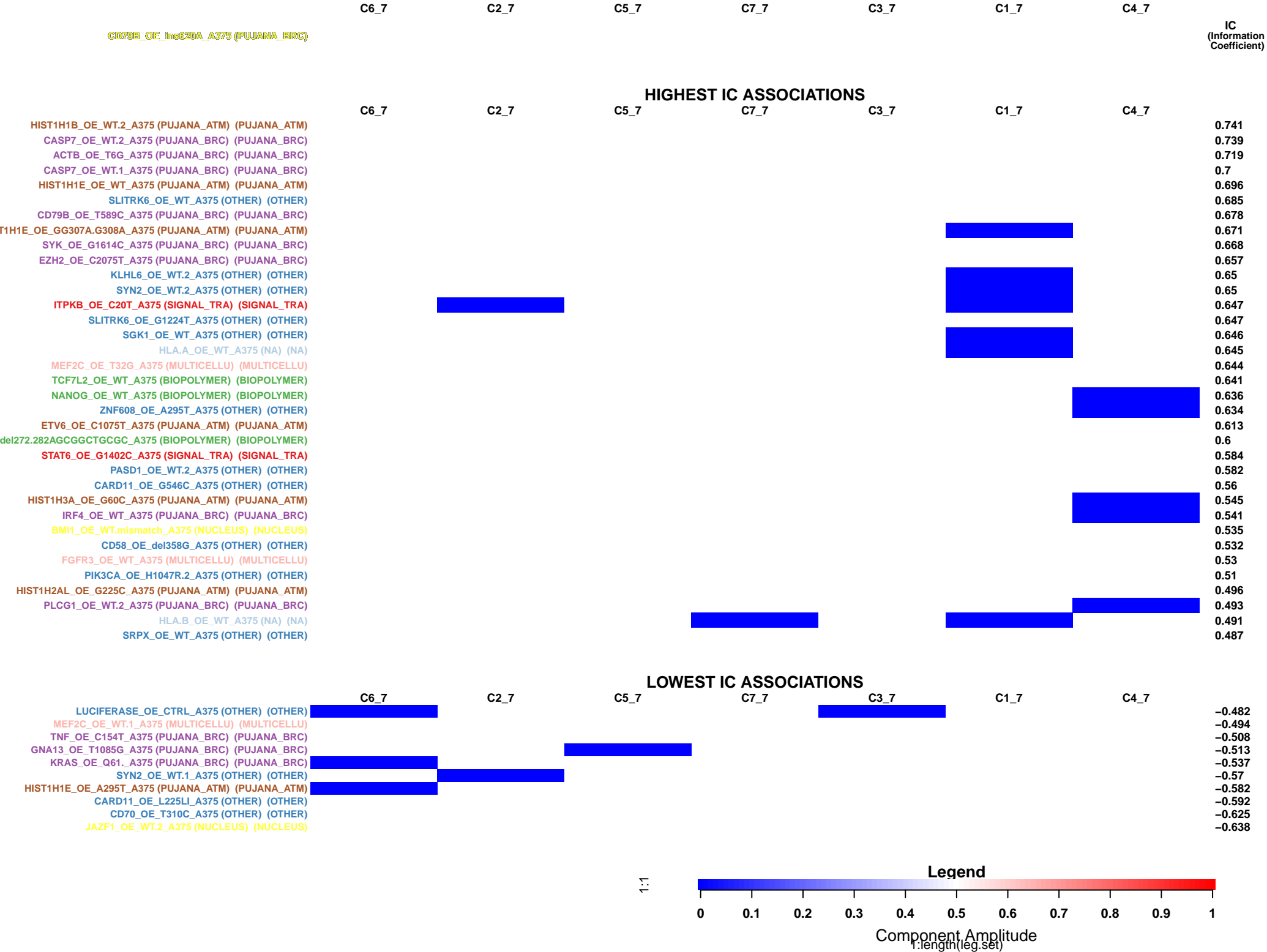
Component Amplitude

1:length(leg.set)

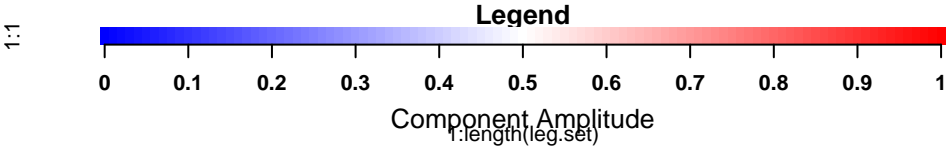
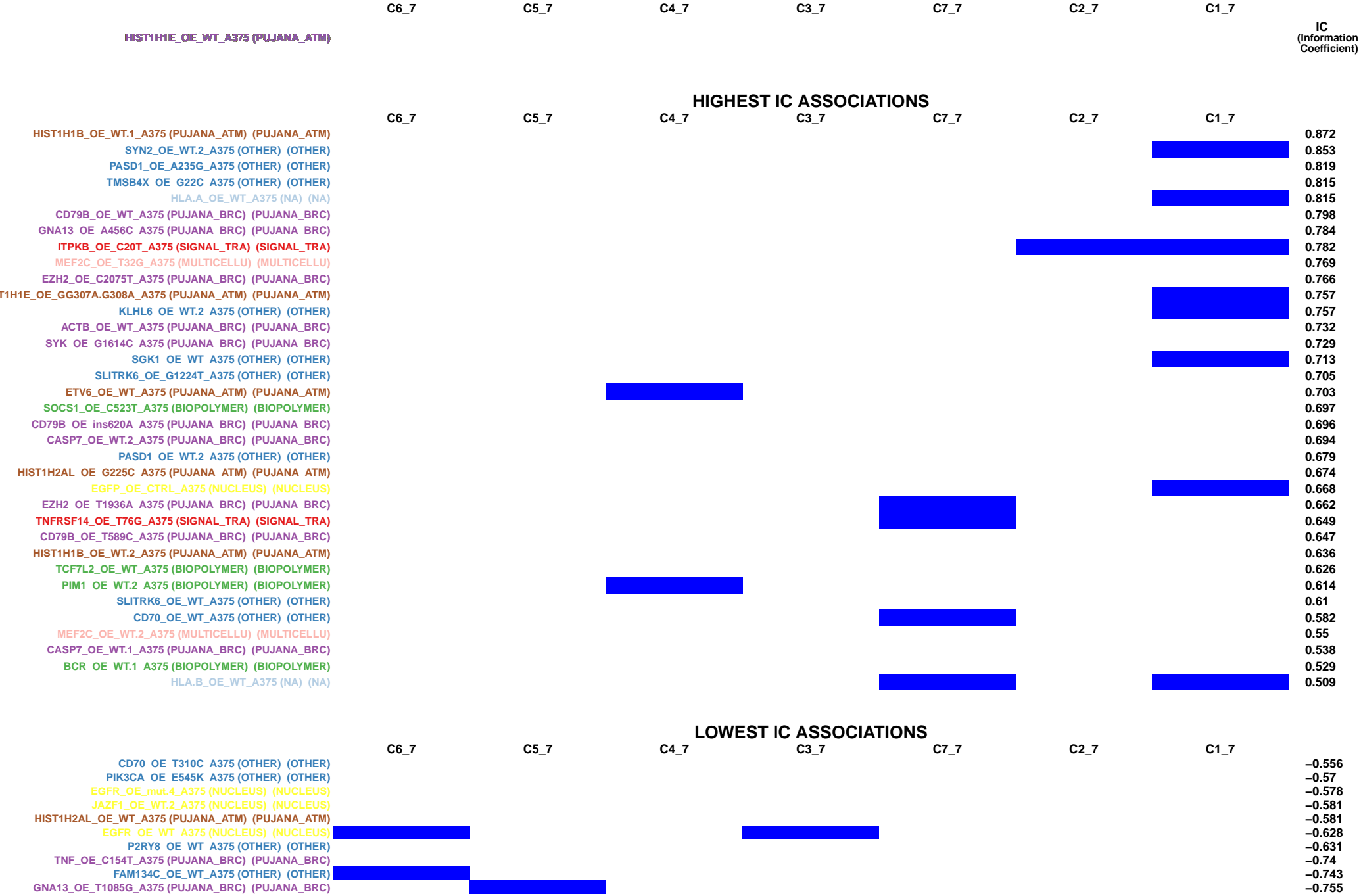
C3_7	C2_7	C1_7	C5_7	C6_7	C7_7	C4_7	IC (Information Coefficient)
------	------	------	------	------	------	------	------------------------------------



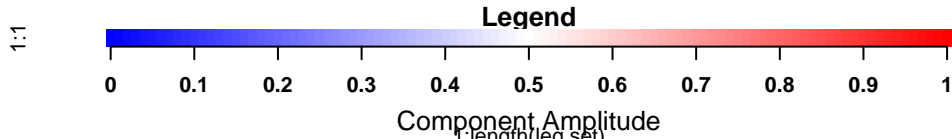
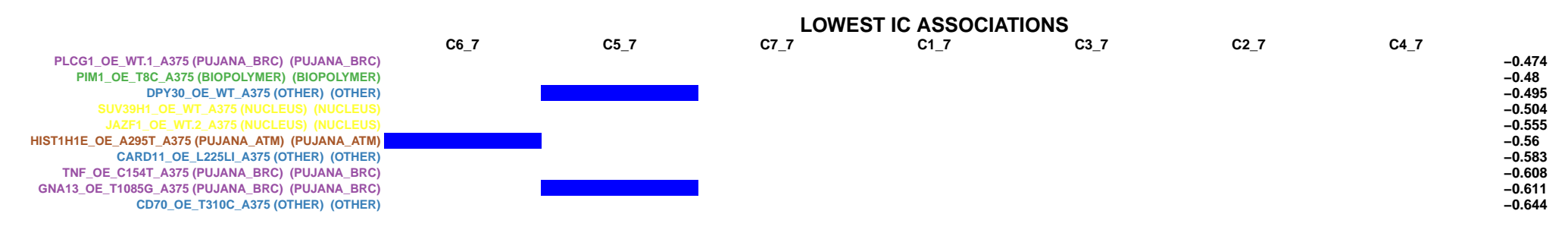
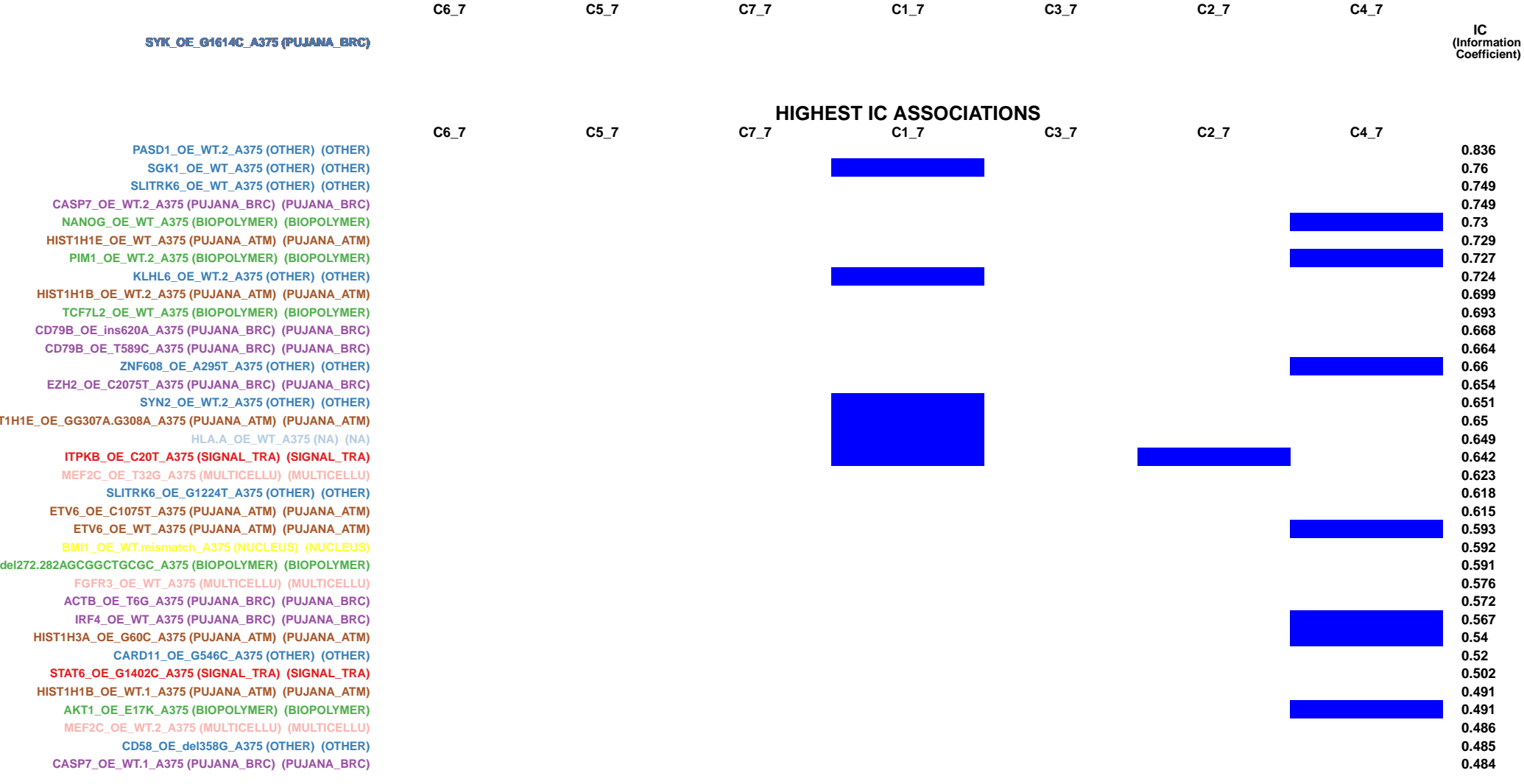
CD79B\_OE\_ins620A\_A375 (PUJANA\_BRC)



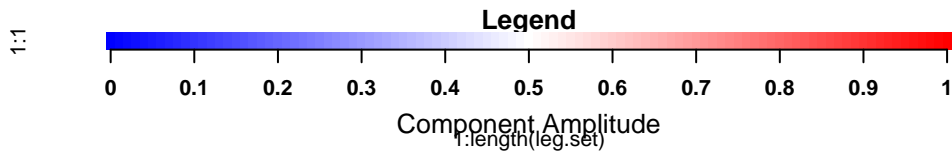
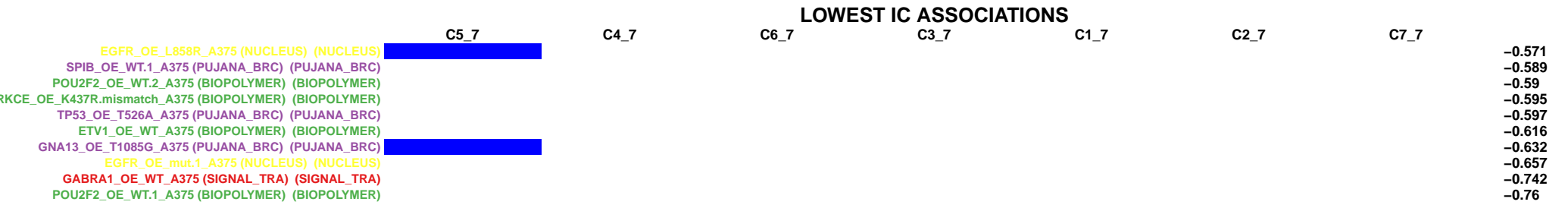
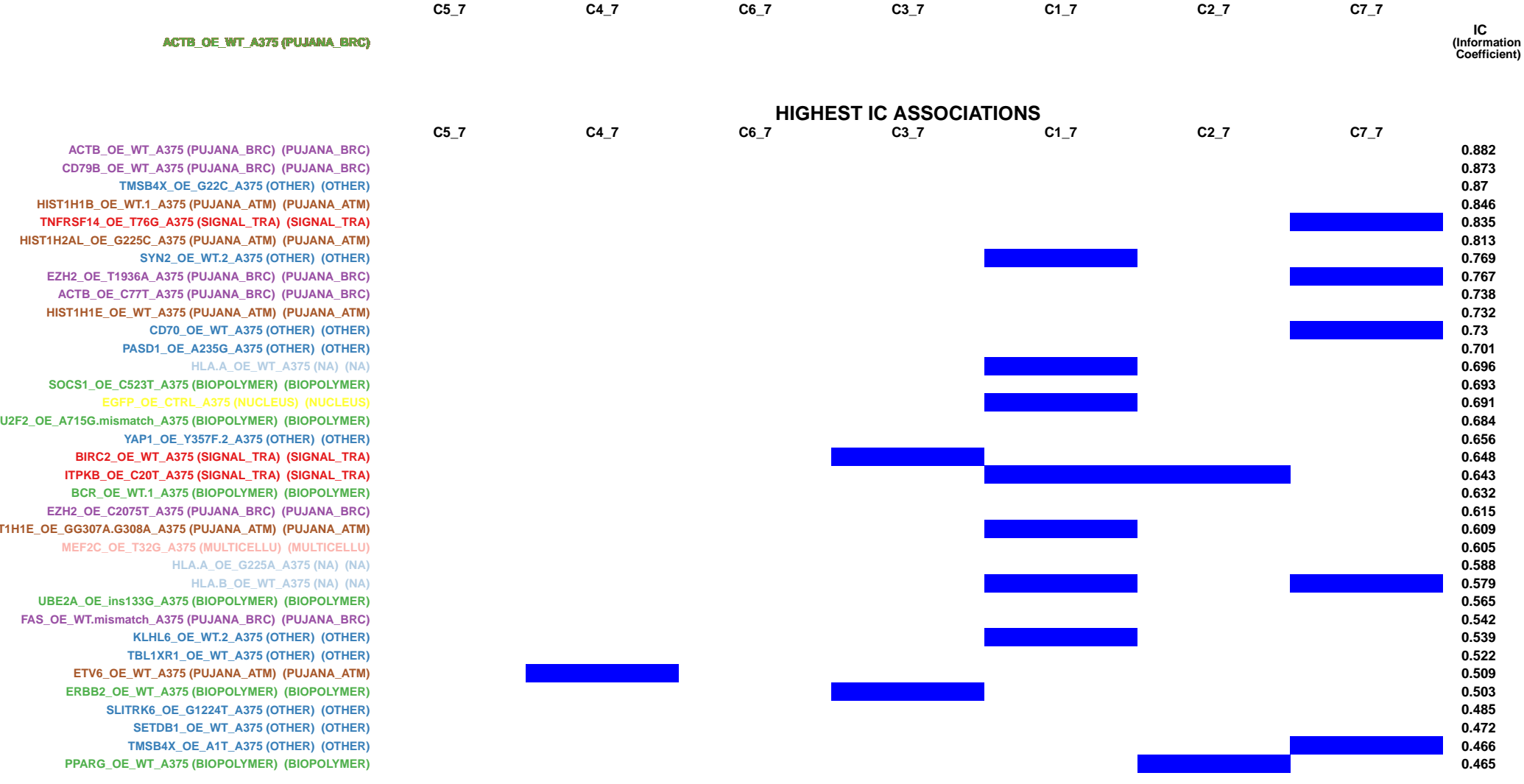
HIST1H1E\_OE\_WT\_A375 (PUJANA\_ATM)



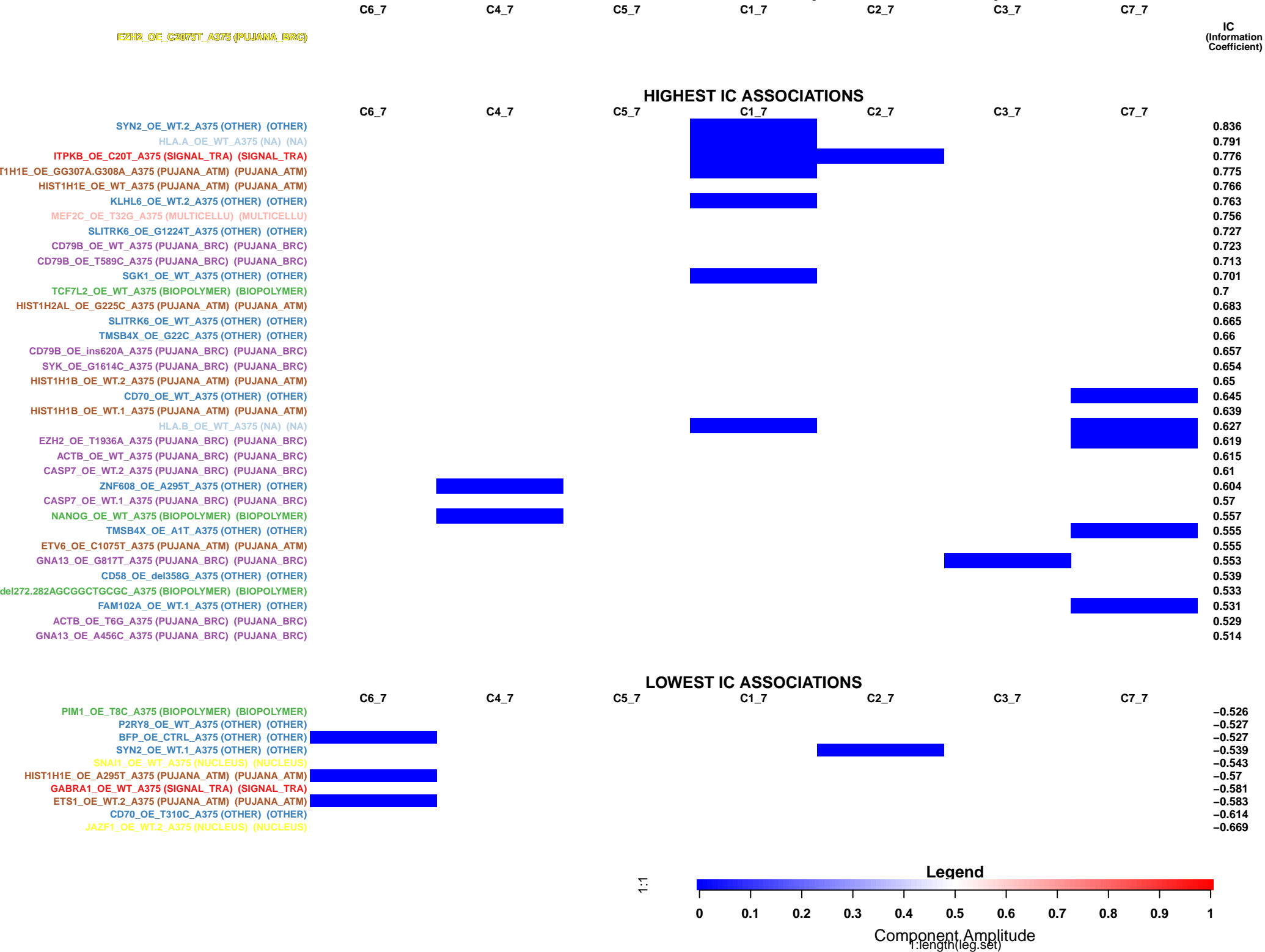
SYK\_OE\_G1614C\_A375 (PUJANA\_BRC)



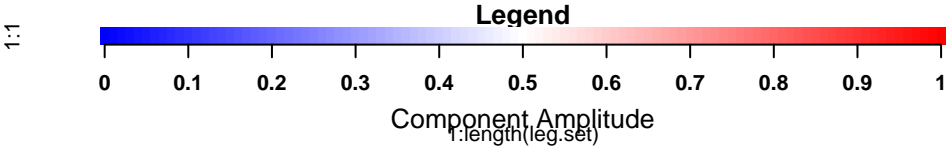
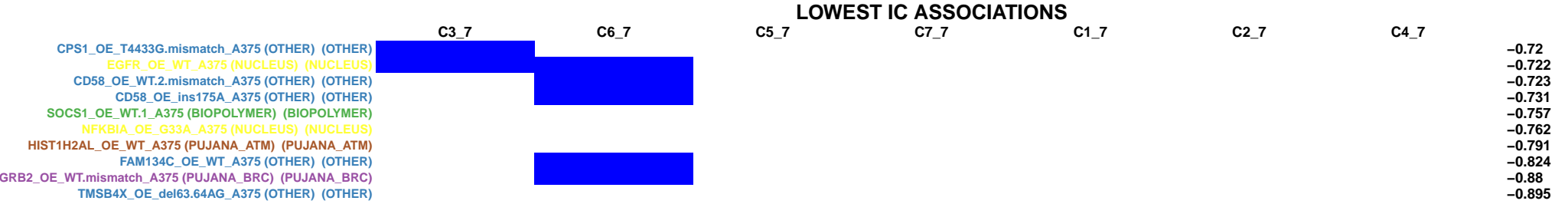
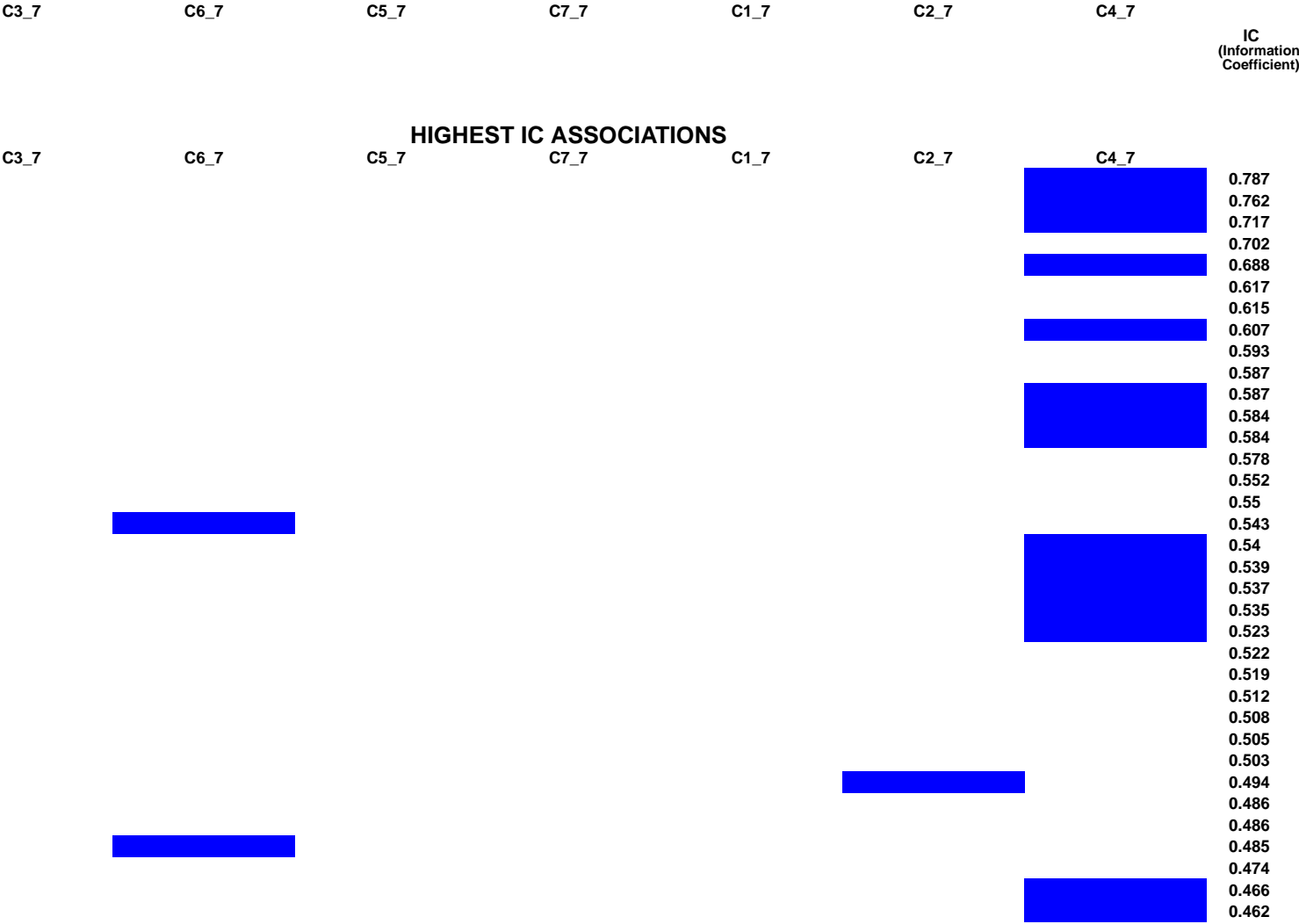
ACTB\_OE\_WT\_A375 (PUJANA\_BRC)



EZH2\_OE\_C2075T\_A375 (PUJANA\_BRC)

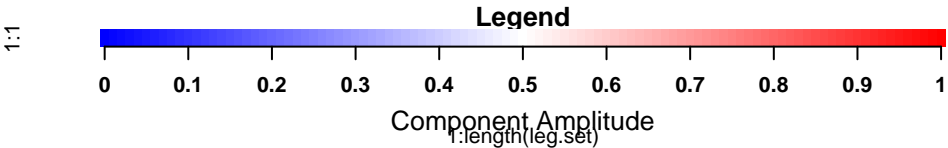
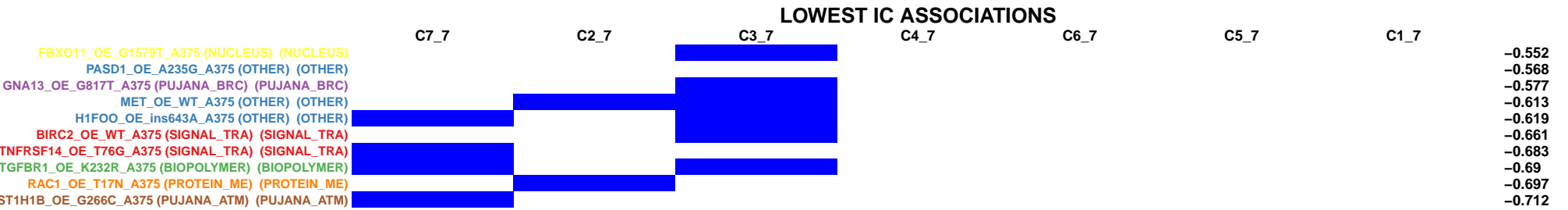
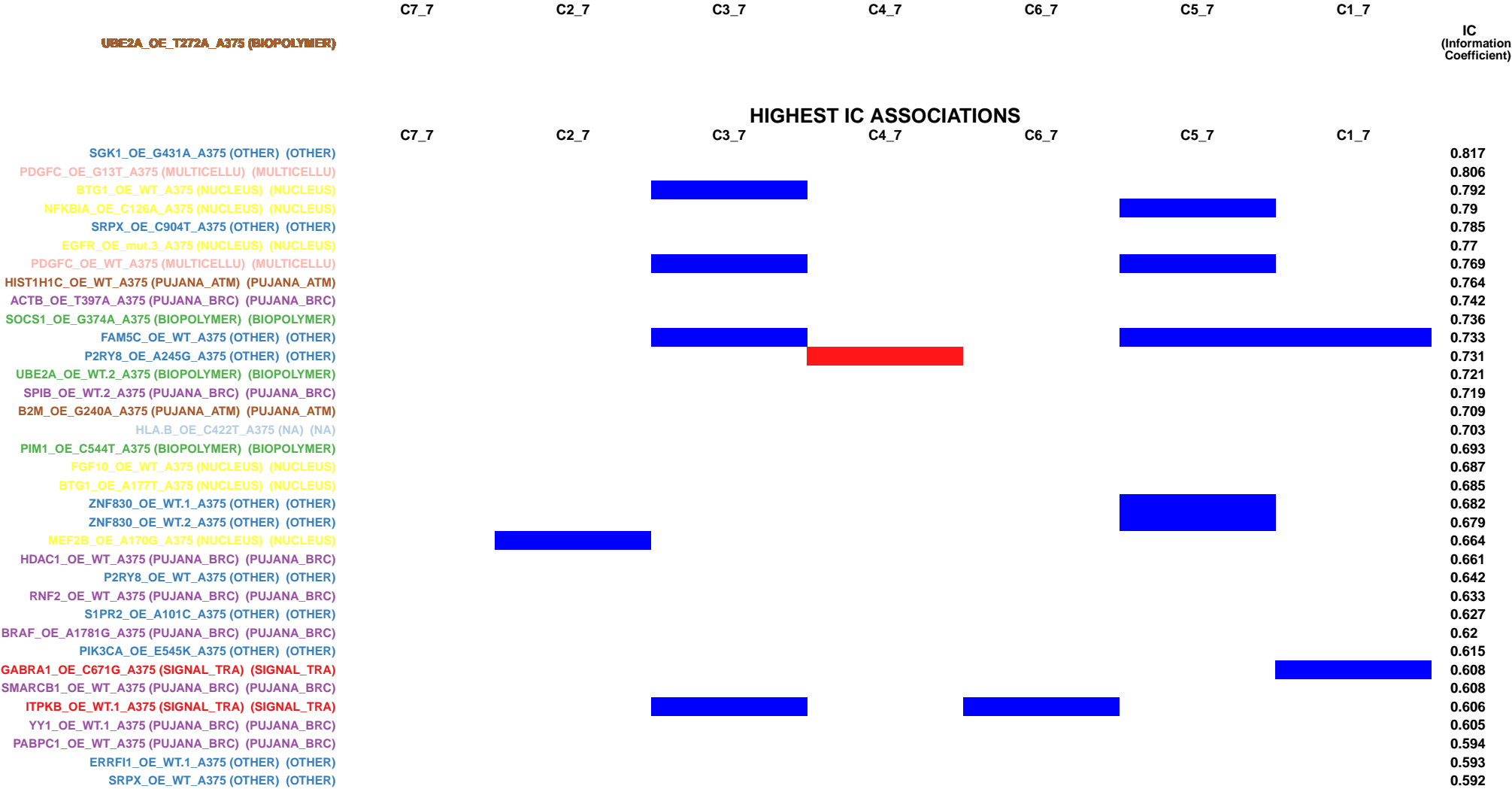


MEF2C\_OE\_WT.2\_A375 (MULTICELLU)

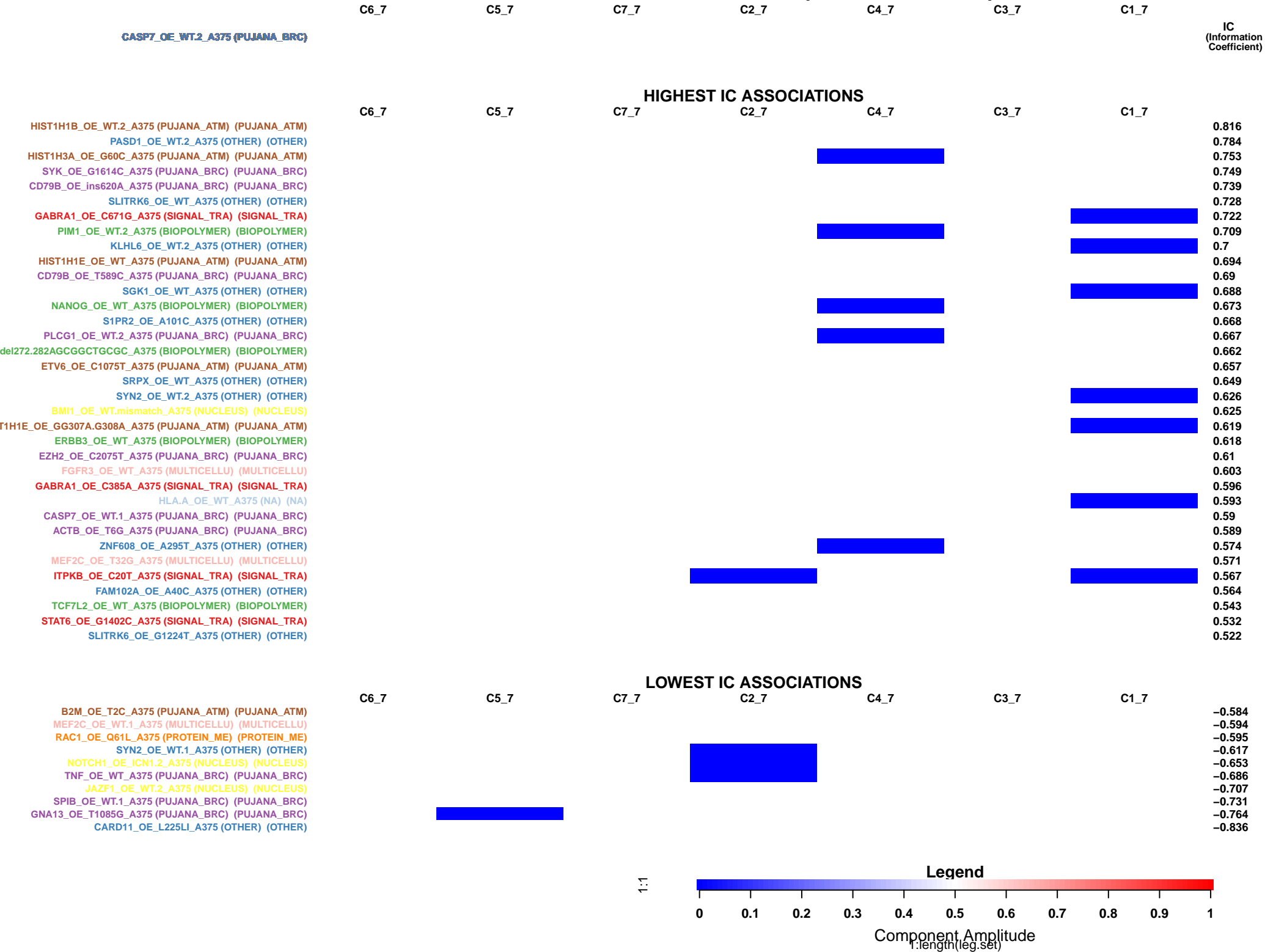




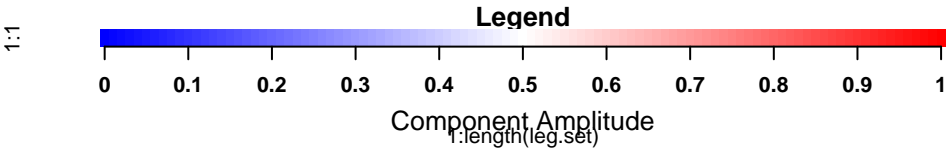
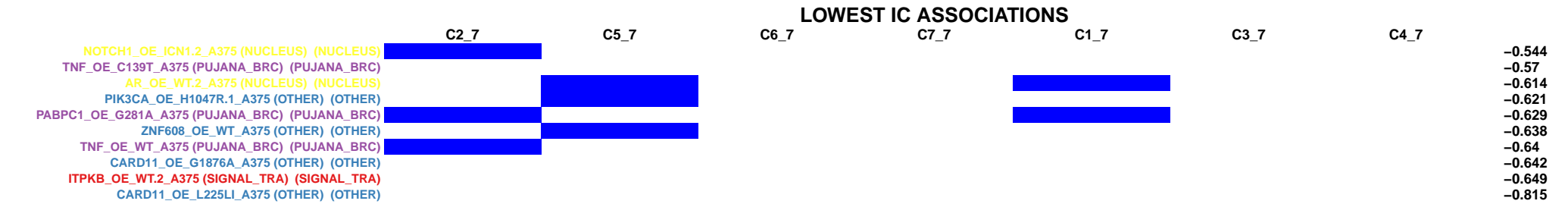
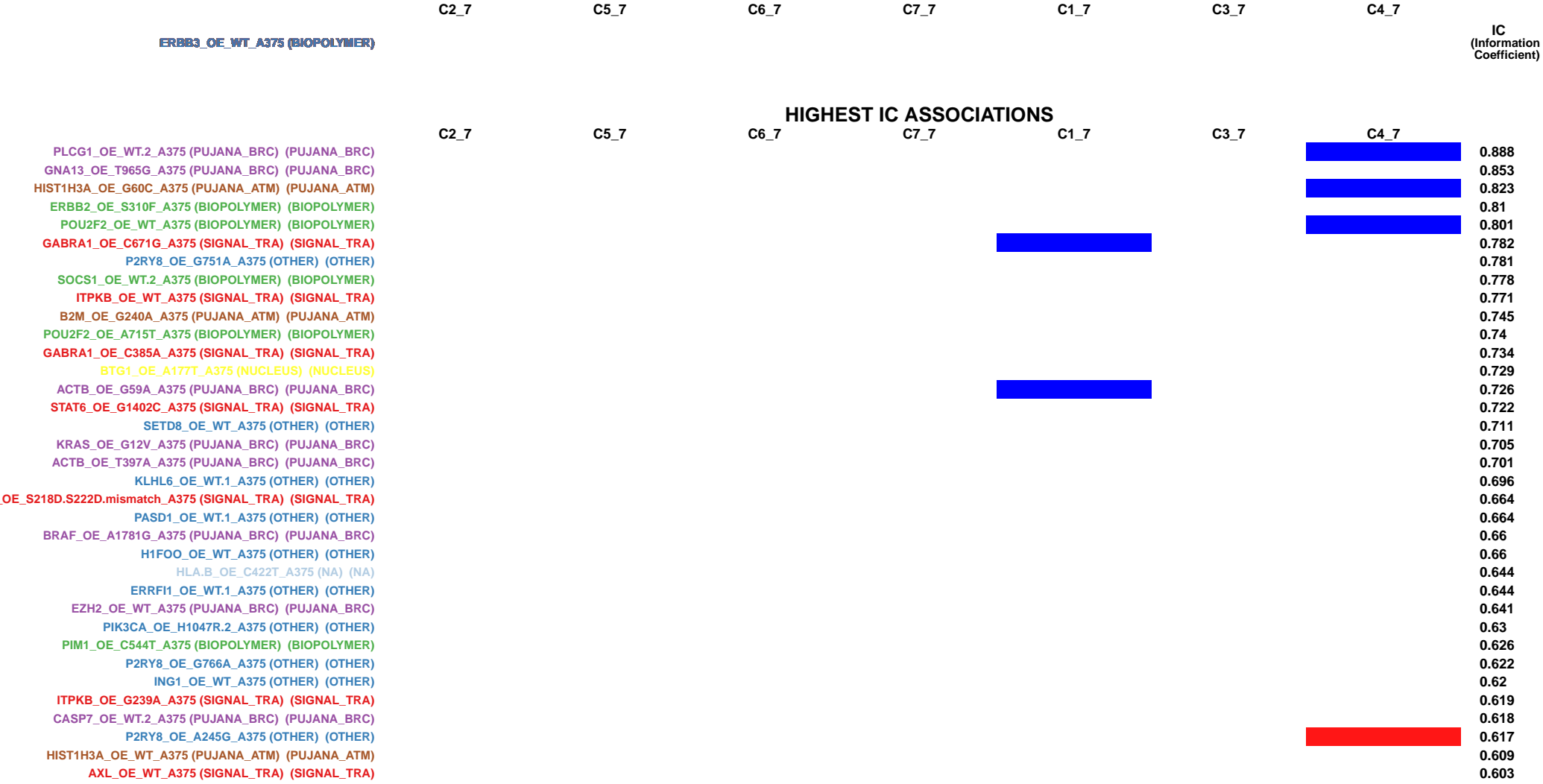
UBE2A\_OE\_T272A\_A375 (BIOPOLYMER)



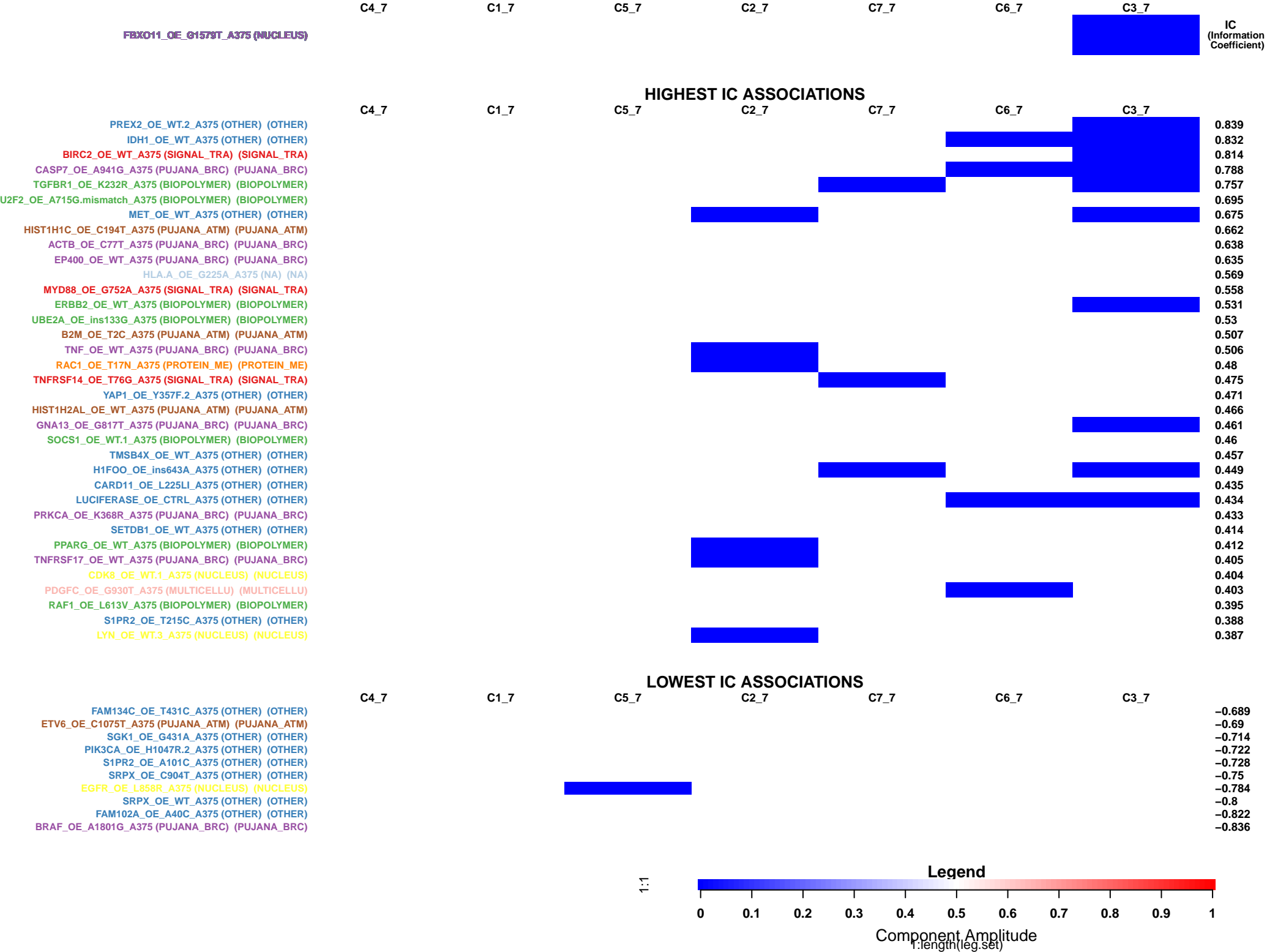
CASP7\_OE\_WT.2\_A375 (PUJANA\_BRC)



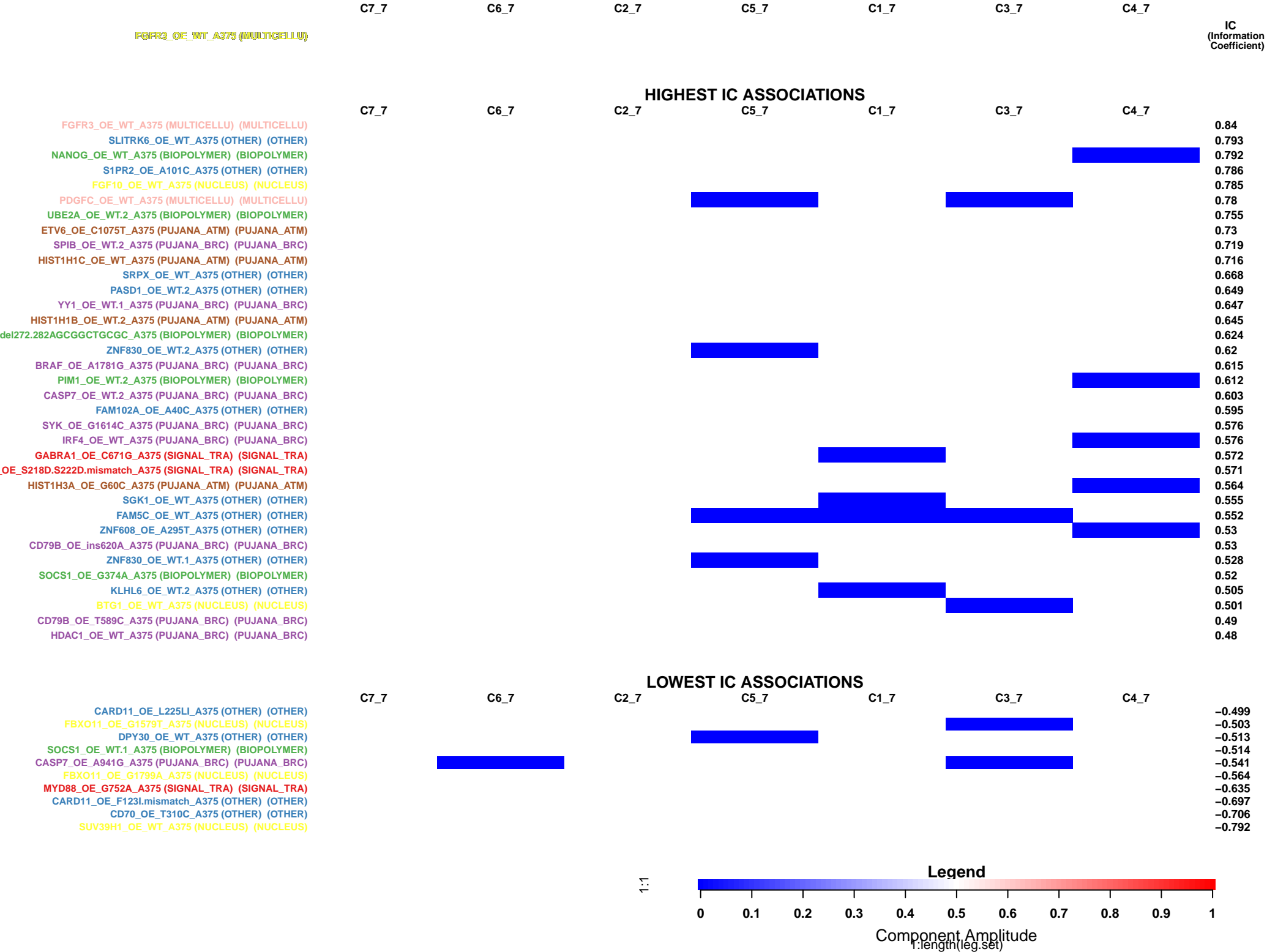
ERBB3\_OE\_WT\_A375 (BIOPOLYMER)



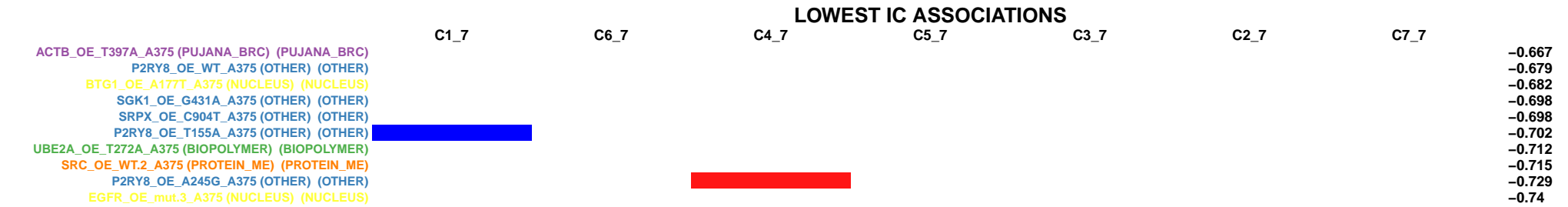
FBXO11\_OE\_G1579T\_A375 (NUCLEUS)



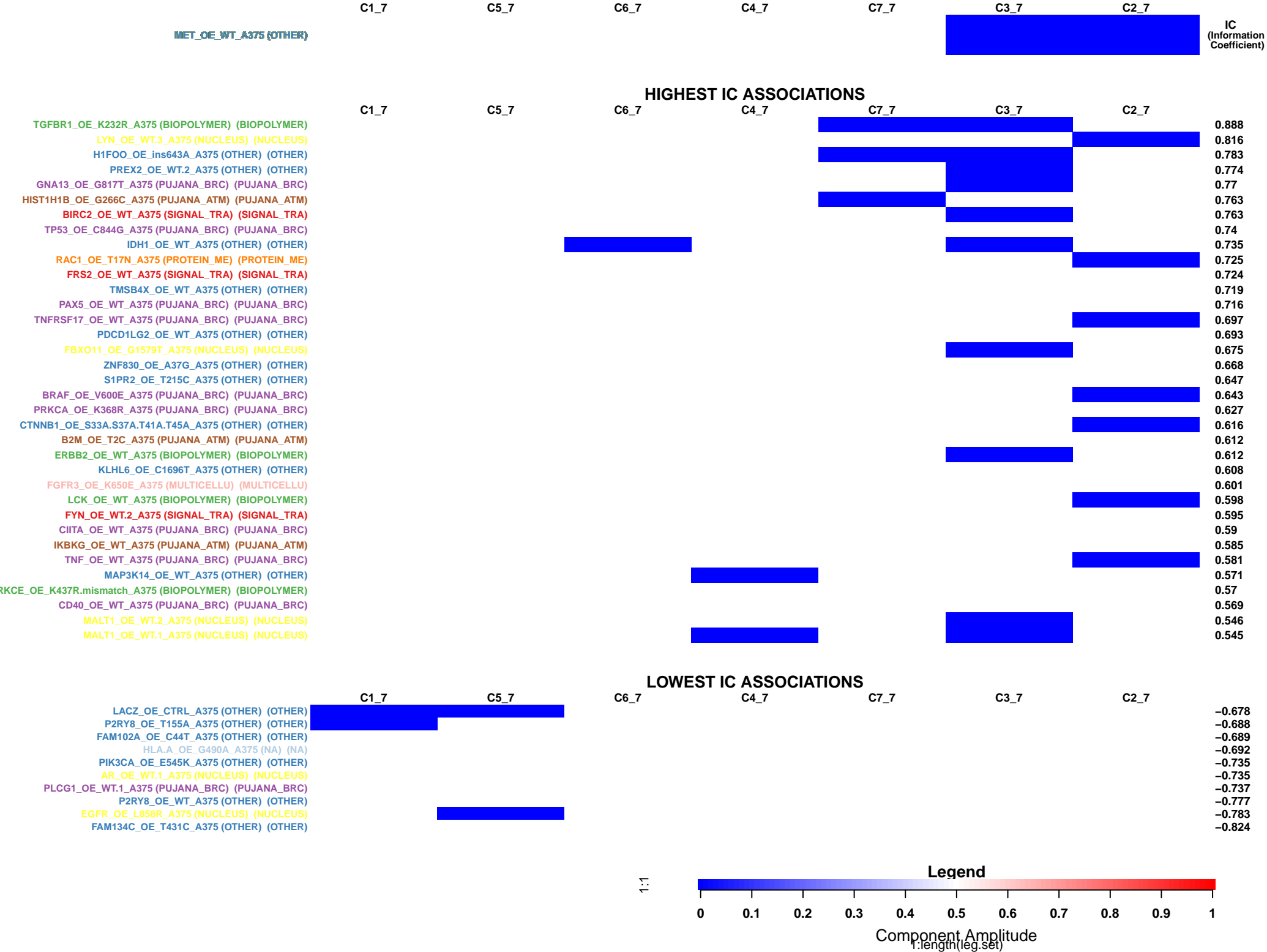
FGFR3\_OE\_WT\_A375 (MULTICELLU)



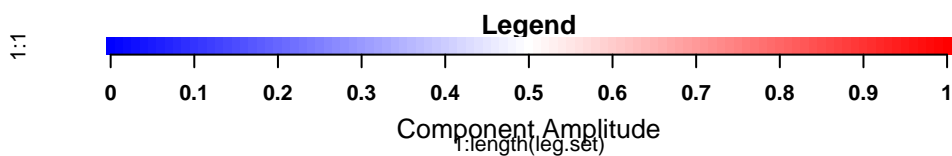
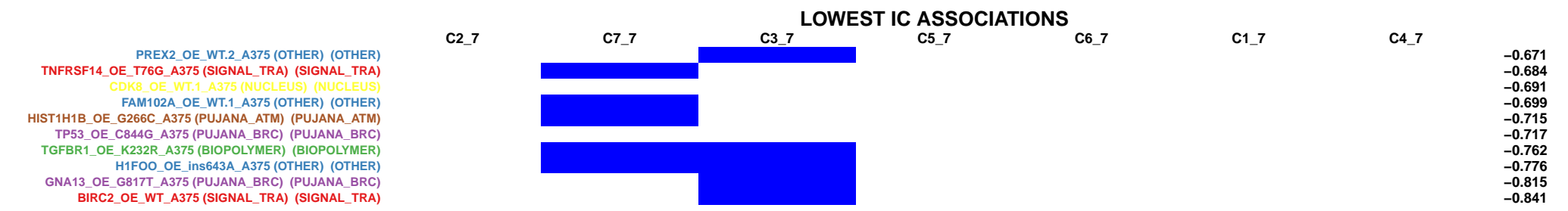
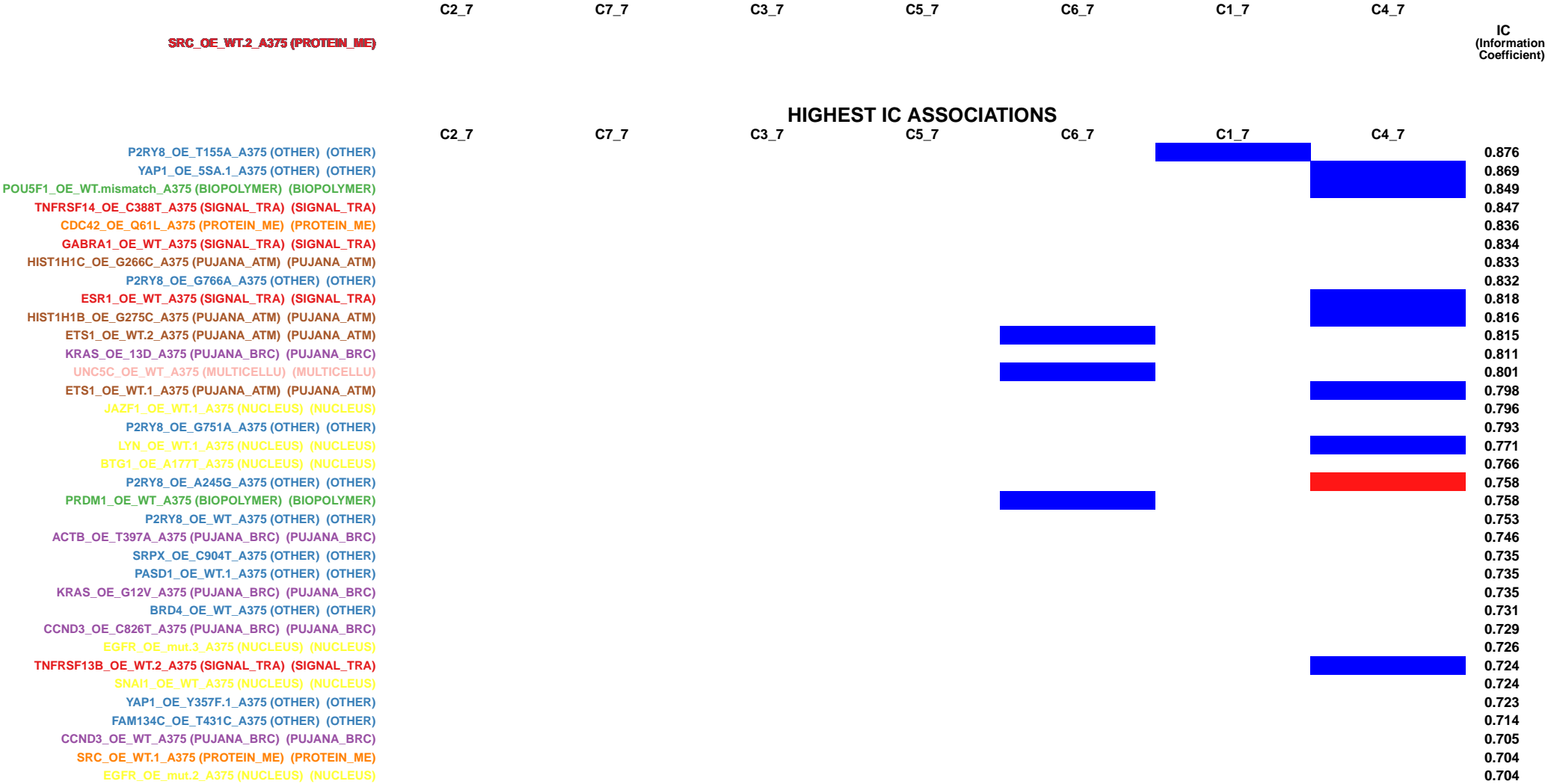
C1_7	C6_7	C4_7	C5_7	C3_7	C2_7	C7_7	IC (Information Coefficient)
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MET\_OE\_WT\_A375 (OTHER)

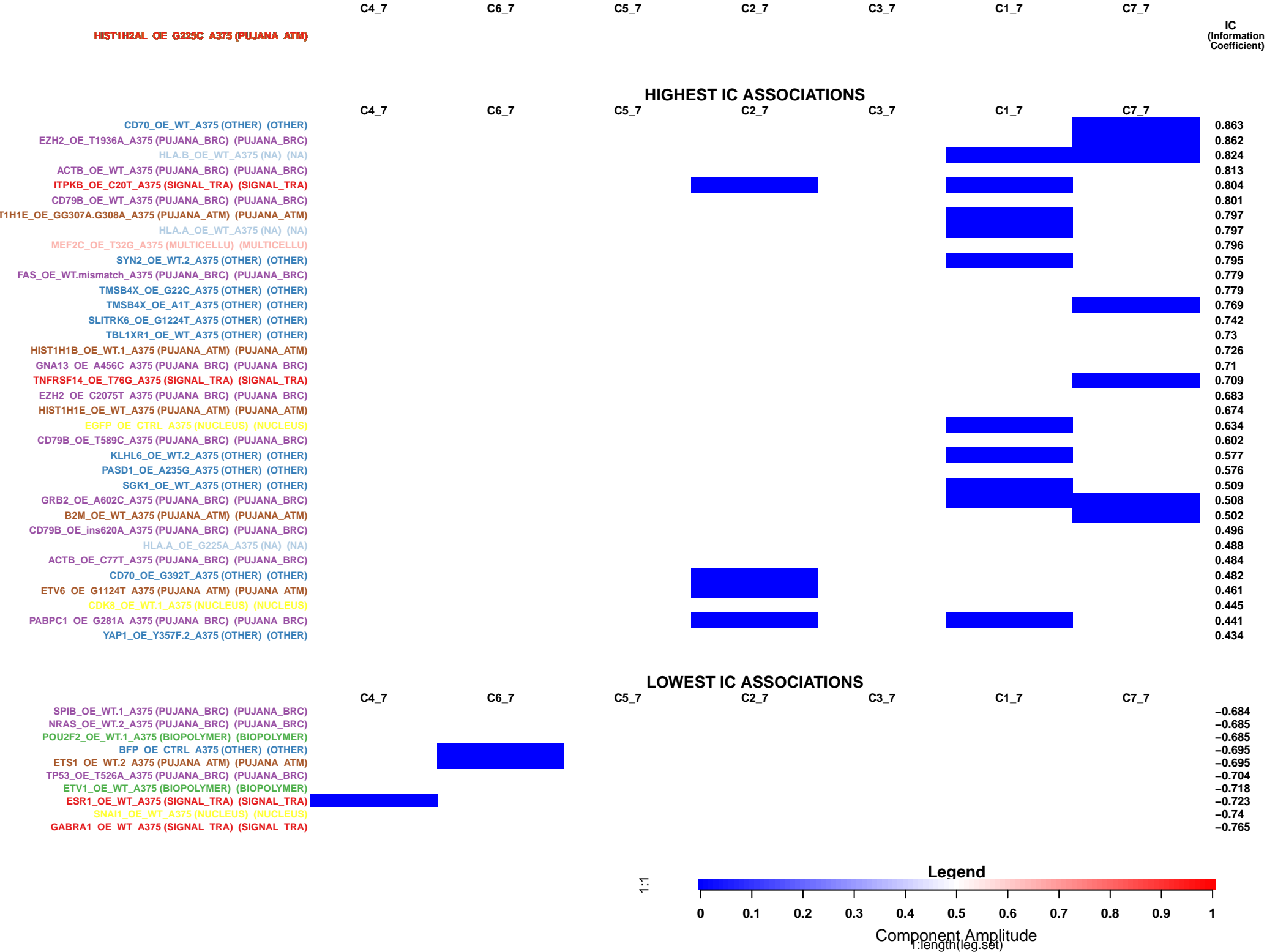


SRC\_OE\_WT.2\_A375 (PROTEIN\_ME)

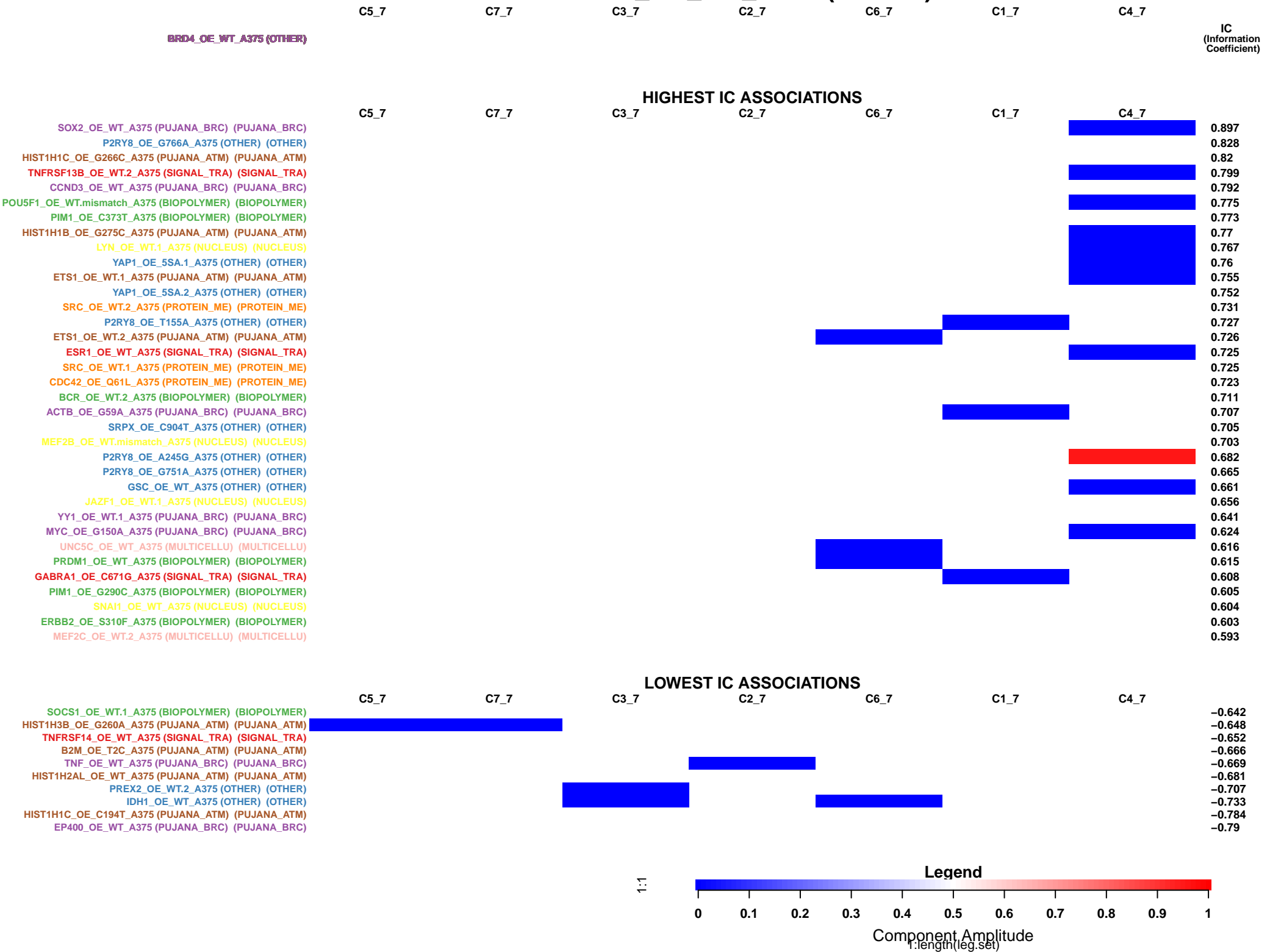




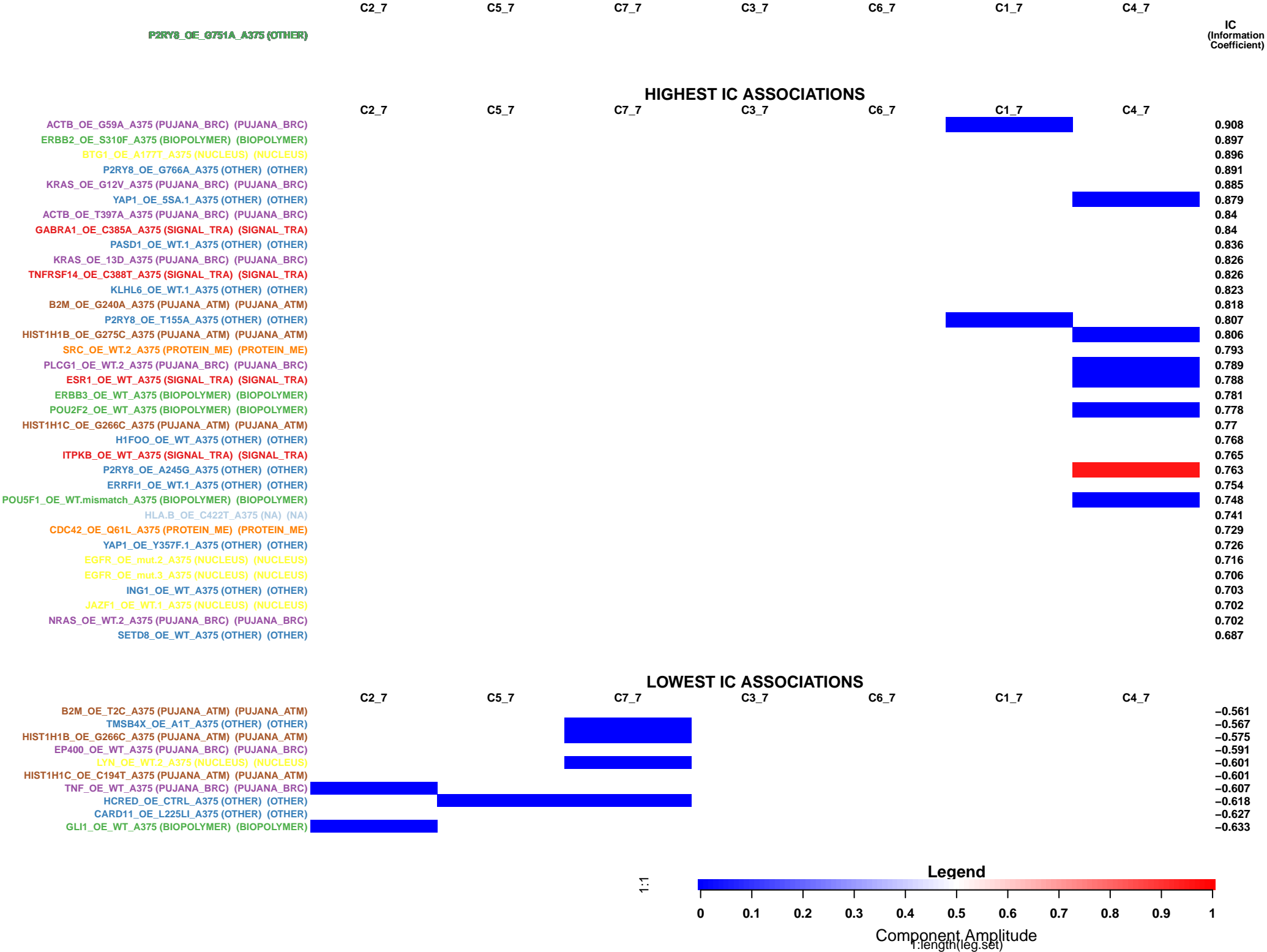
HIST1H2AL\_OE\_G225C\_A375 (PUJANA\_ATM)



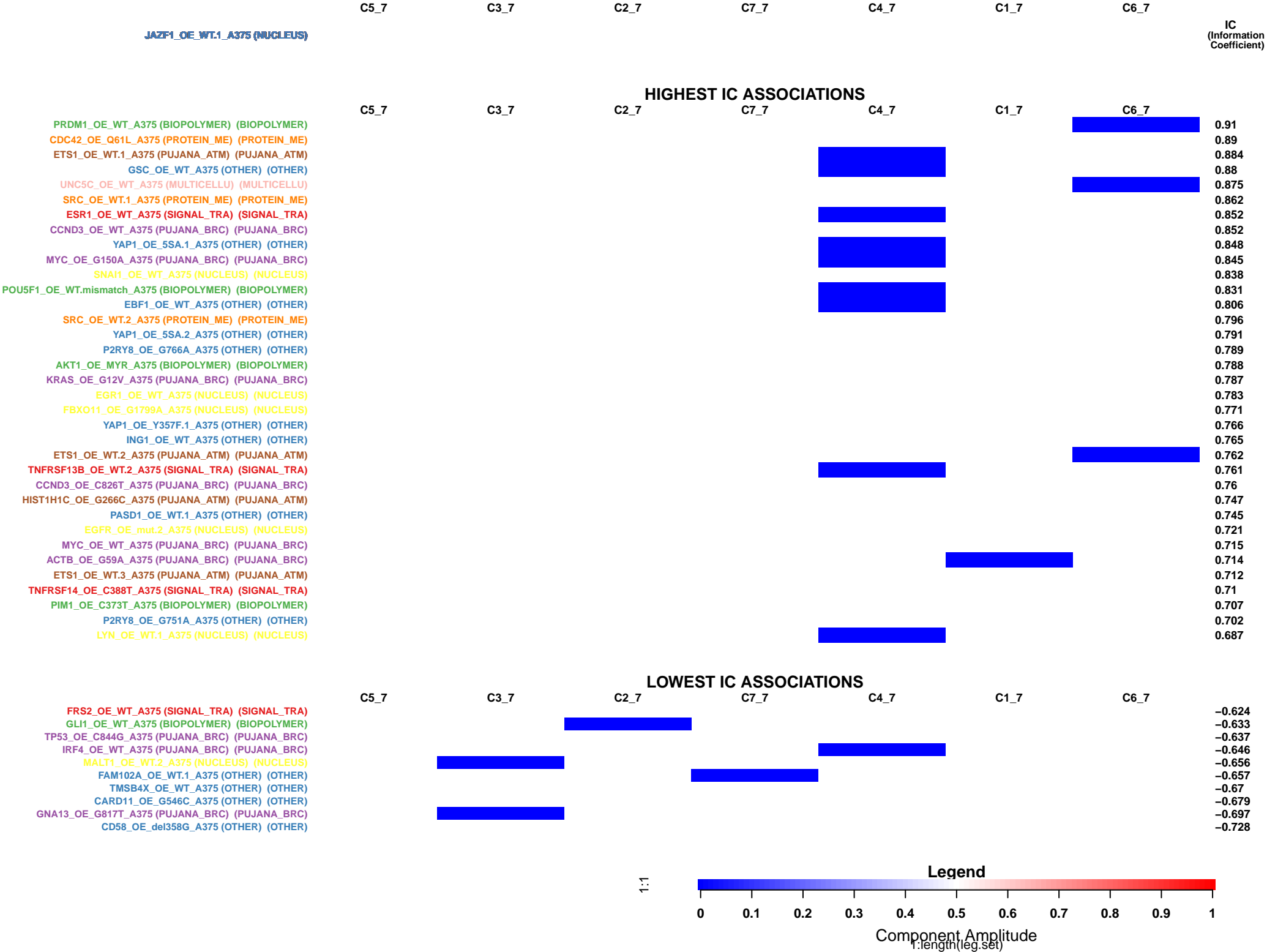
**BRD4\_OE\_WT\_A375 (OTHER)**



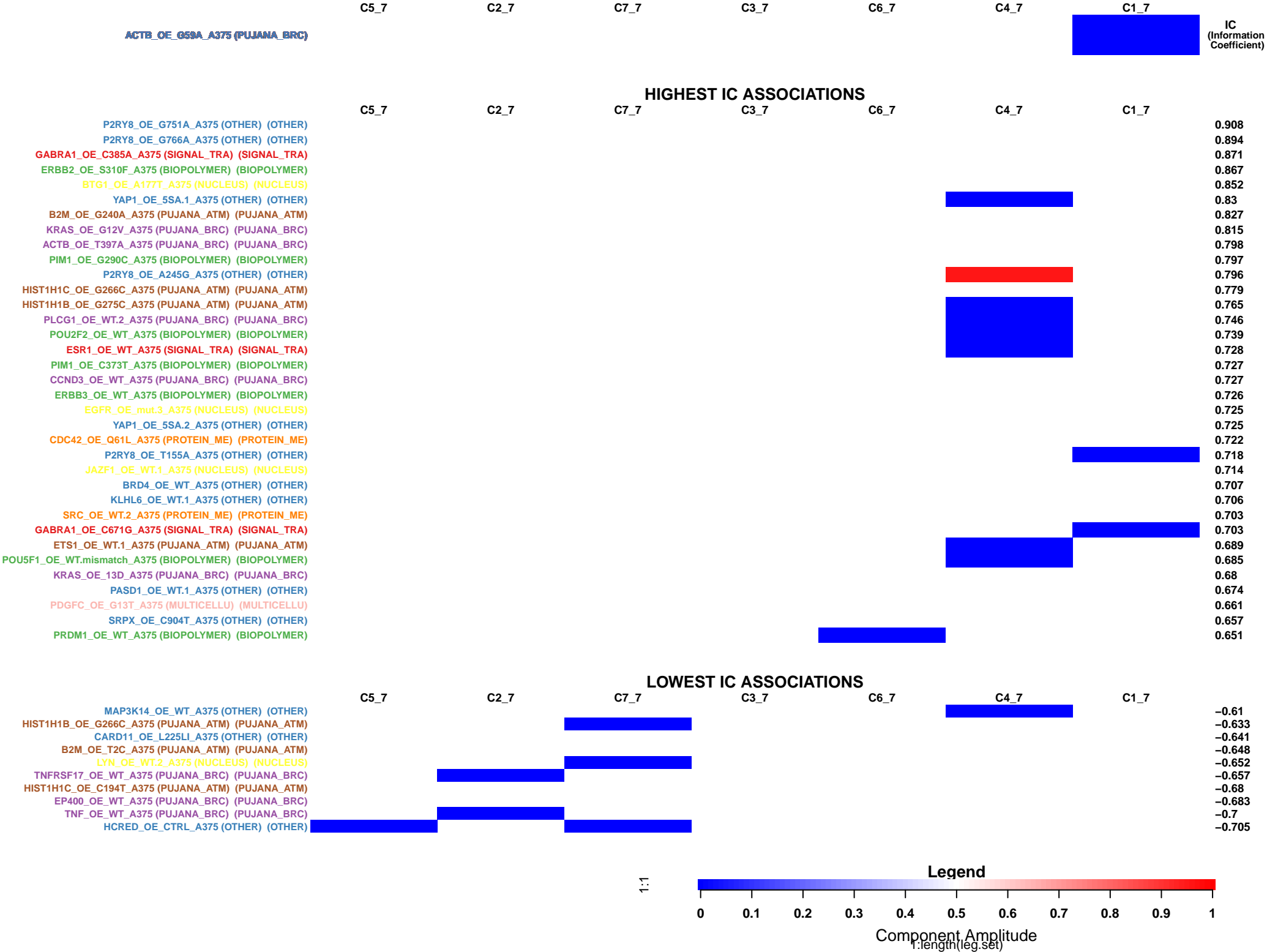
P2RY8\_OE\_G751A\_A375 (OTHER)



JAZF1\_OE\_WT.1\_A375 (NUCLEUS)



ACTB\_OE\_G59A\_A375 (PUJANA\_BRC)

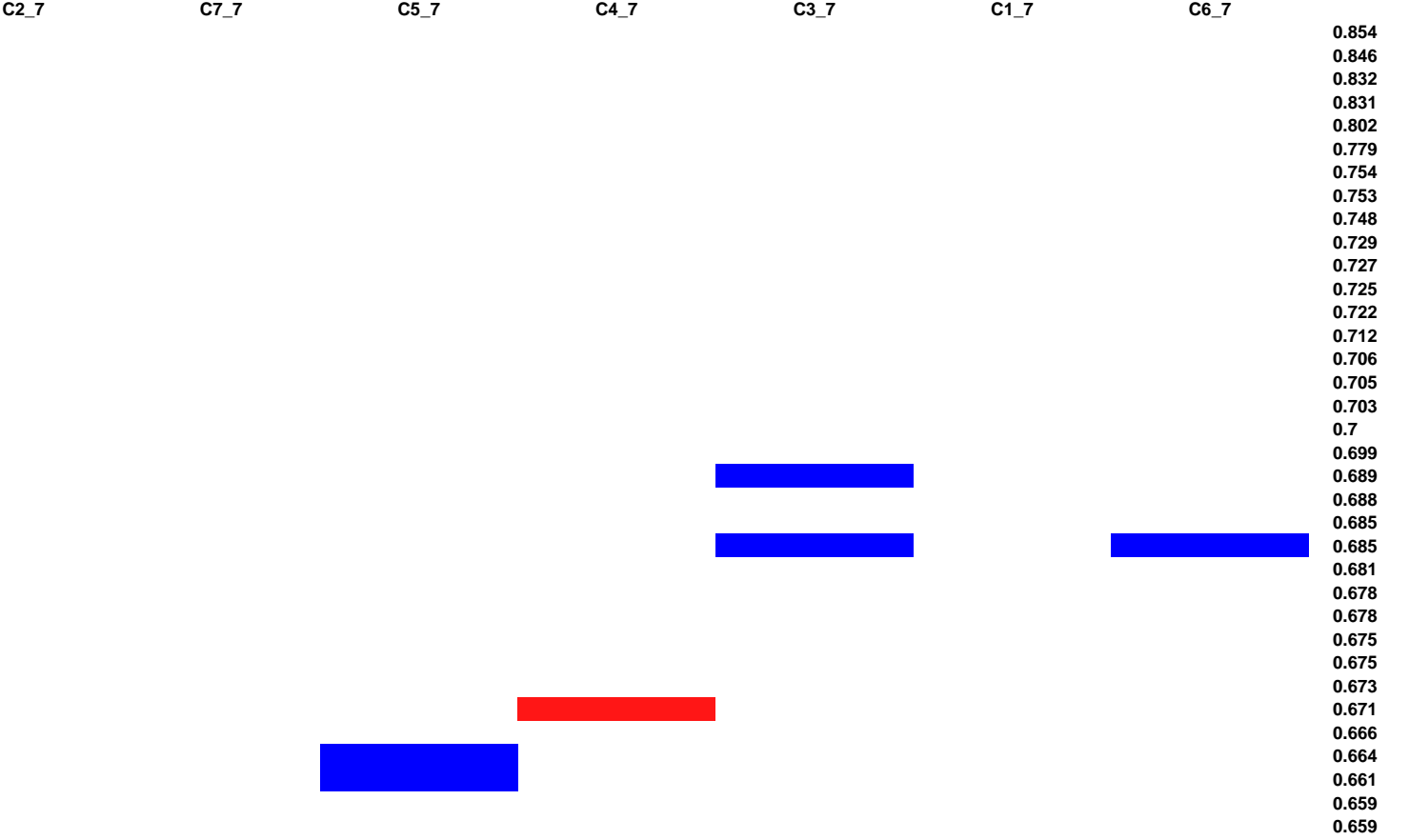


ERRFI1\_OE\_WT.1\_A375 (OTHER)

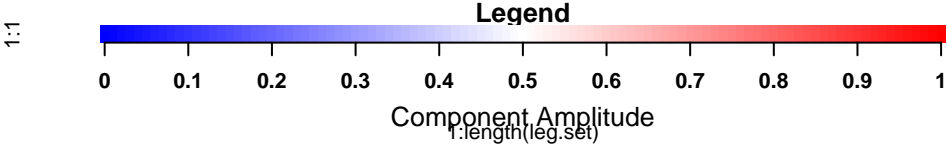
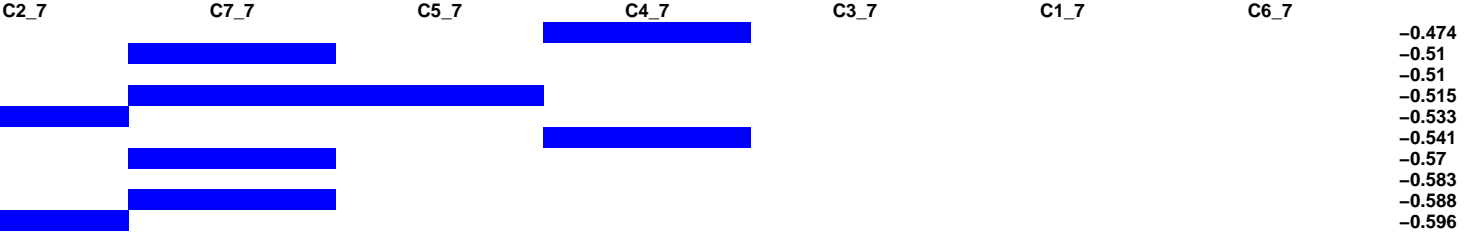
C2\_7 C7\_7 C5\_7 C4\_7 C3\_7 C1\_7 C6\_7 IC (Information Coefficient)

ERRFI1\_OE\_WT.1\_A375 (OTHER)

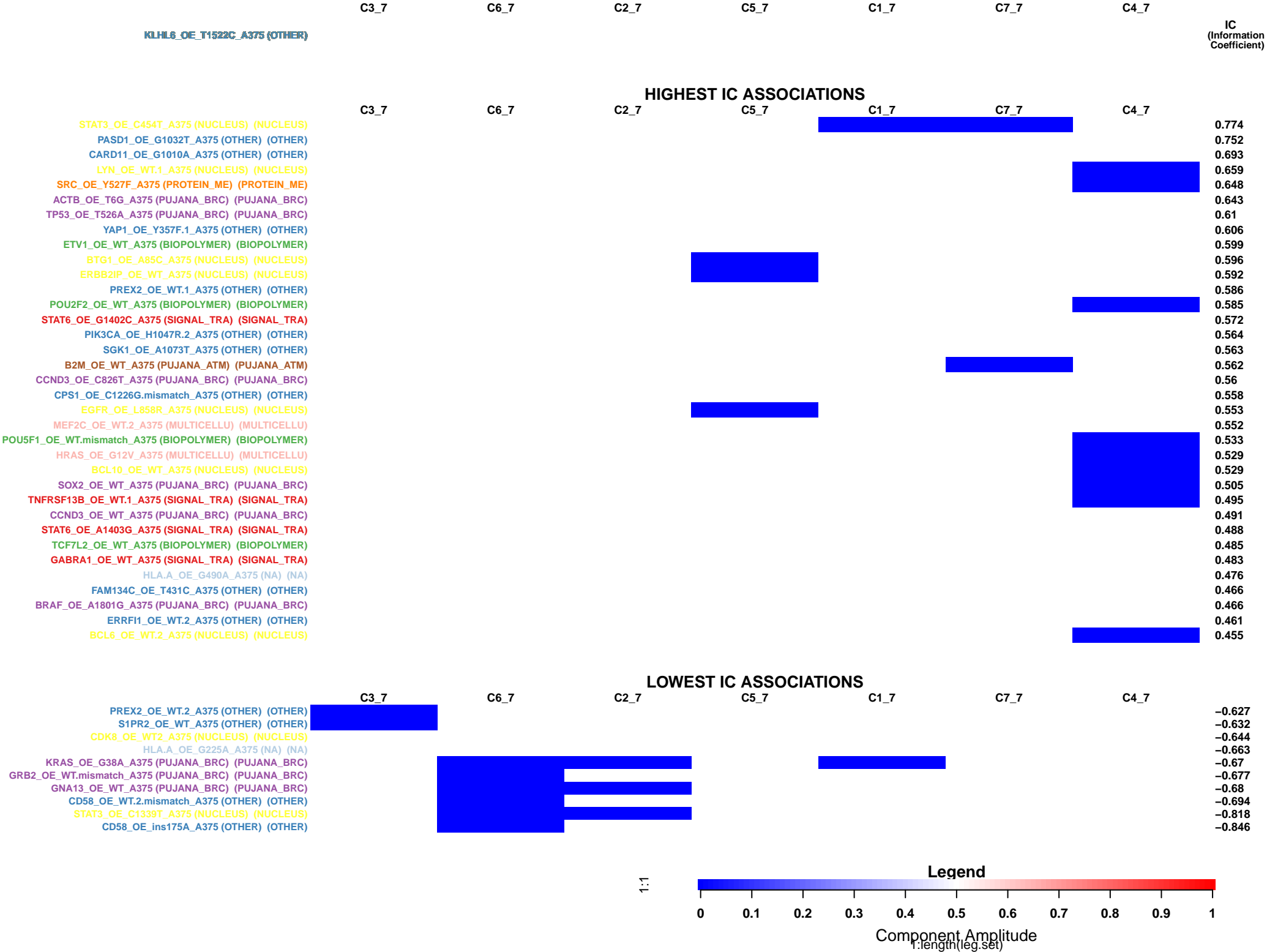
HIGHEST IC ASSOCIATIONS



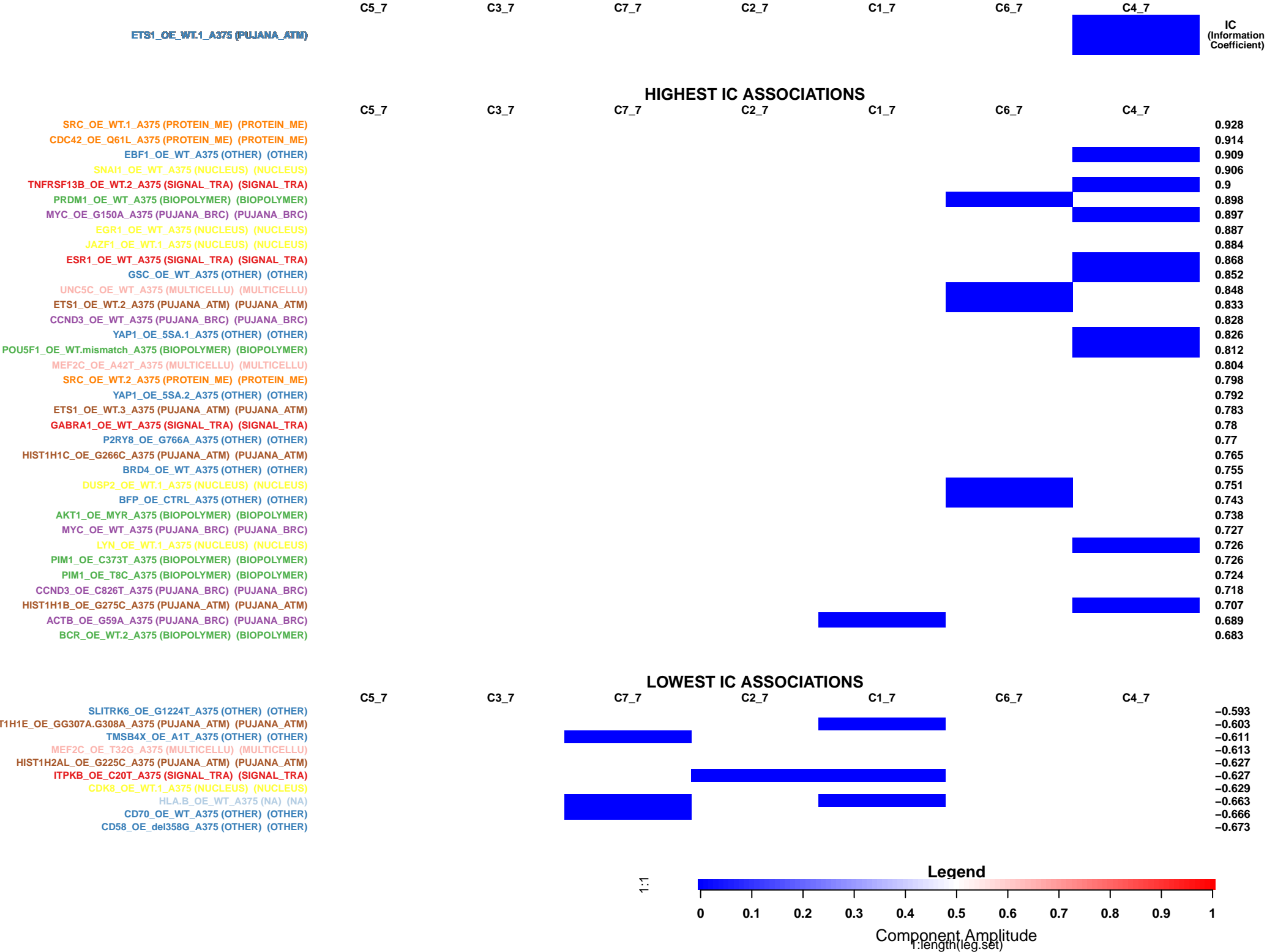
LOWEST IC ASSOCIATIONS



KLHL6\_OE\_T1522C\_A375 (OTHER)

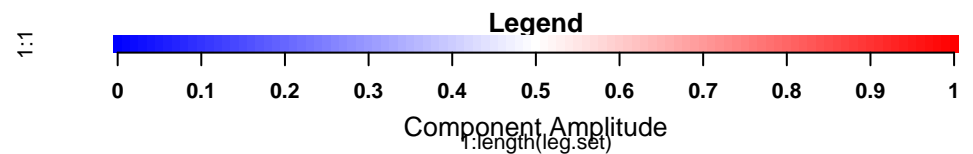
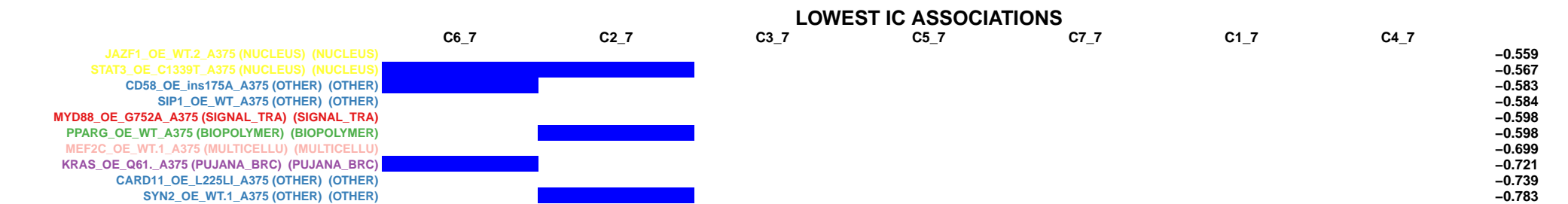
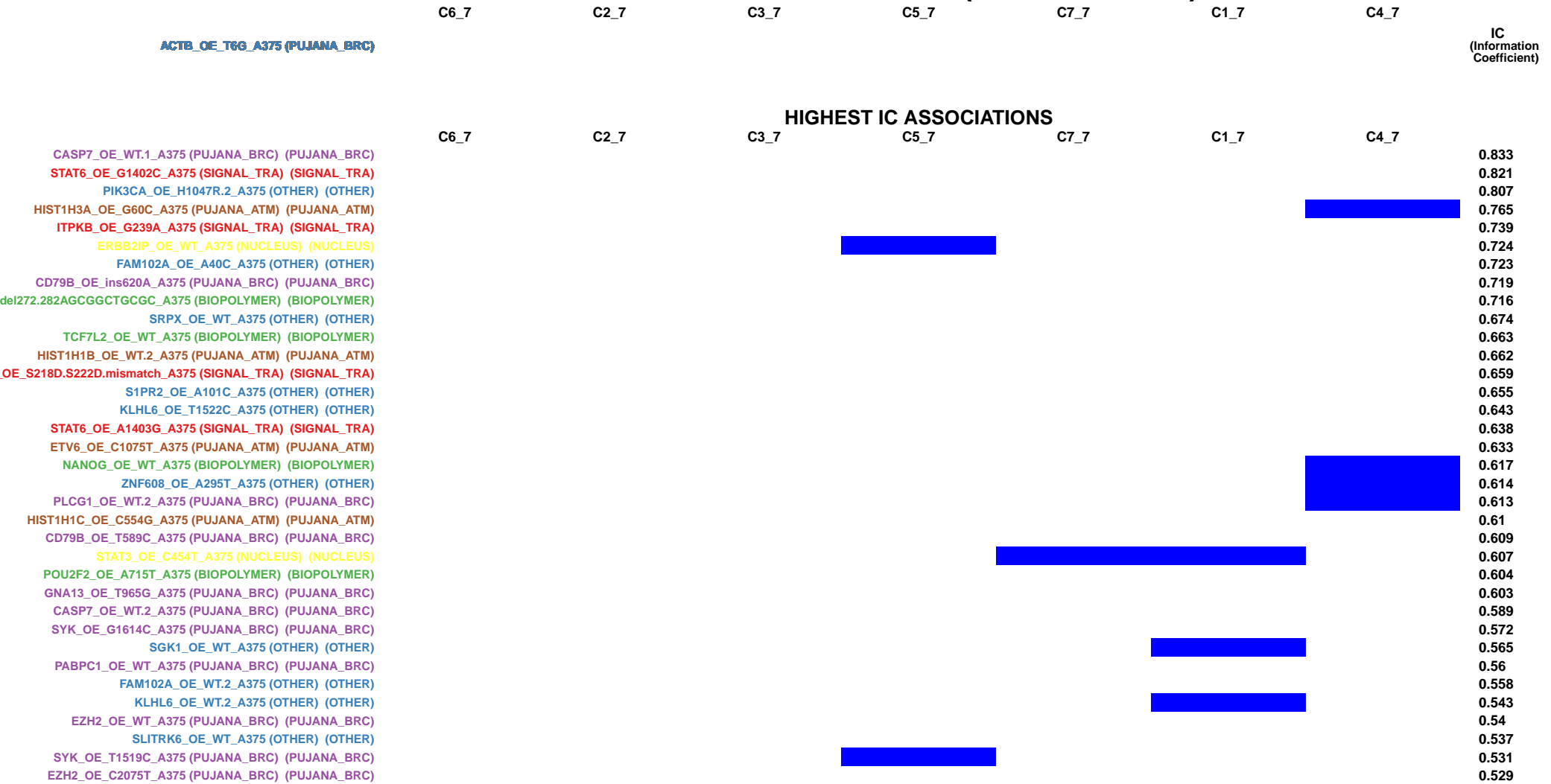


ETS1\_OE\_WT.1\_A375 (PUJANA\_ATM)

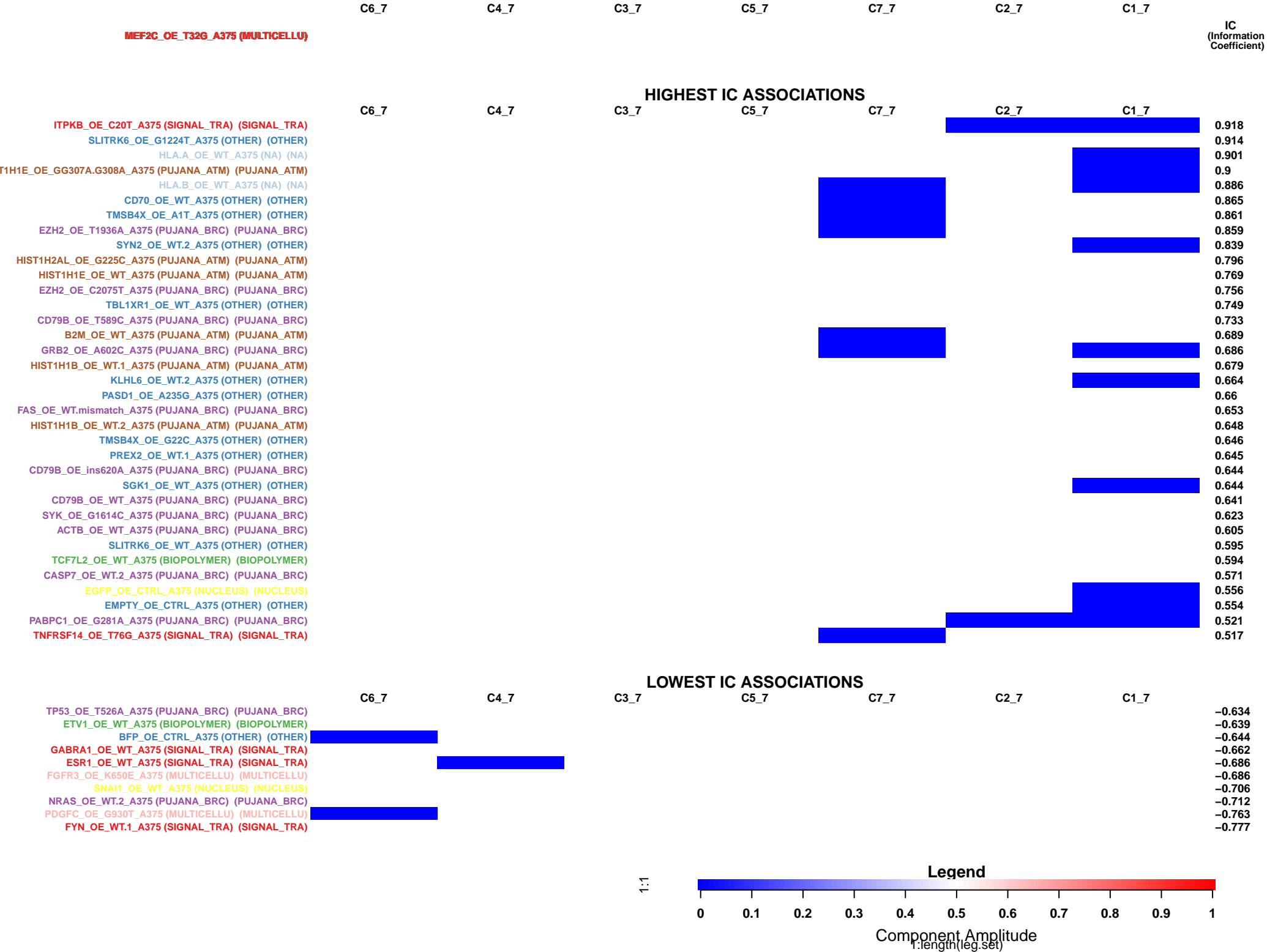




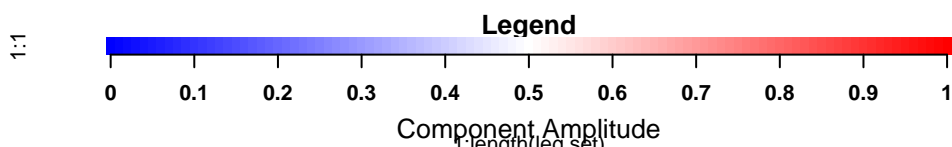
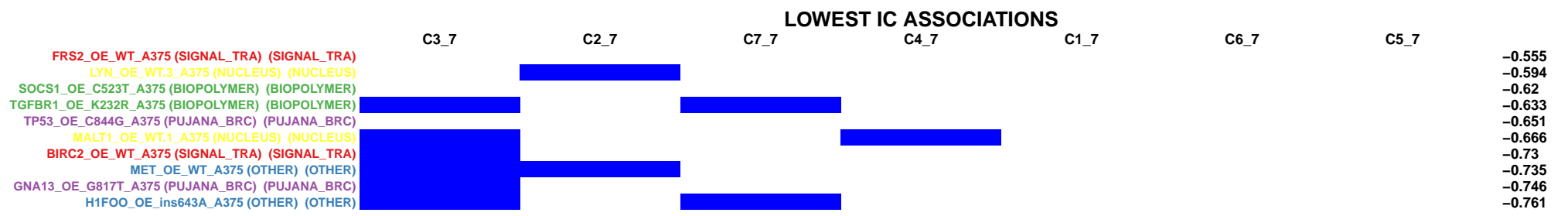
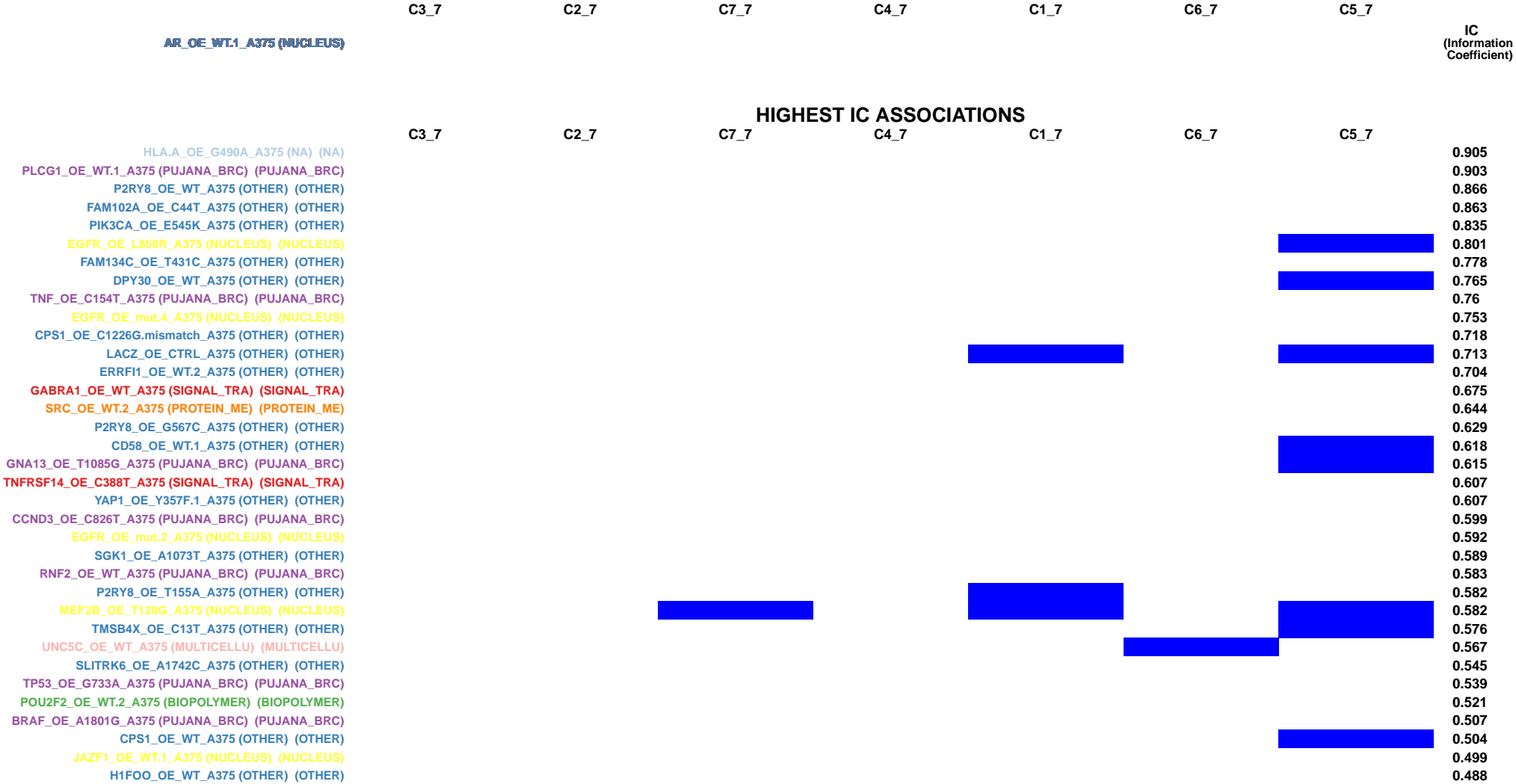
# ACTB\_OE\_T6G\_A375 (PUJANA\_BRC)



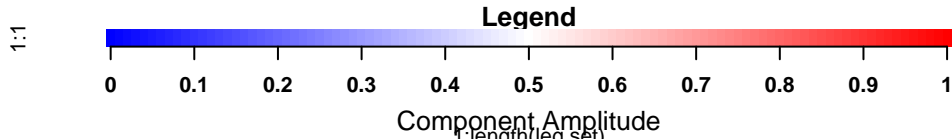
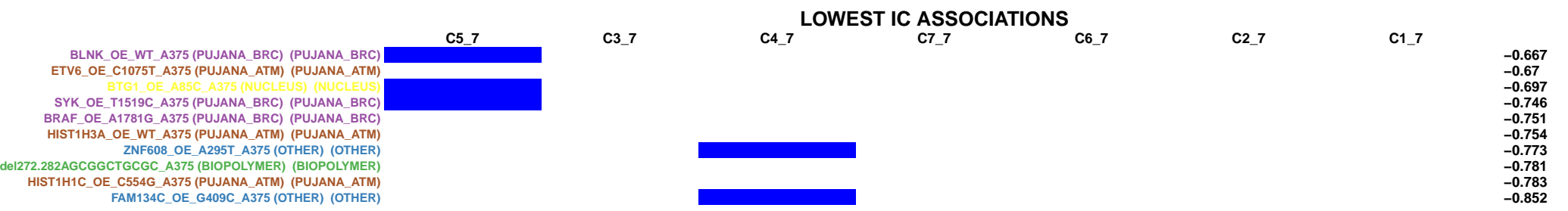
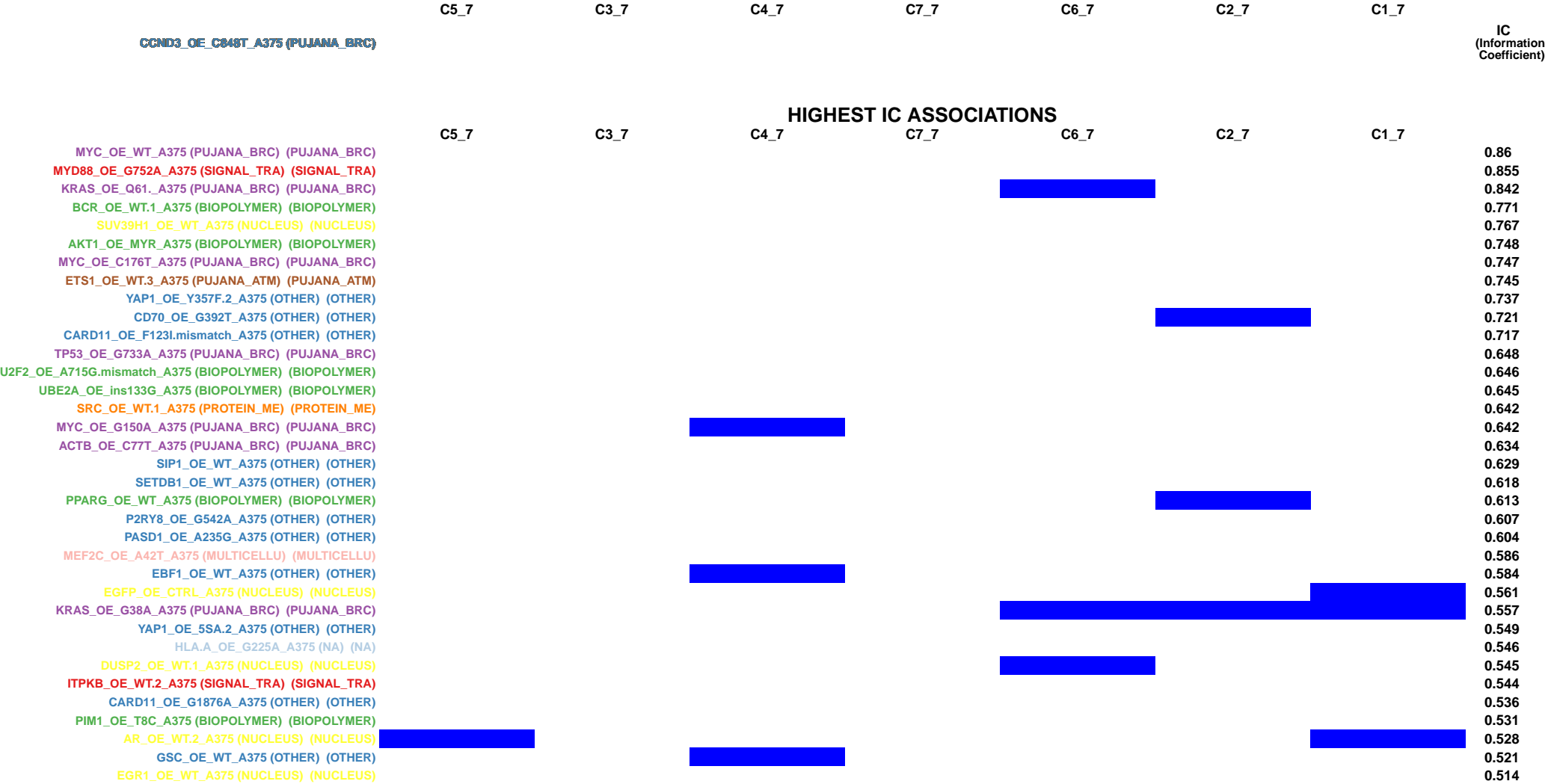
MEF2C\_OE\_T32G\_A375 (MULTICELLU)



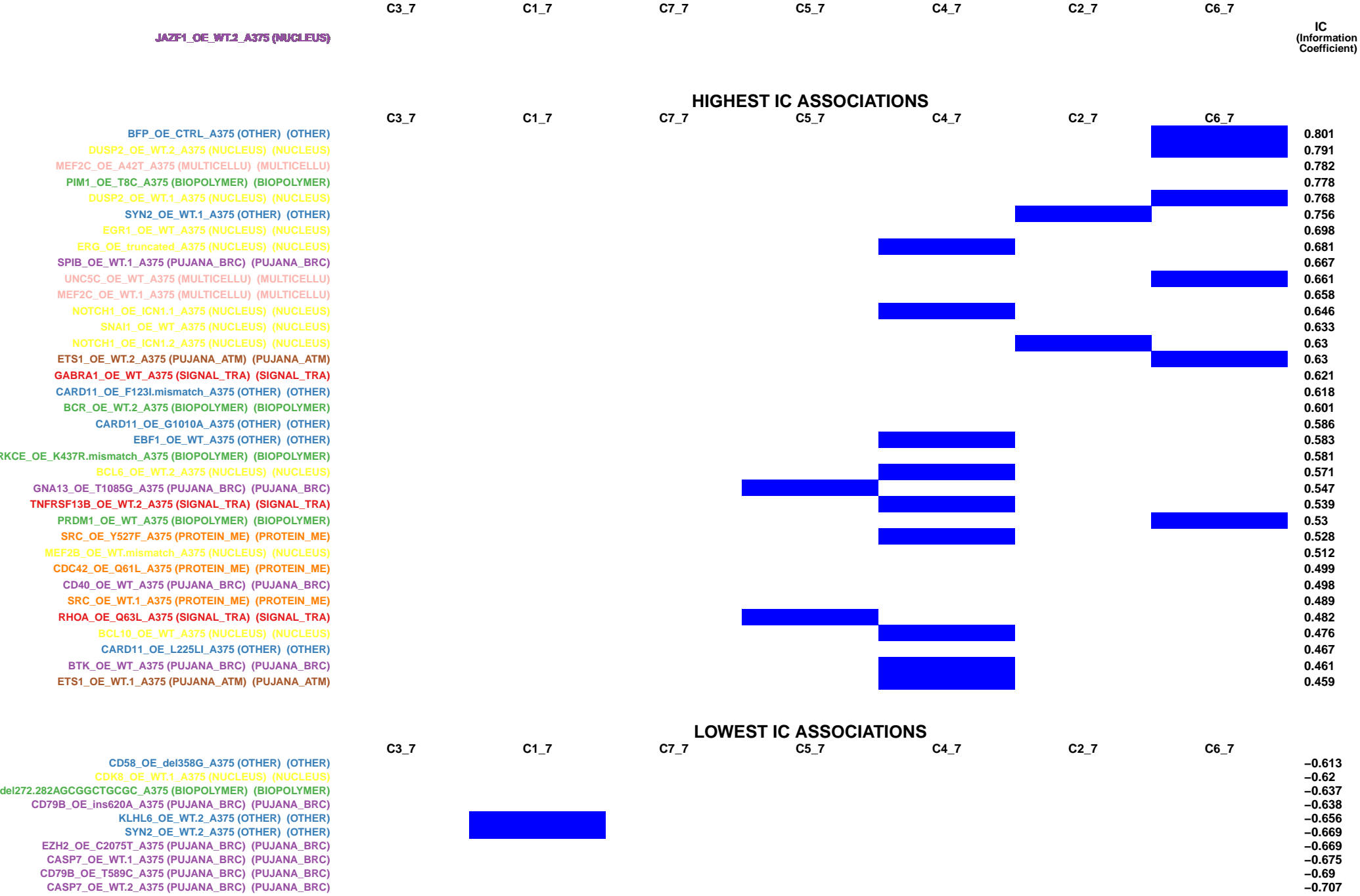
AR\_OE\_WT.1\_A375 (NUCLEUS)



CCND3\_OE\_C848T\_A375 (PUJANA\_BRC)



JAZF1\_OE\_WT.2\_A375 (NUCLEUS)



C6_7	C5_7	C3_7	C4_7	C1_7	C7_7	C2_7
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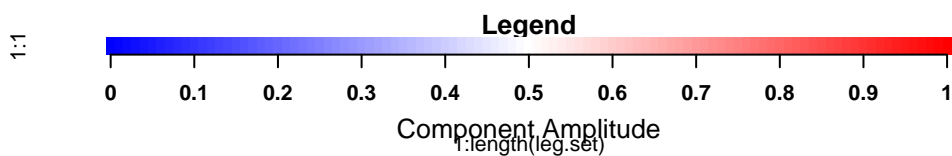
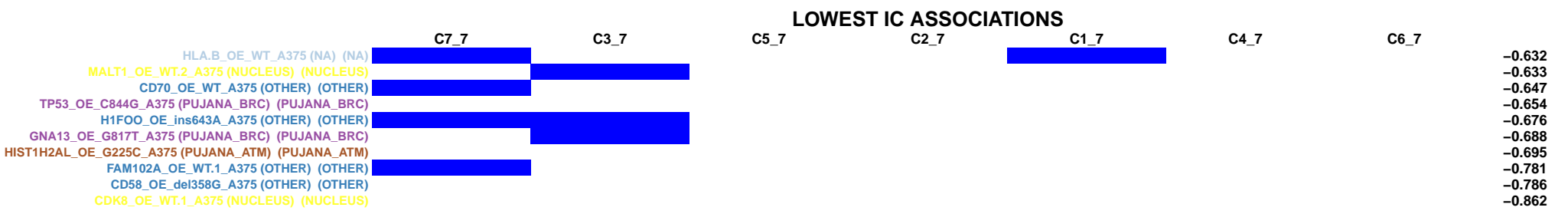
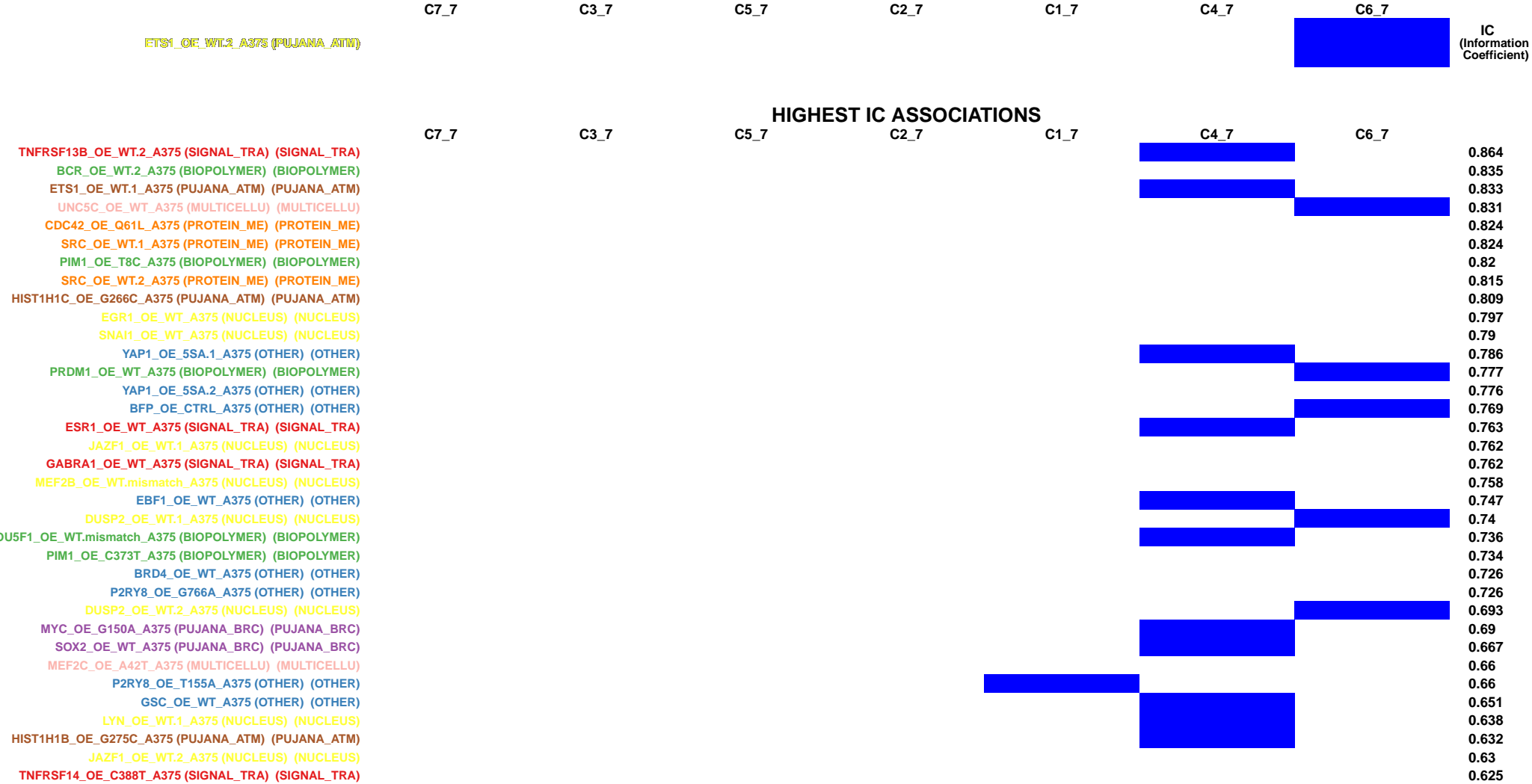
IC  
Information  
Coefficient)

C6\_7                      C5\_7                      C3\_7                      C4\_7                      C1\_7                      C7\_7                      C2\_7

Component Amplitude  
1:length(leg.set)

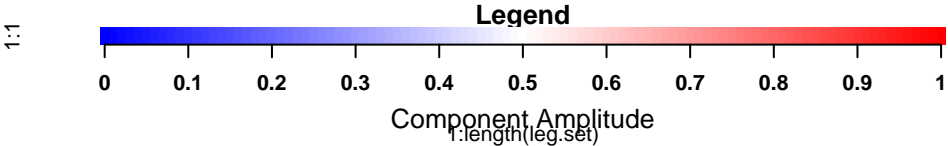
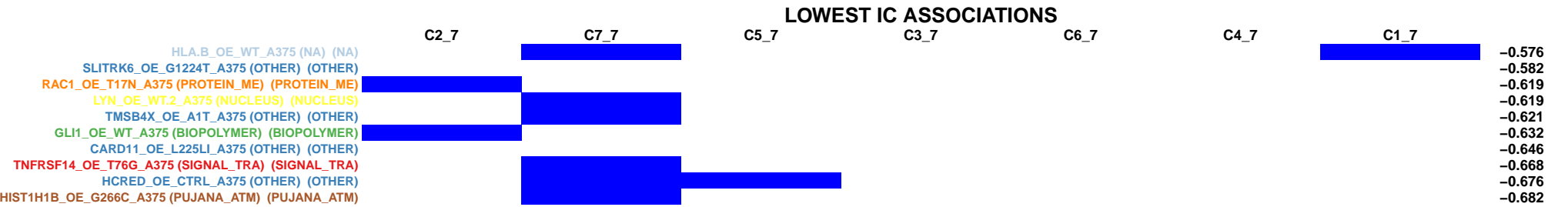
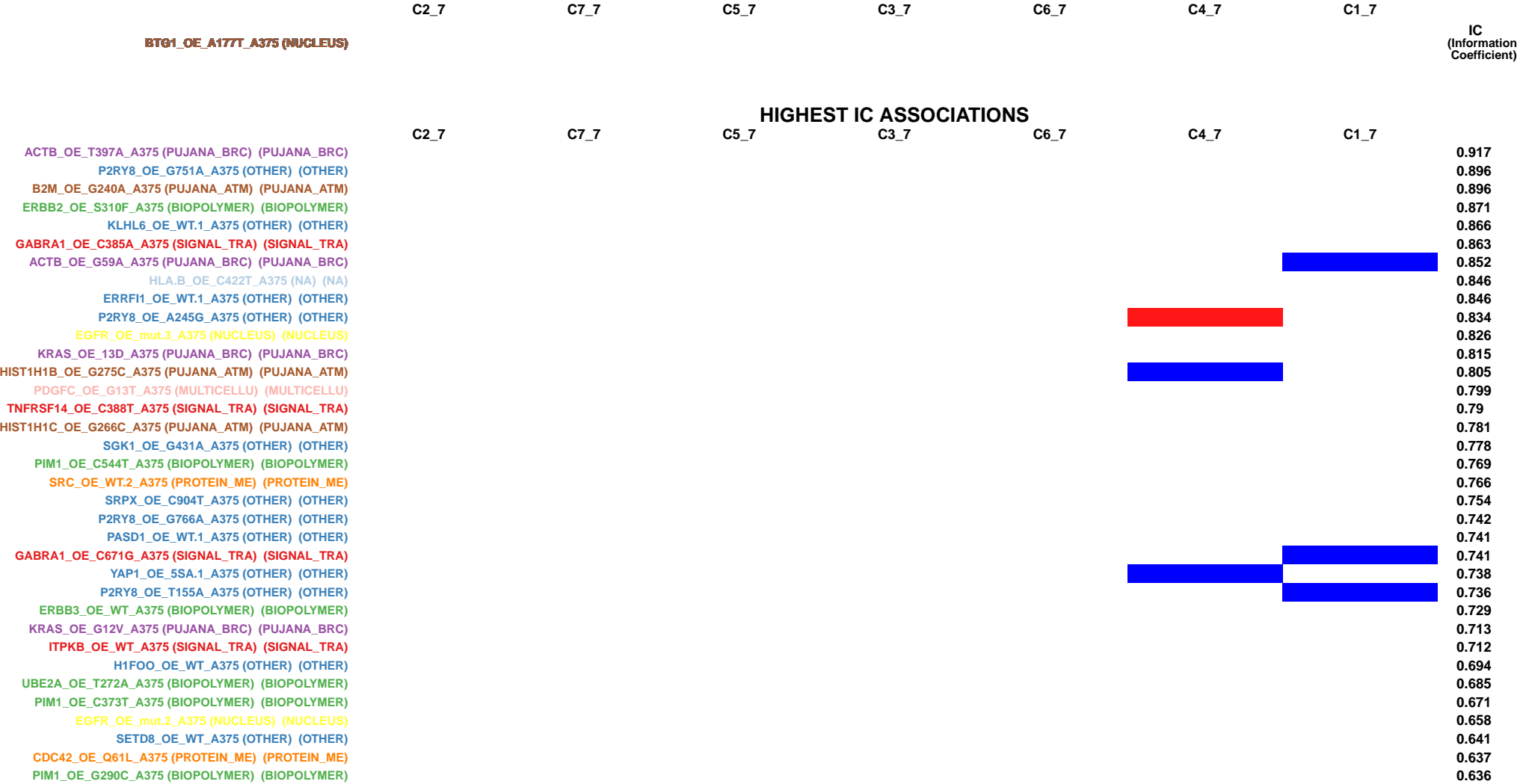
```
1:length(leg.set)
```

ETS1\_OE\_WT.2\_A375 (PUJANA\_ATM)



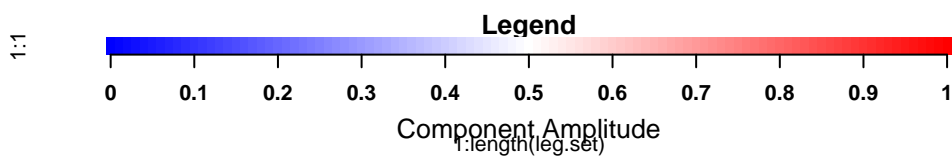
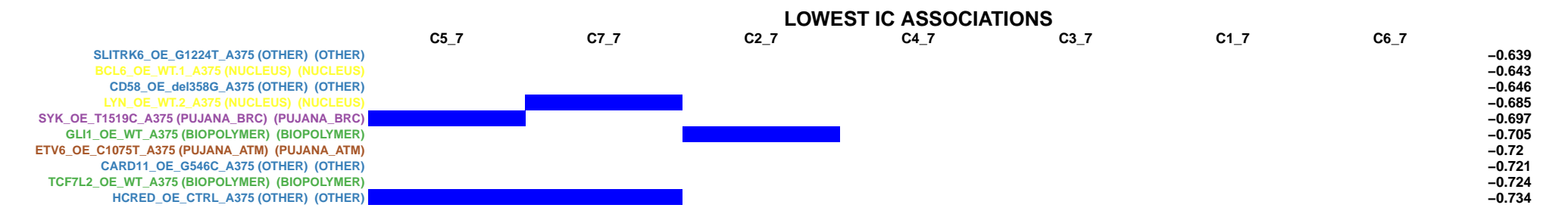
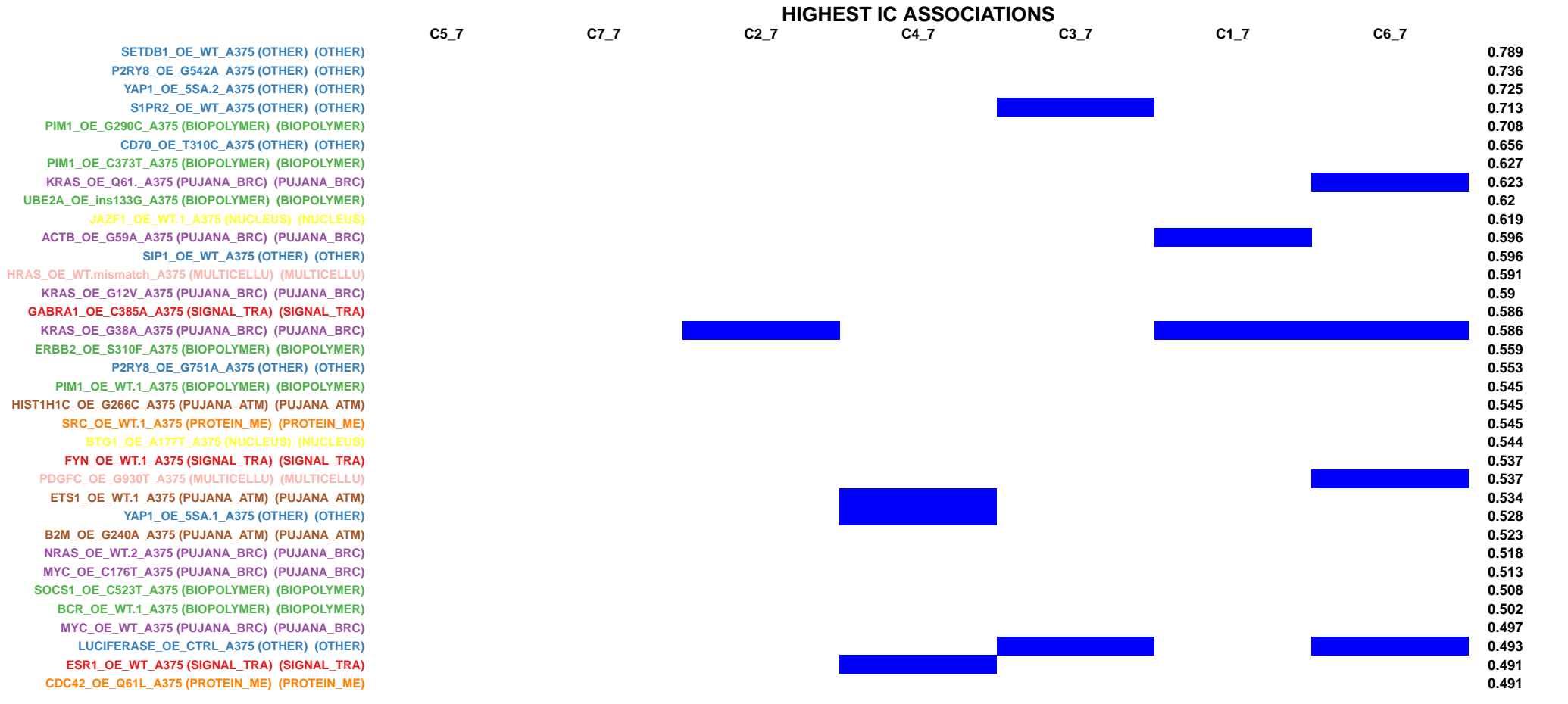
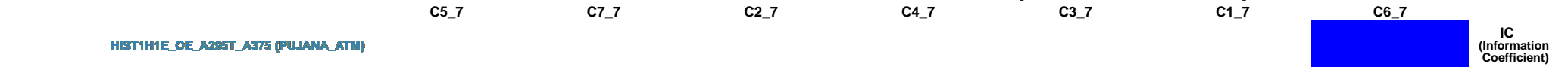
BTG1\_OE\_A177T\_A375 (NUCLEUS)

BTG1\_OE\_A177T\_A375 (NUCLEUS)

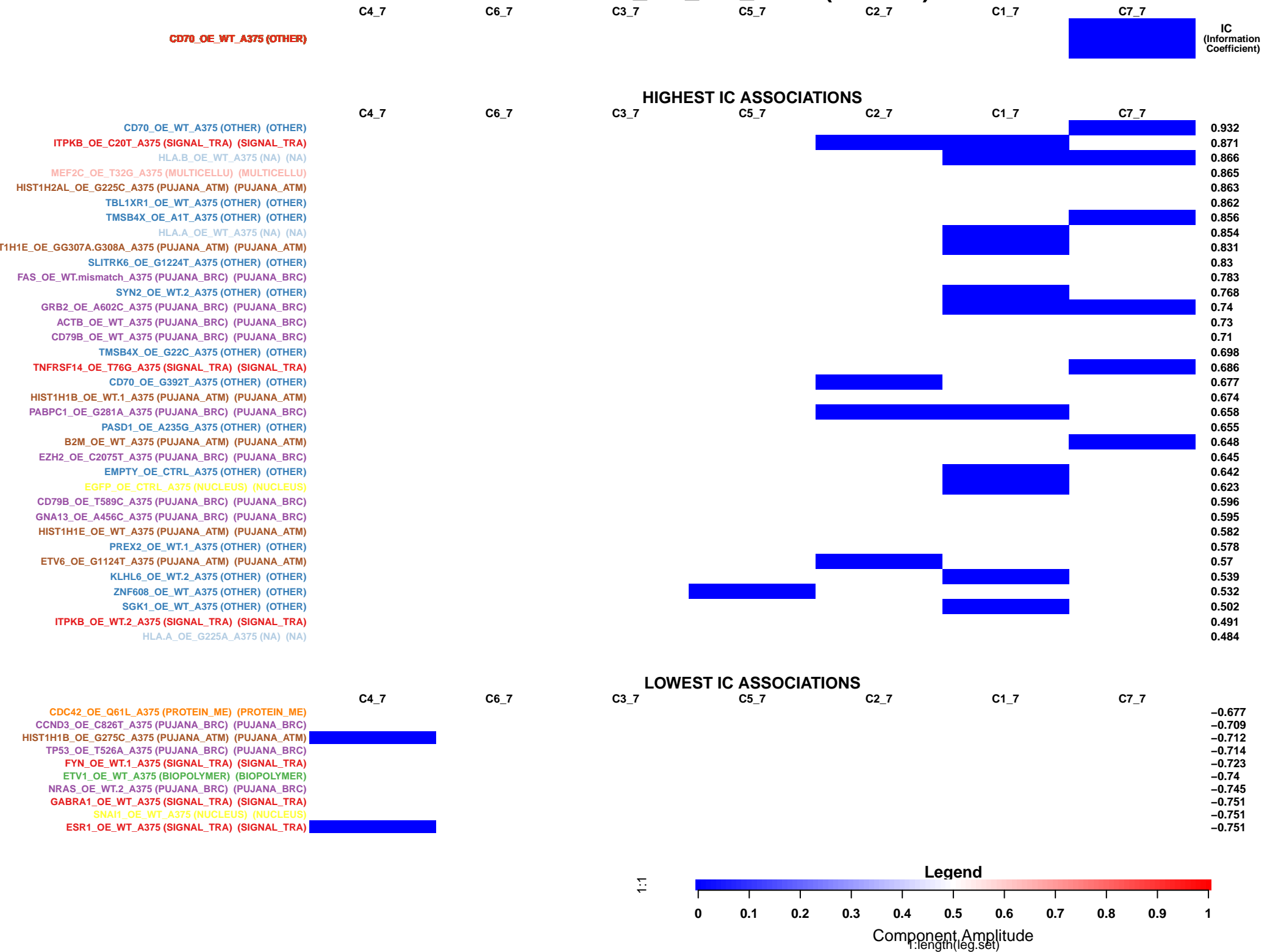




HIST1H1E\_OE\_A295T\_A375 (PUJANA\_ATM)

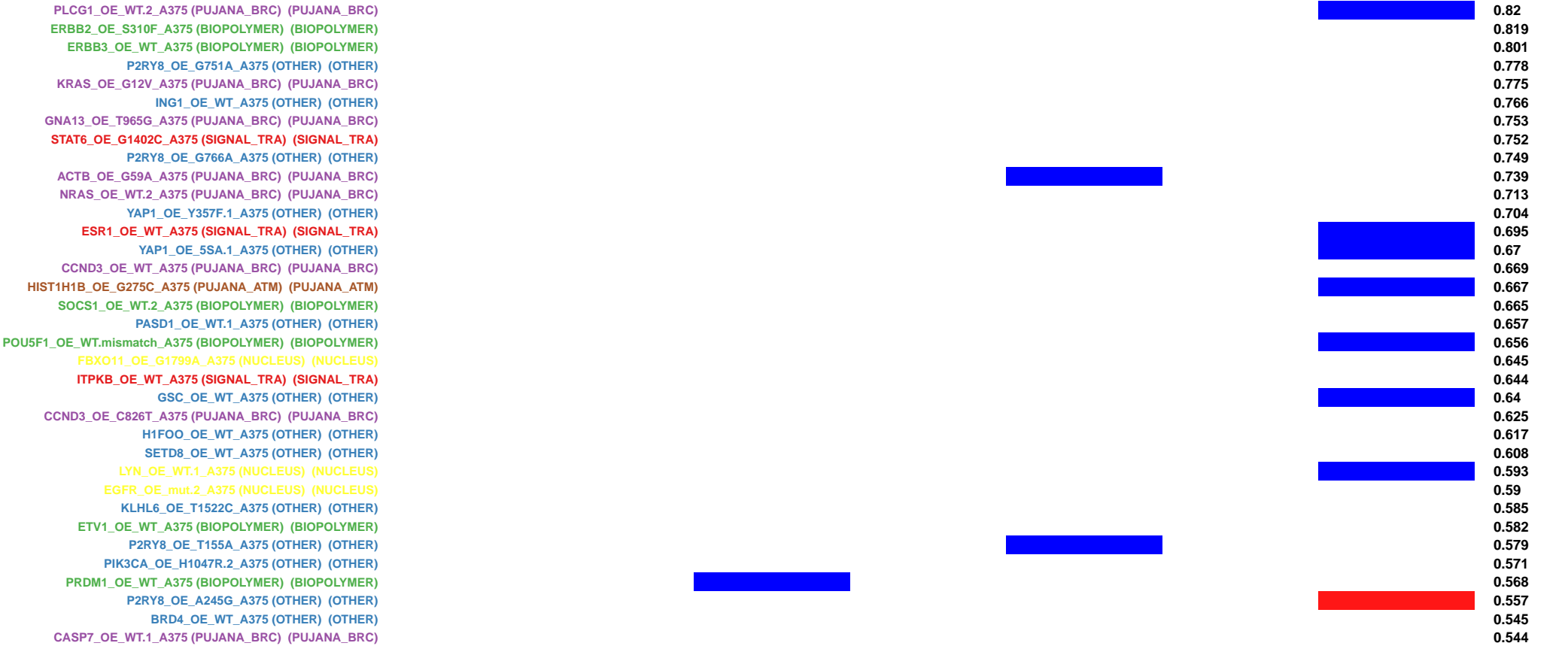


**CD70\_OE\_WT\_A375 (OTHER)**



C2_7	C5_7	C6_7	C3_7	C1_7	C7_7	C4_7	IC (Information Coefficient)
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C2\_7                      C5\_7                      C6\_7                      C3\_7                      C1\_7                      C7\_7                      C4\_7

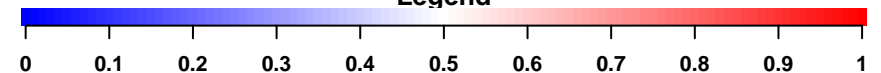


C2\_7                      C5\_7                      C6\_7                      C3\_7                      C1\_7                      C7\_7                      C4\_7



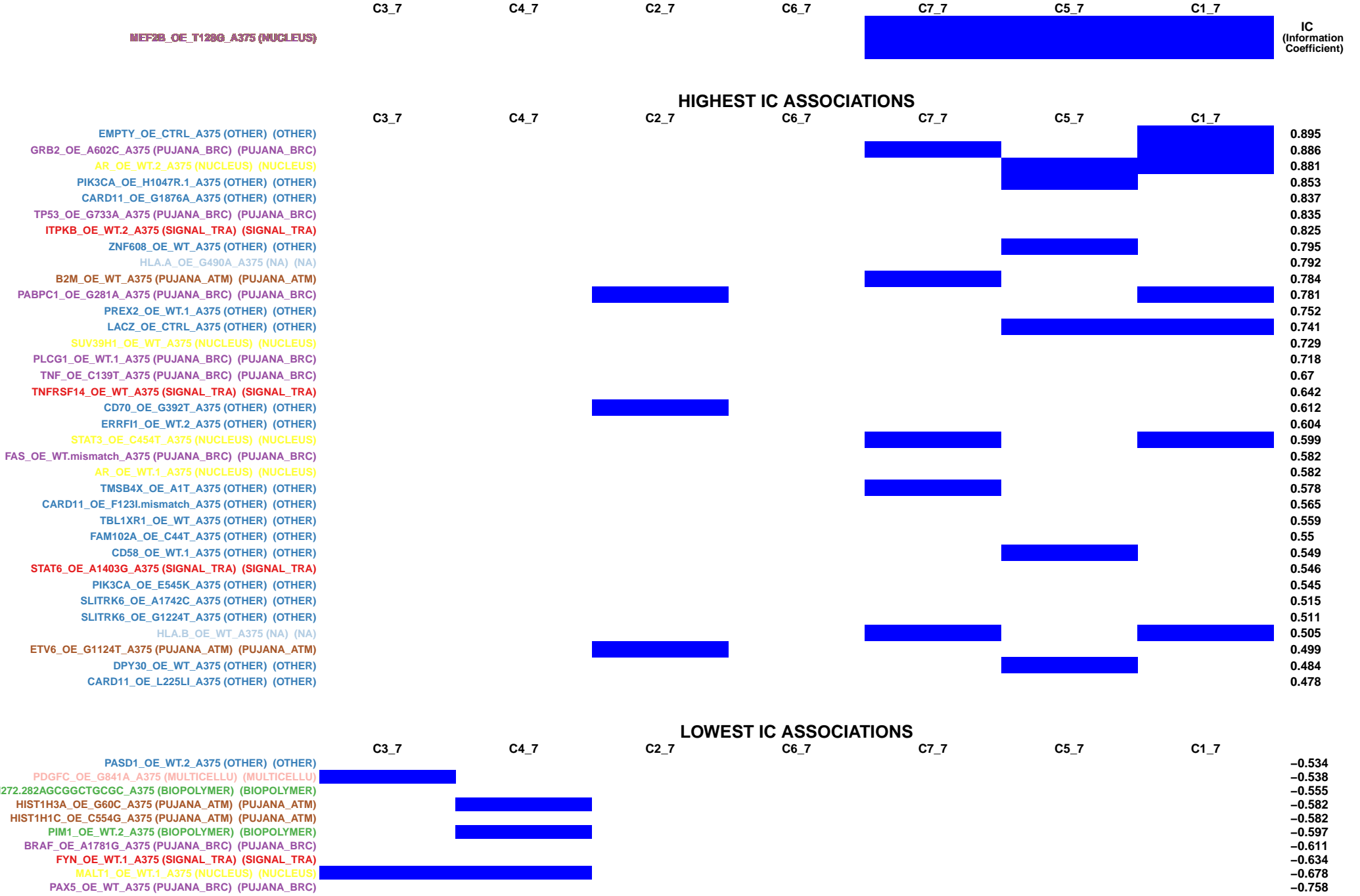
1:1

### Legend

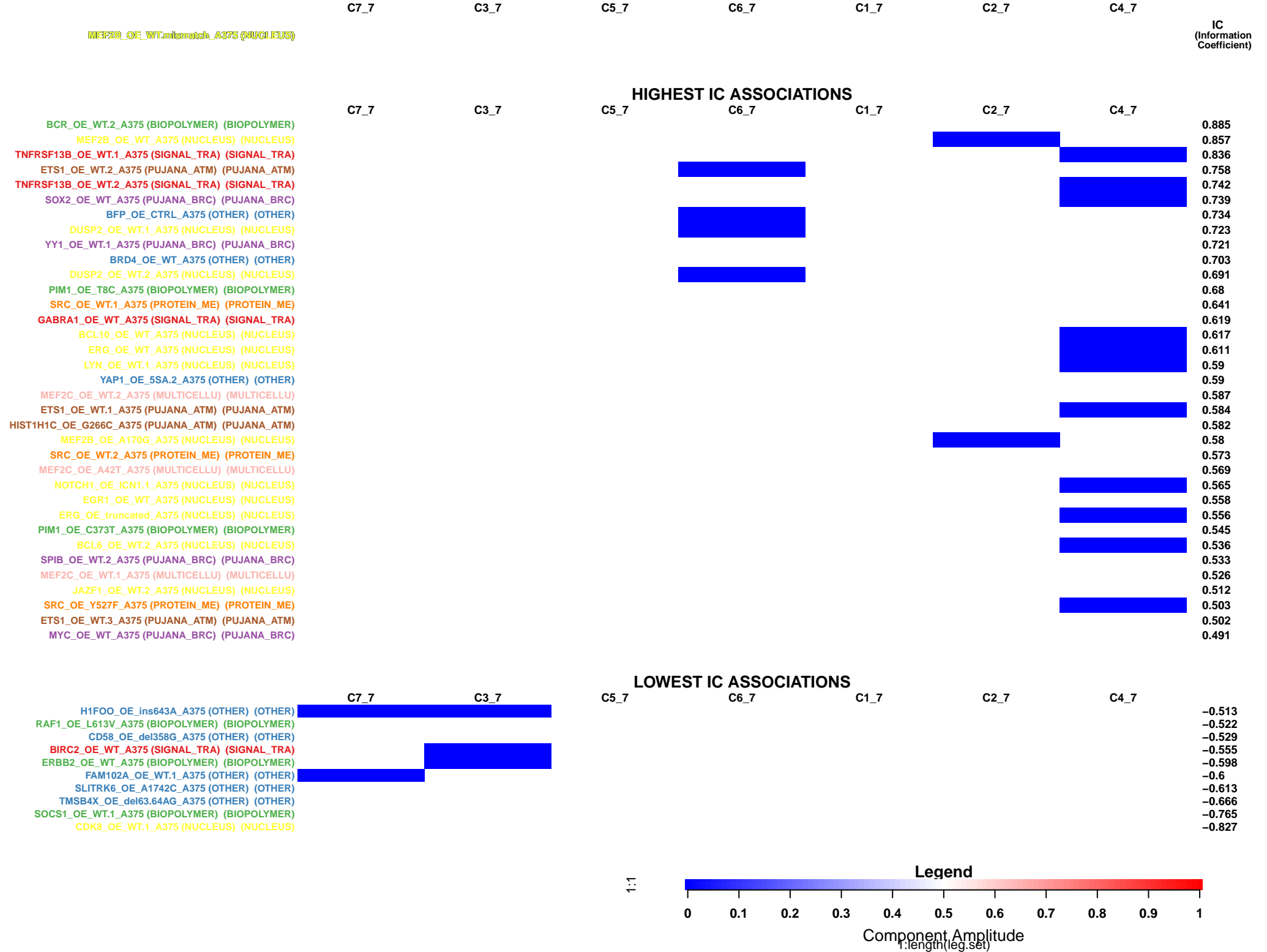
Component Amplitude  
 $1/\text{length}(\log \text{ cct})$ 

```
1:length(leg.set)
```

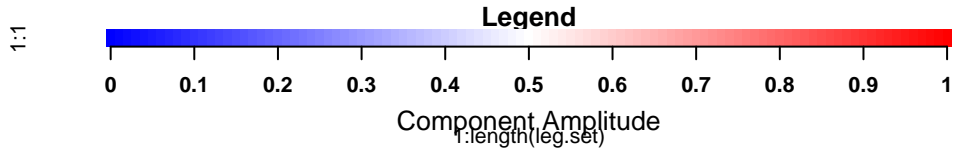
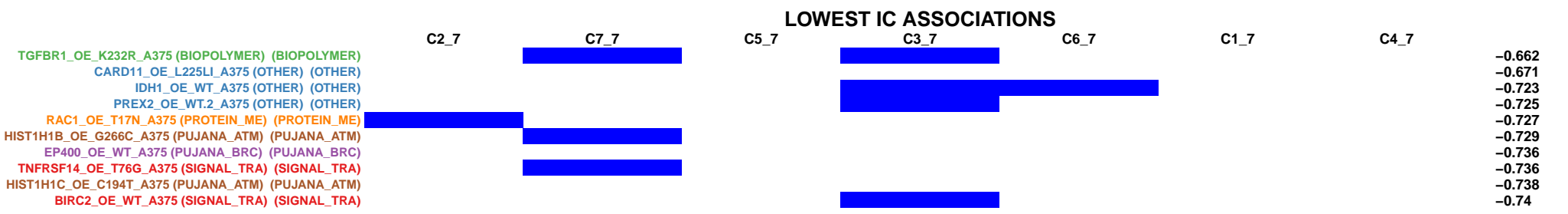
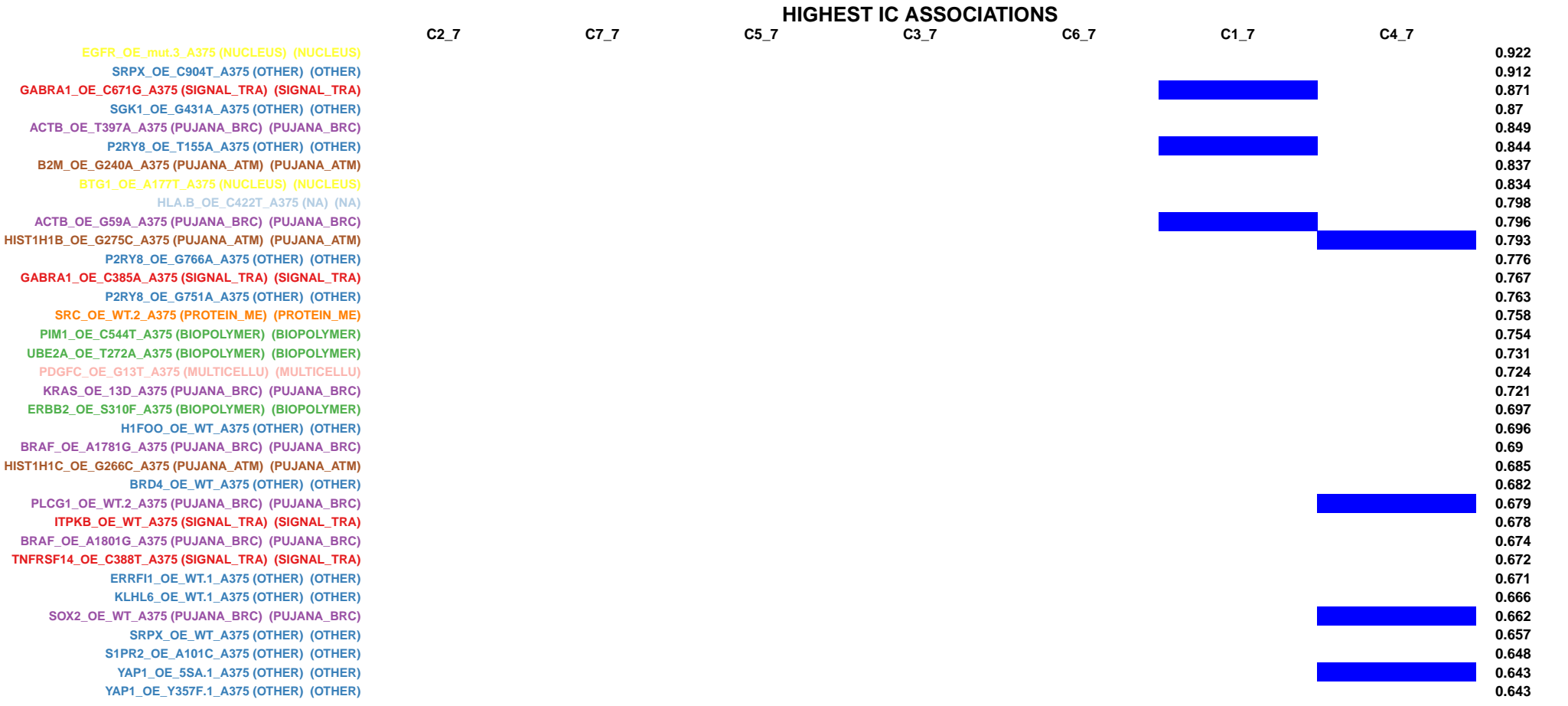
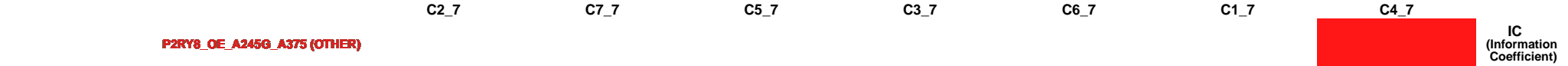
MEF2B\_OE\_T128G\_A375 (NUCLEUS)



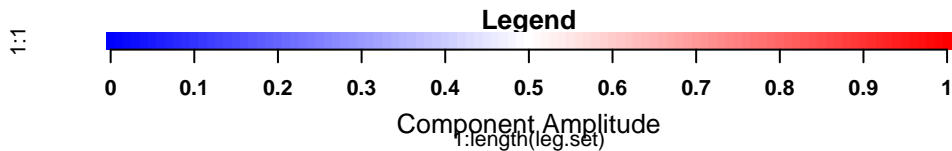
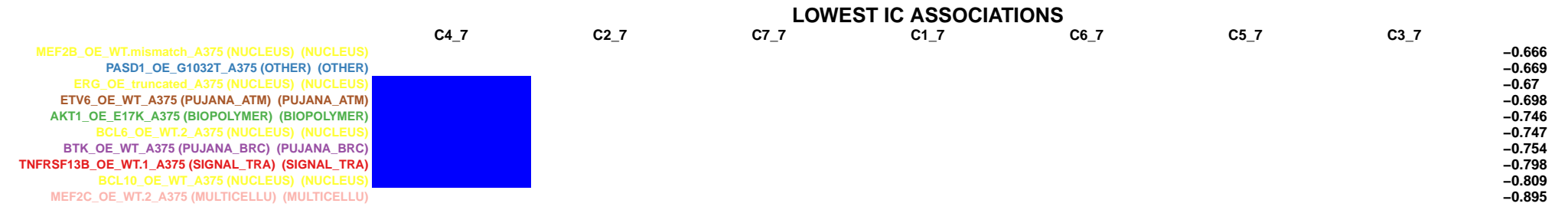
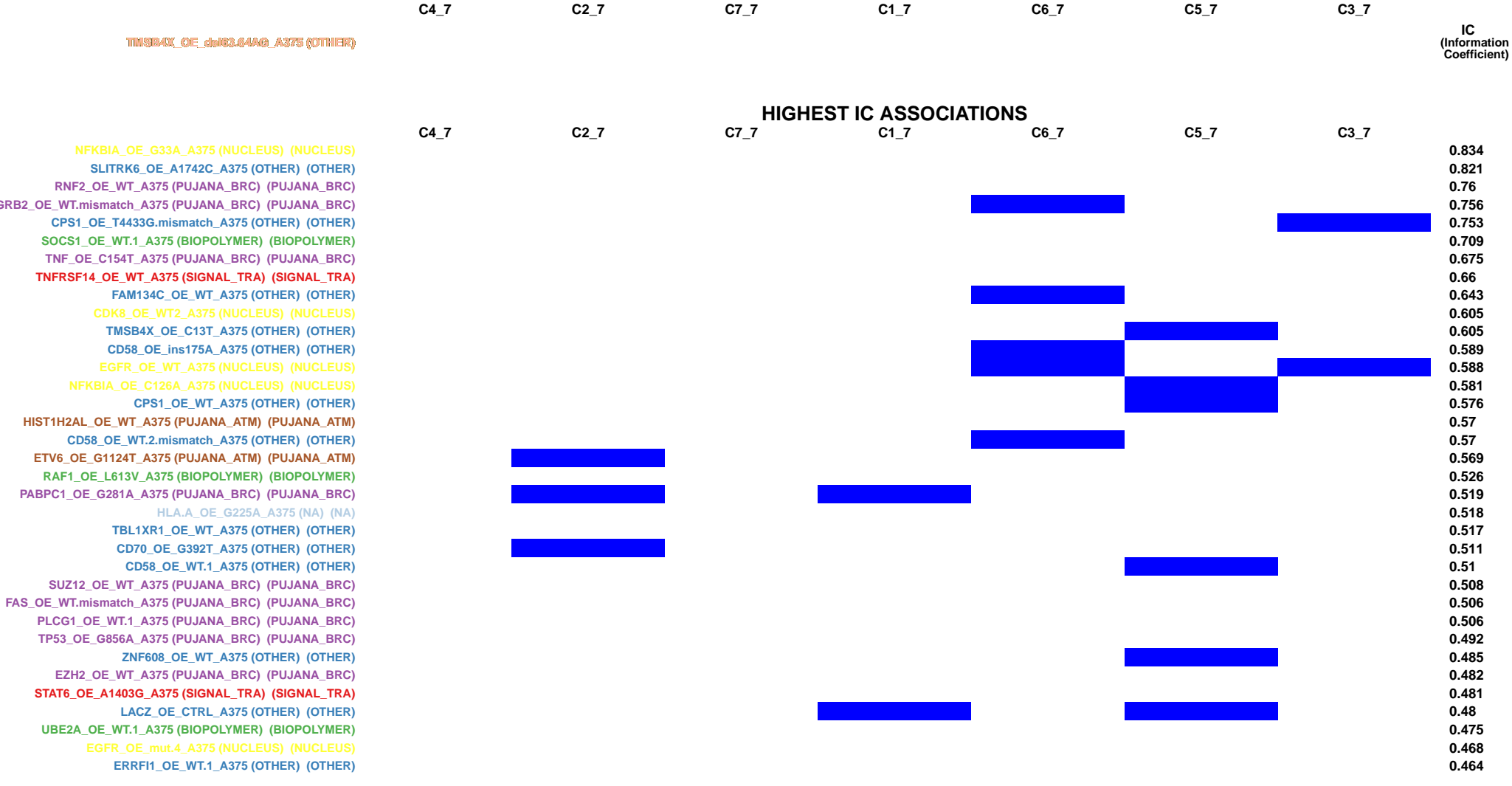
MEF2B\_OE\_WT.mismatch\_A375 (NUCLEUS)



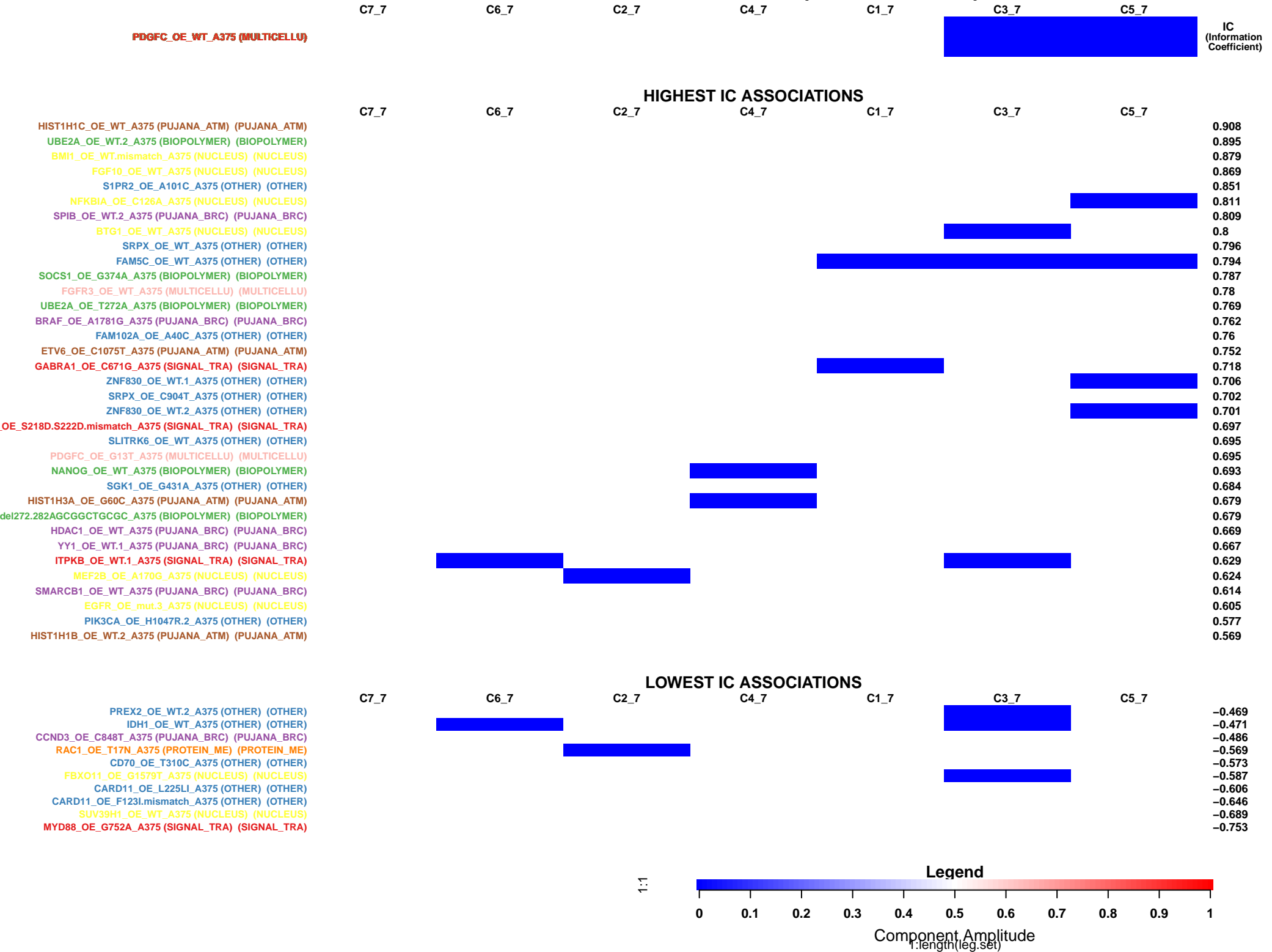
P2RY8\_OE\_A245G\_A375 (OTHER)



TMSB4X\_OE\_del63.64AG\_A375 (OTHER)

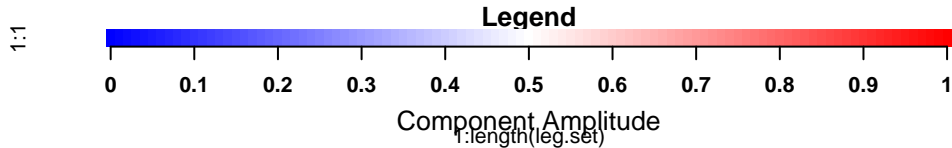
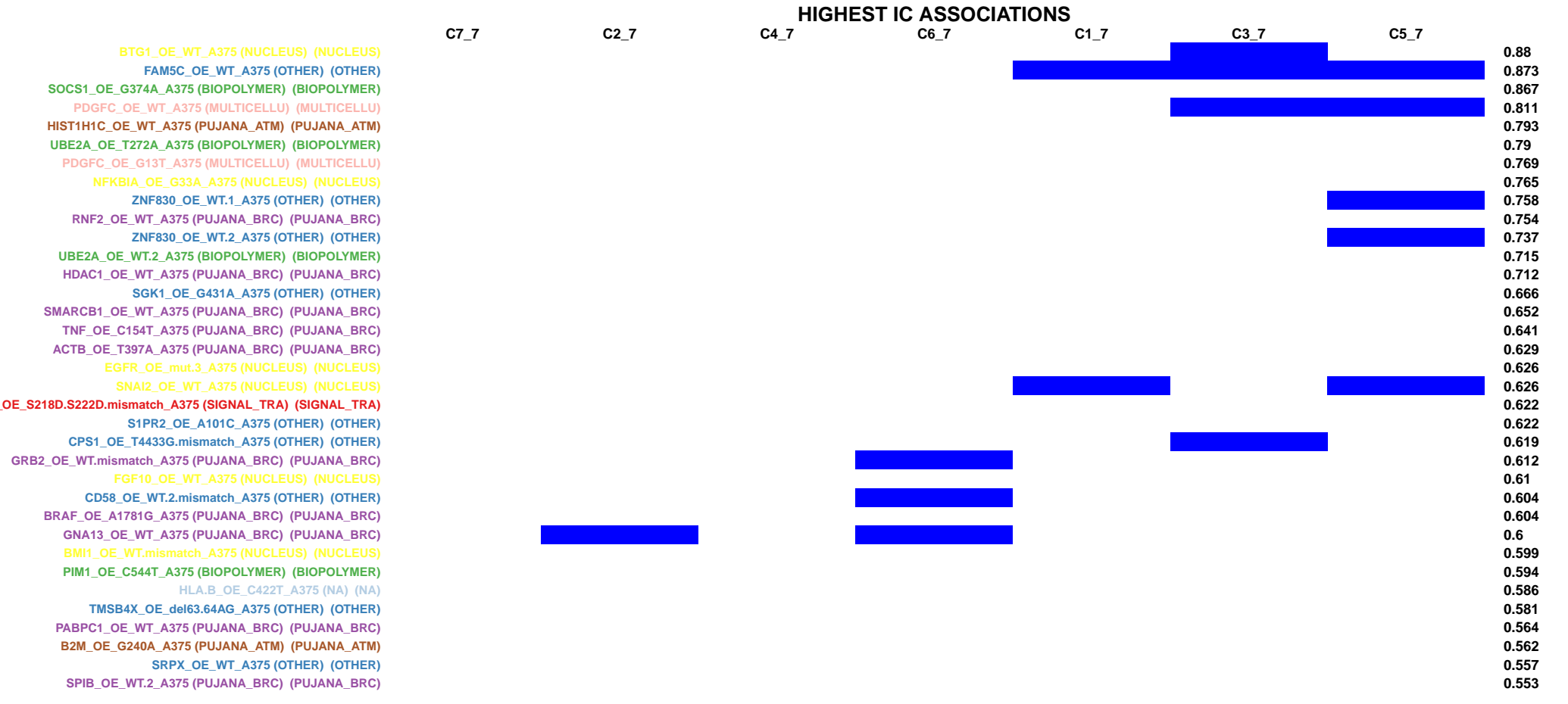
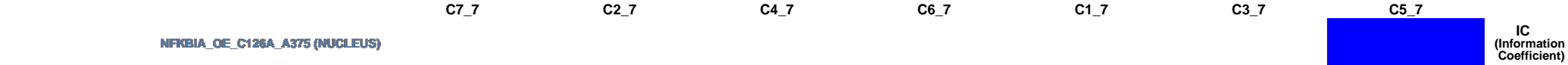


# PDGFC\_OE\_WT\_A375 (MULTICELLU)

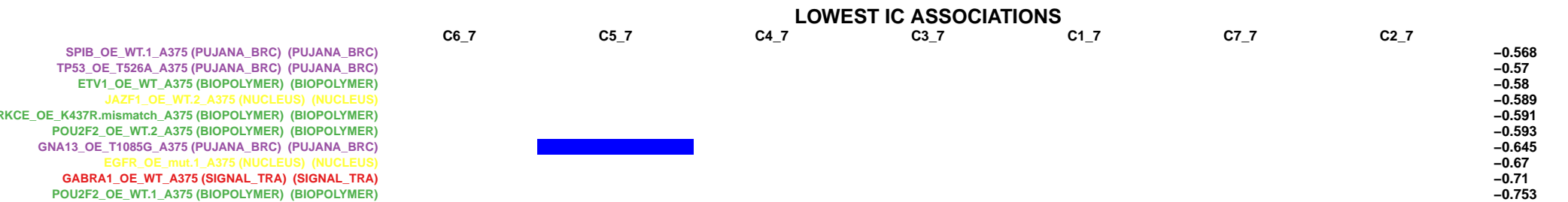




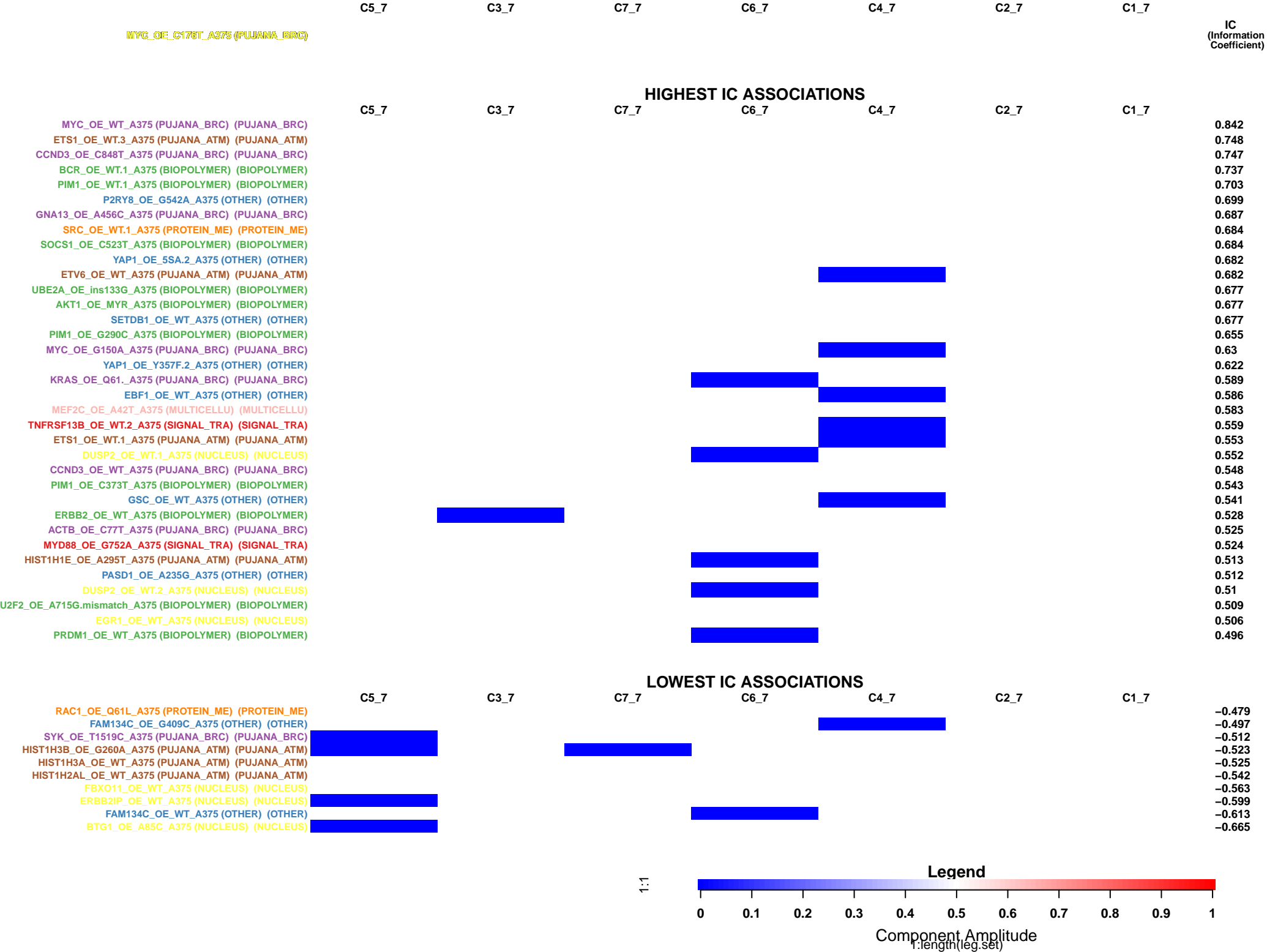
NFKBIA\_OE\_C126A\_A375 (NUCLEUS)



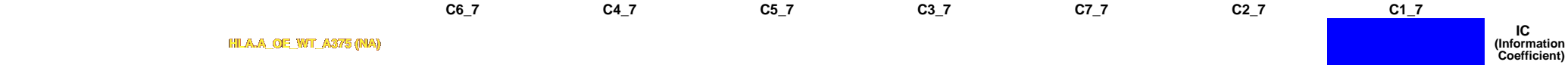
C6_7	C5_7	C4_7	C3_7	C1_7	C7_7	C2_7	IC (Information Coefficient)
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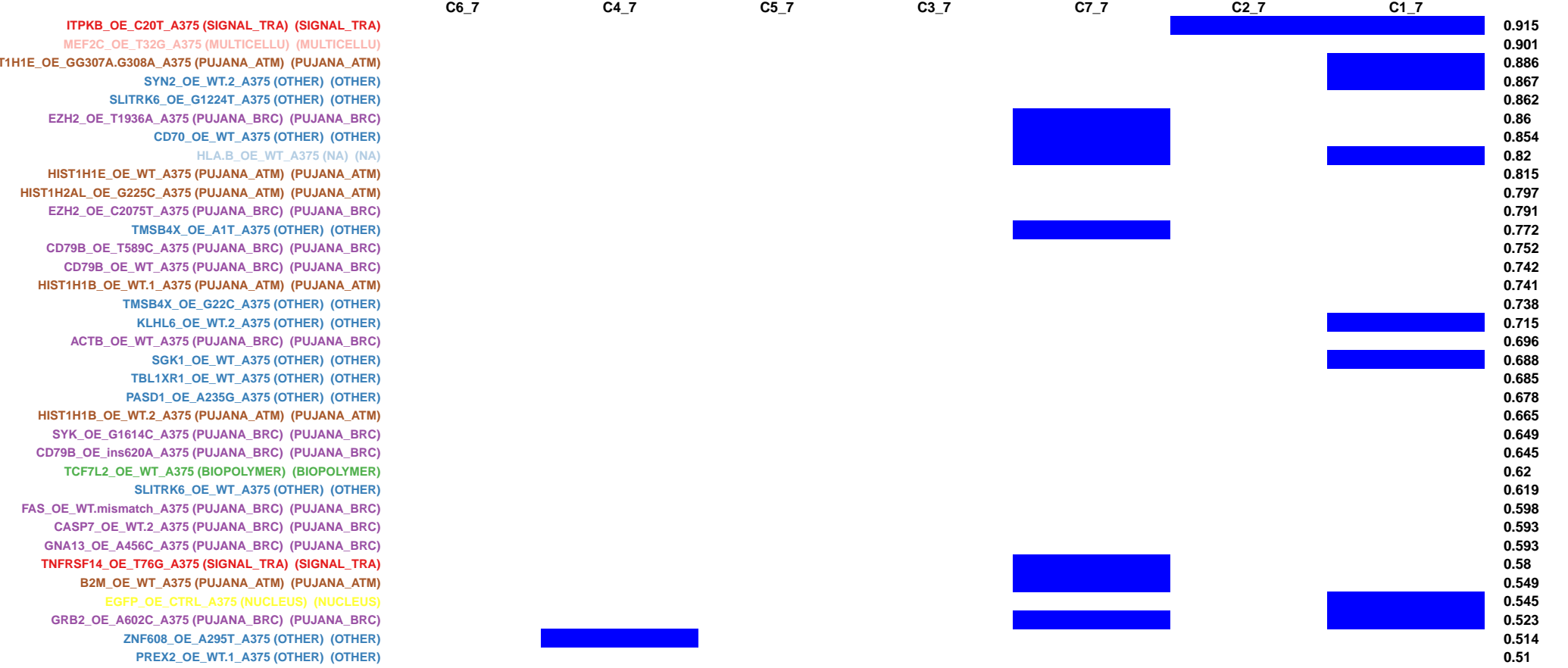
MYC\_OE\_C176T\_A375 (PUJANA\_BRC)



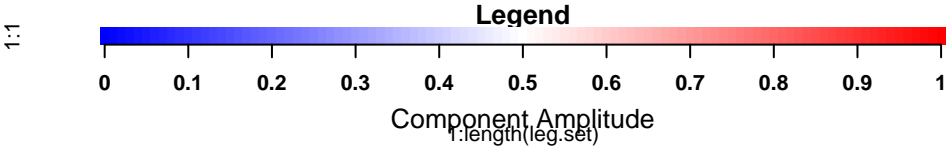
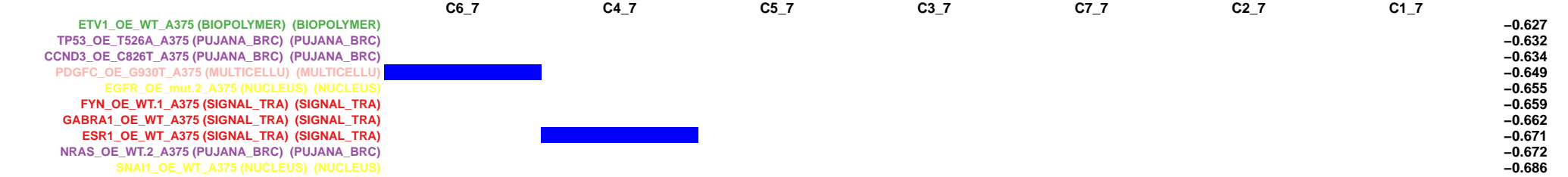
HLA.A\_OE\_WT\_A375 (NA)



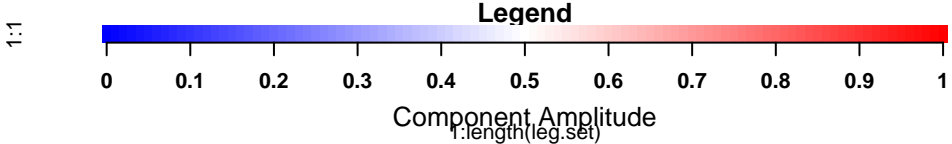
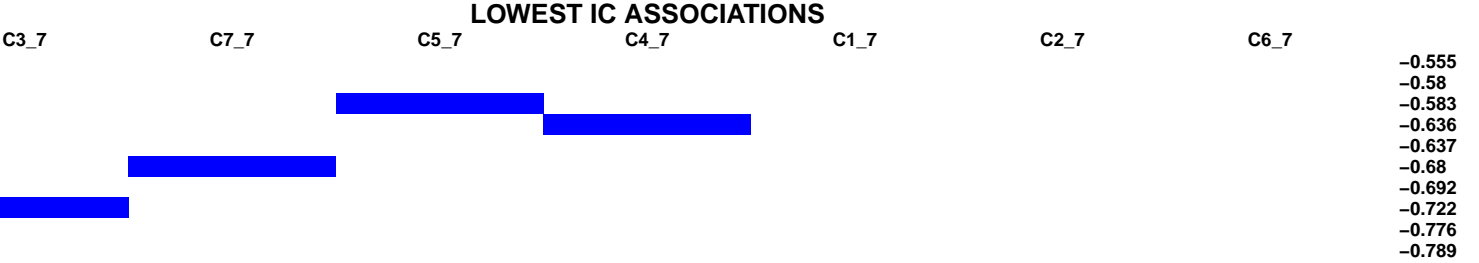
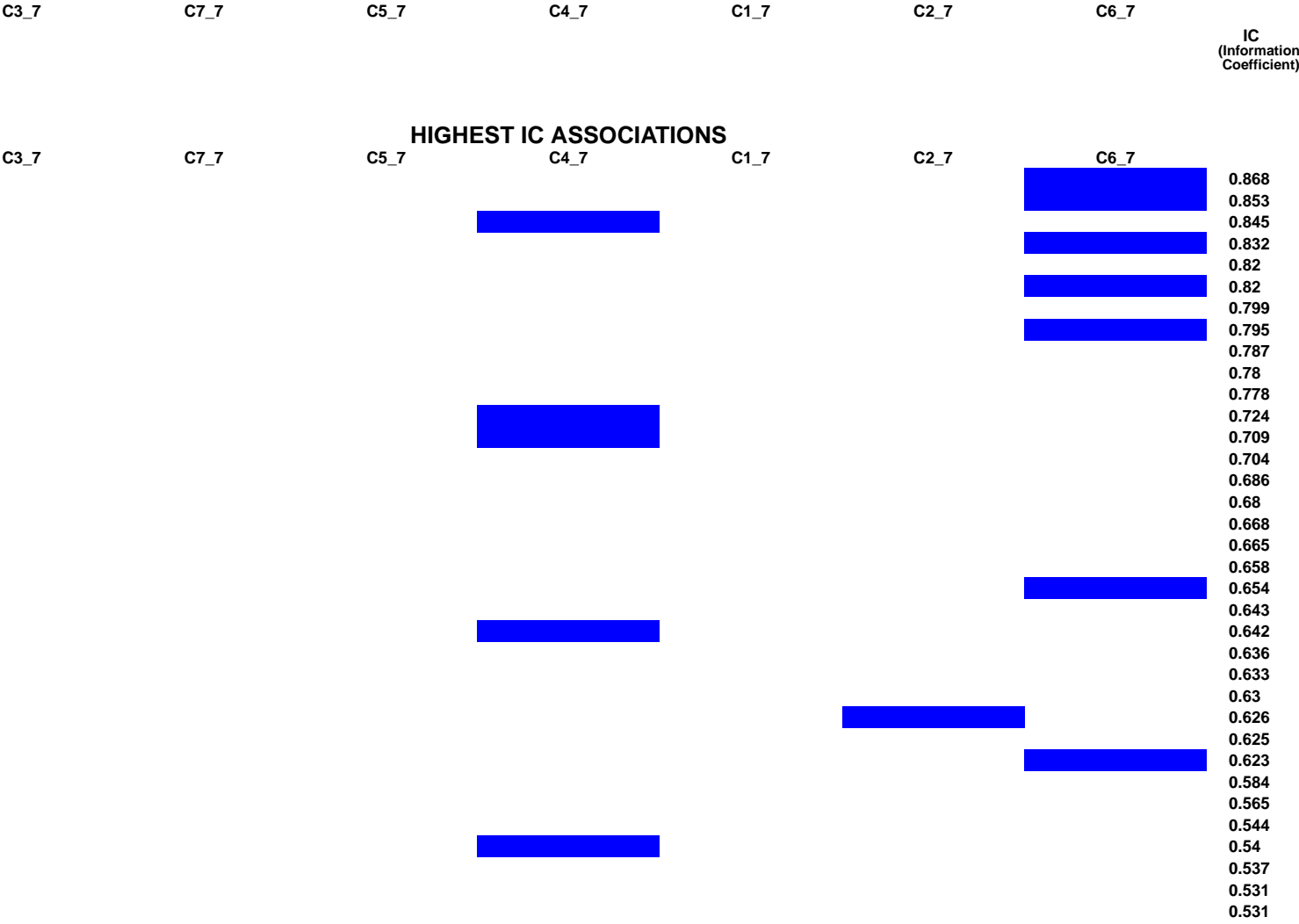
HIGHEST IC ASSOCIATIONS



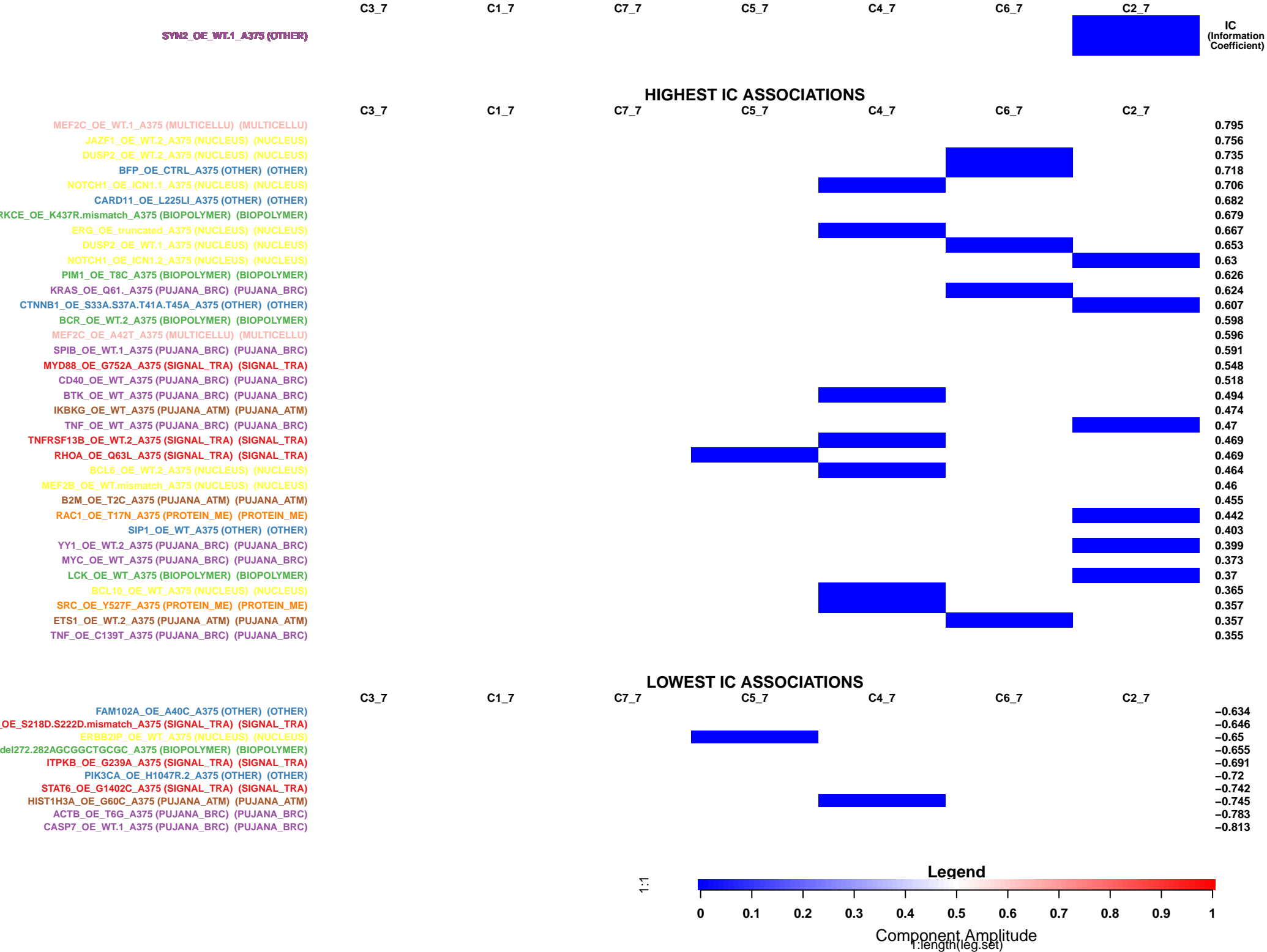
LOWEST IC ASSOCIATIONS



PIM1\_OE\_T8C\_A375 (BIOPOLYMER)



SYN2\_OE\_WT.1\_A375 (OTHER)



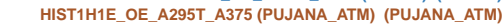
C6_7	C4_7	C2_7	C5_7	C7_7	C3_7	C1_7
------	------	------	------	------	------	------

**0.756**

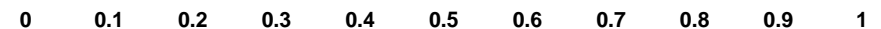
C6\_7                  C4\_7                  C2\_7                  C5\_7                  C7\_7                  C3\_7                  C1\_7



**C6\_7                      C4\_7                      C2\_7                      C5\_7                      C7\_7                      C3\_7                      C1\_7**



1:1

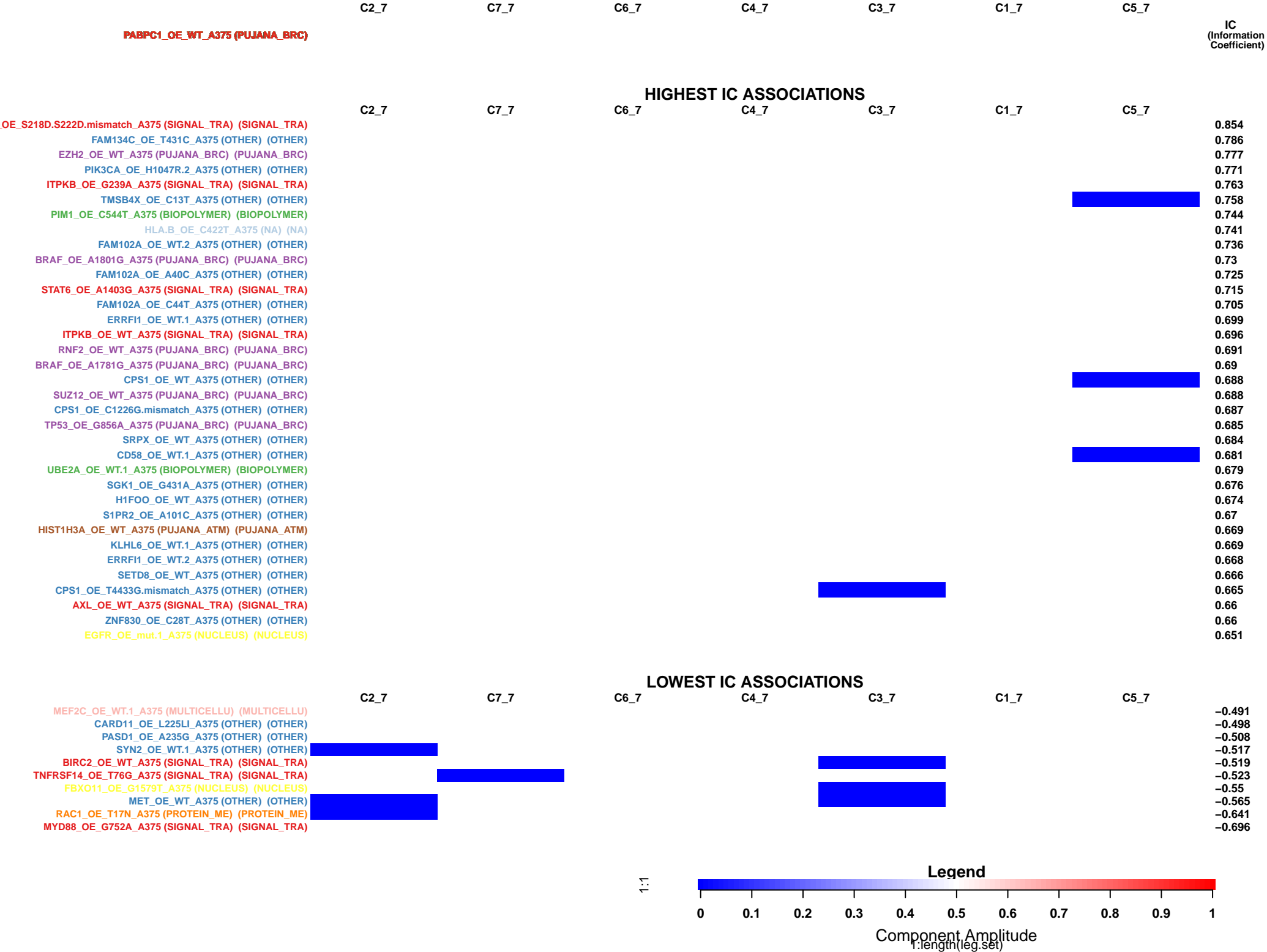
Component Amplitude  
1:length(leg.set)

C6_7	C4_7	C5_7	C3_7	C2_7	C7_7	C1_7	IC (Information Coefficient)

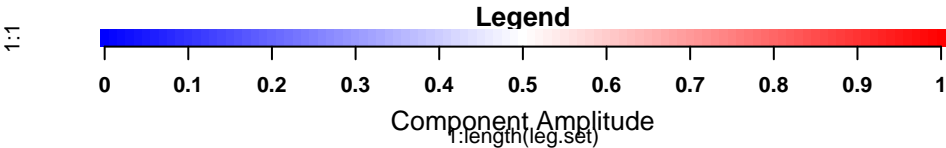
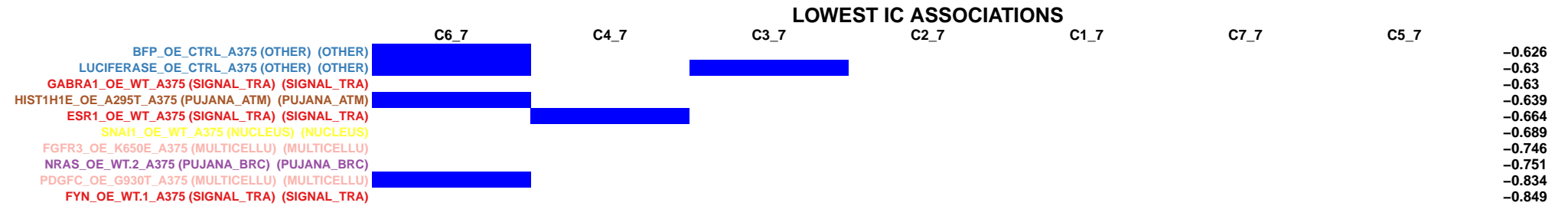
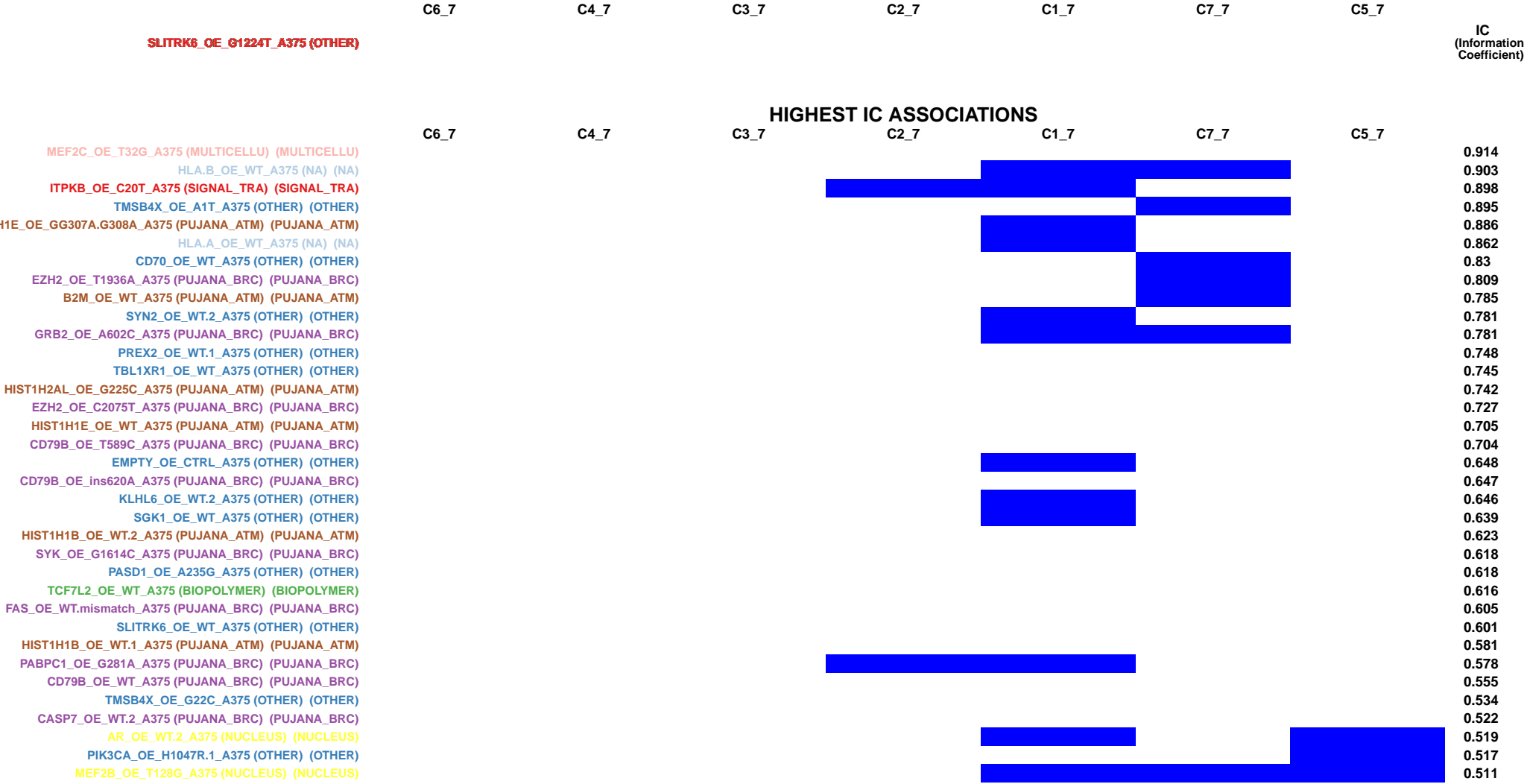




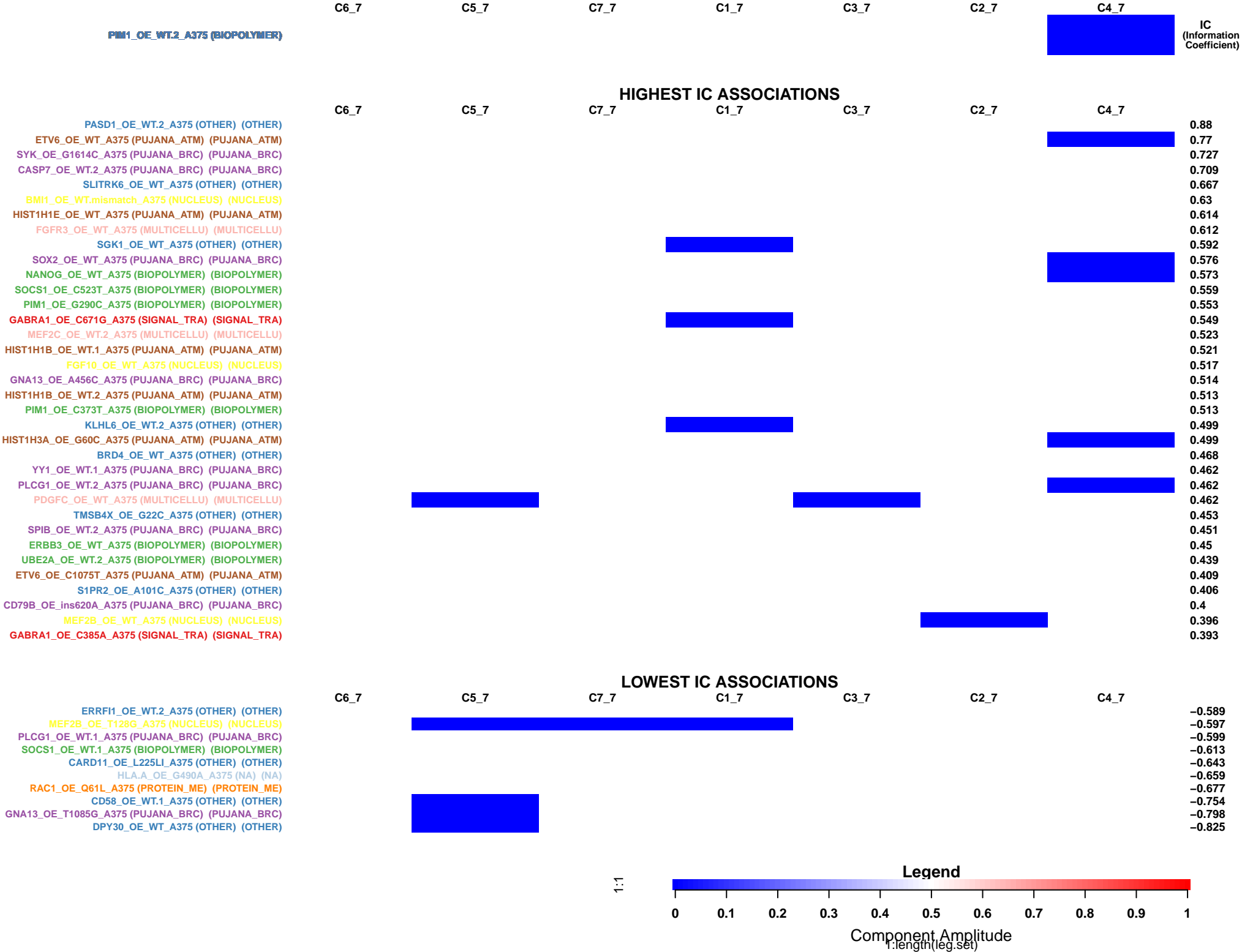
PABPC1\_OE\_WT\_A375 (PUJANA\_BRC)



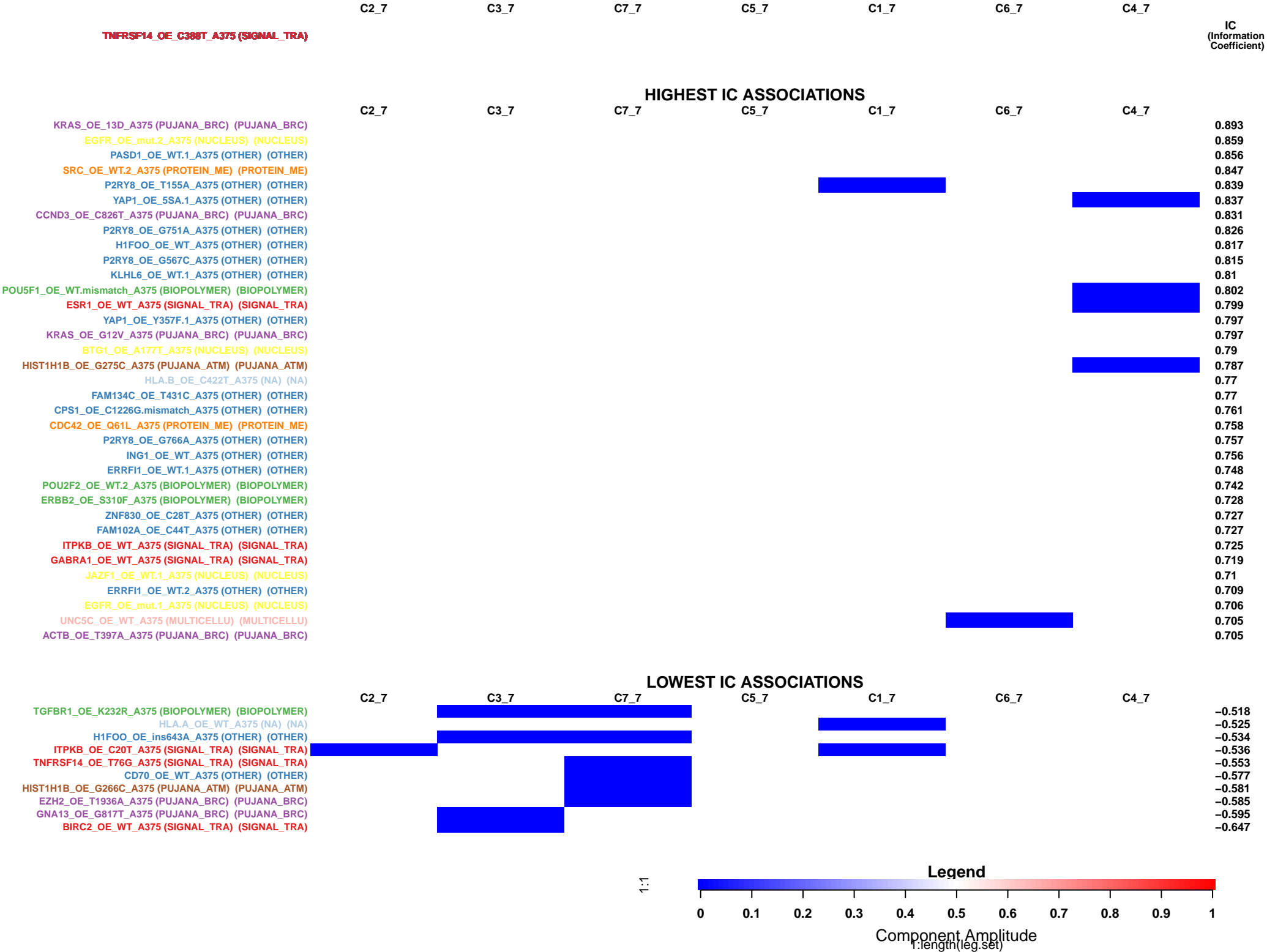
SLITRK6\_OE\_G1224T\_A375 (OTHER)



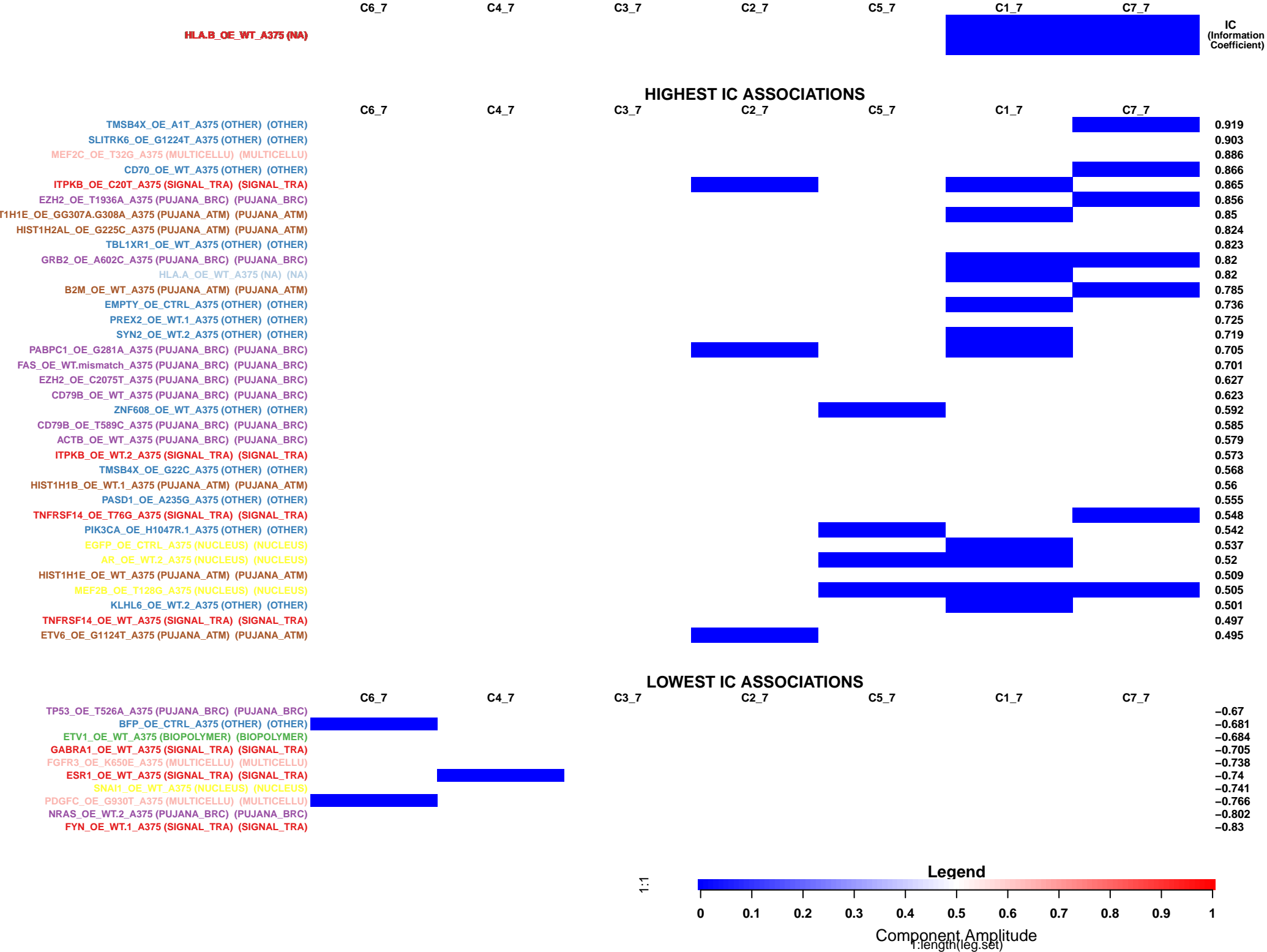
PIM1\_OE\_WT.2\_A375 (BIOPOLYMER)



TNFRSF14\_OE\_C388T\_A375 (SIGNAL\_TRA)



HLA.B\_OE\_WT\_A375 (NA)

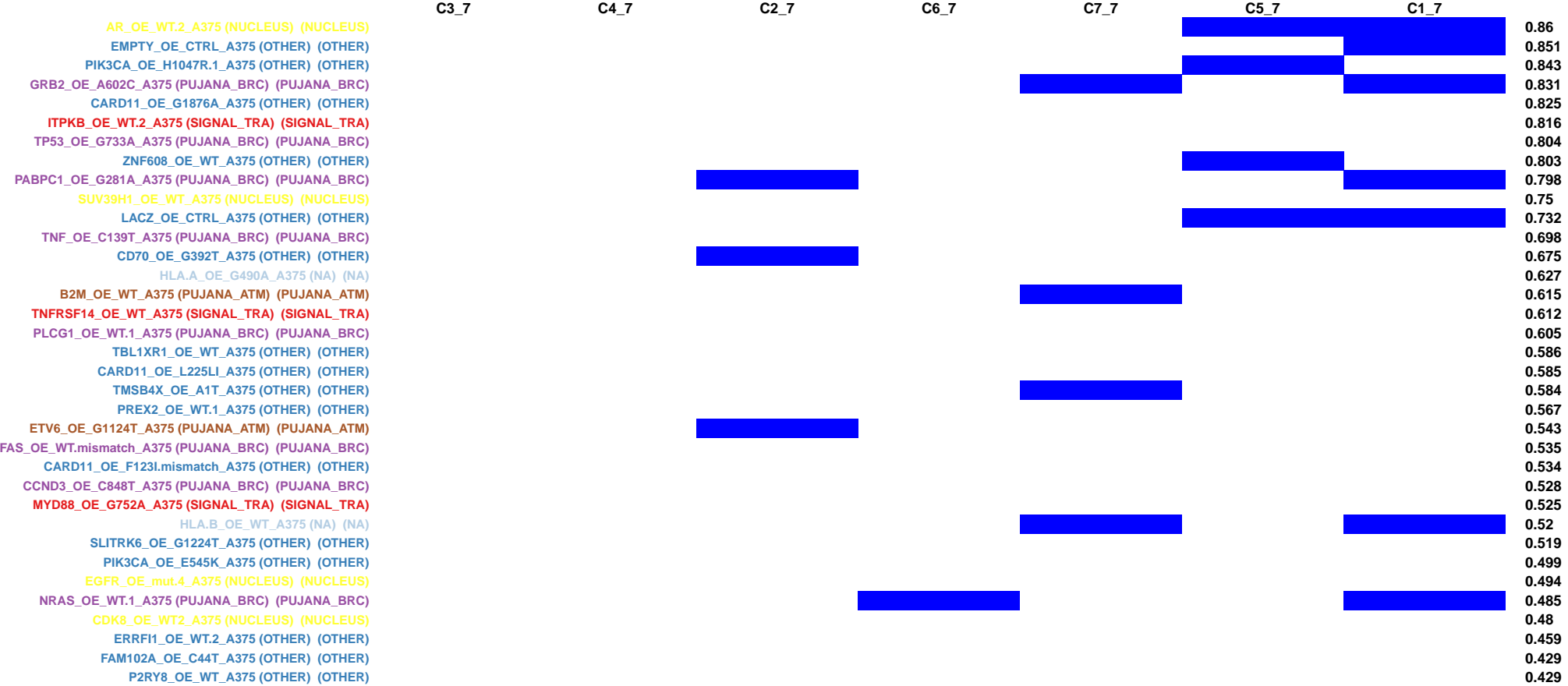


AR\_OE\_WT.2\_A375 (NUCLEUS)

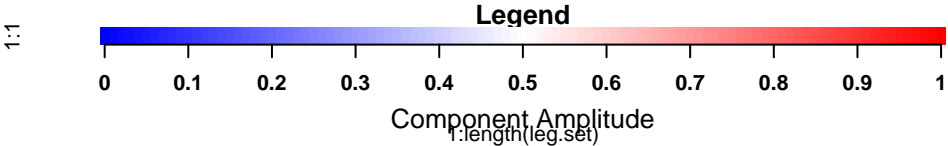
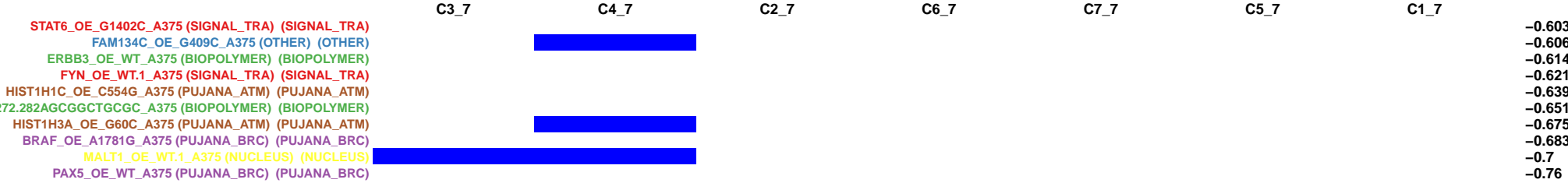
AR\_OE\_WT.2\_A375 (NUCLEUS)

IC  
(Information  
Coefficient)

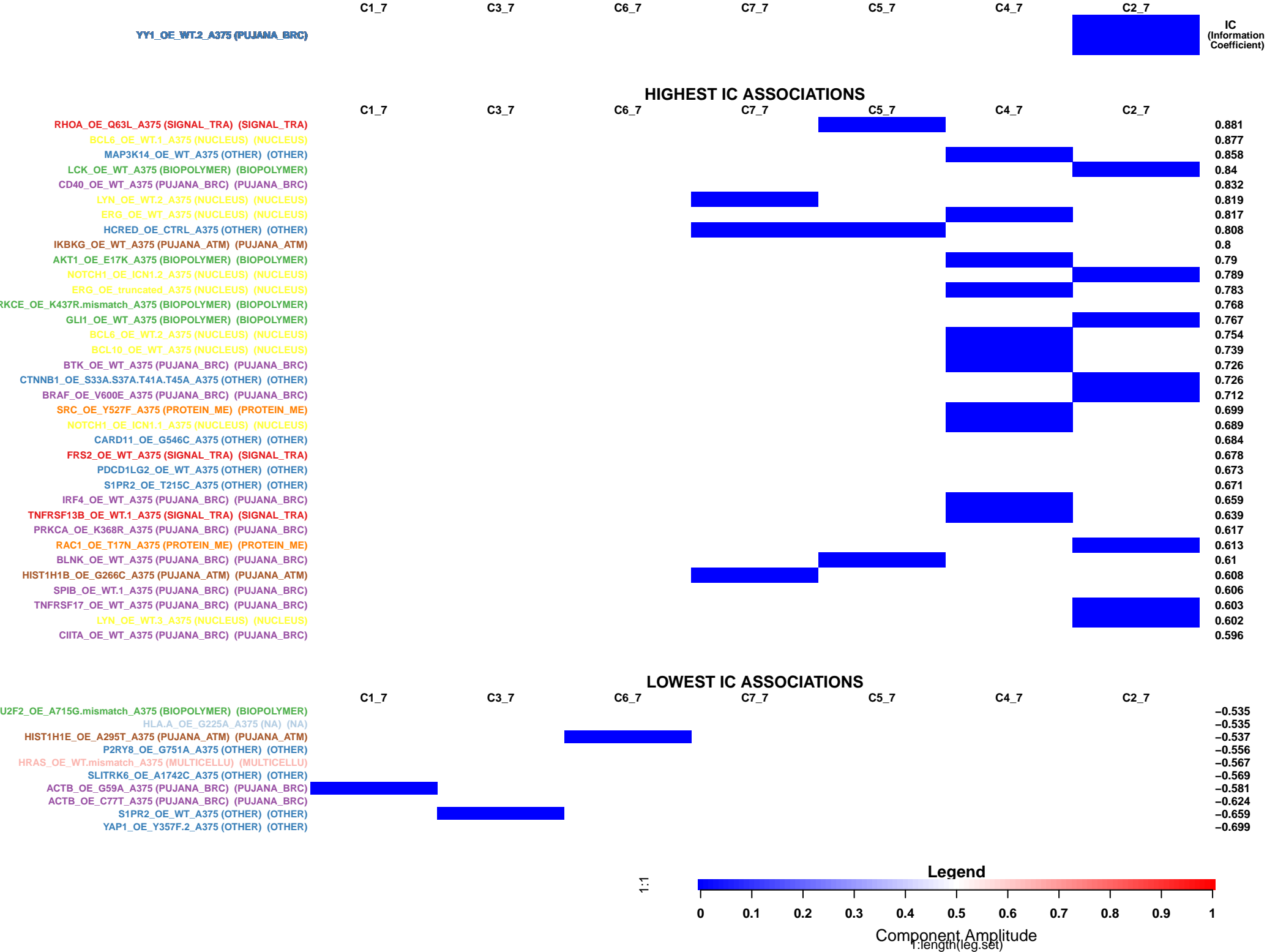
HIGHEST IC ASSOCIATIONS



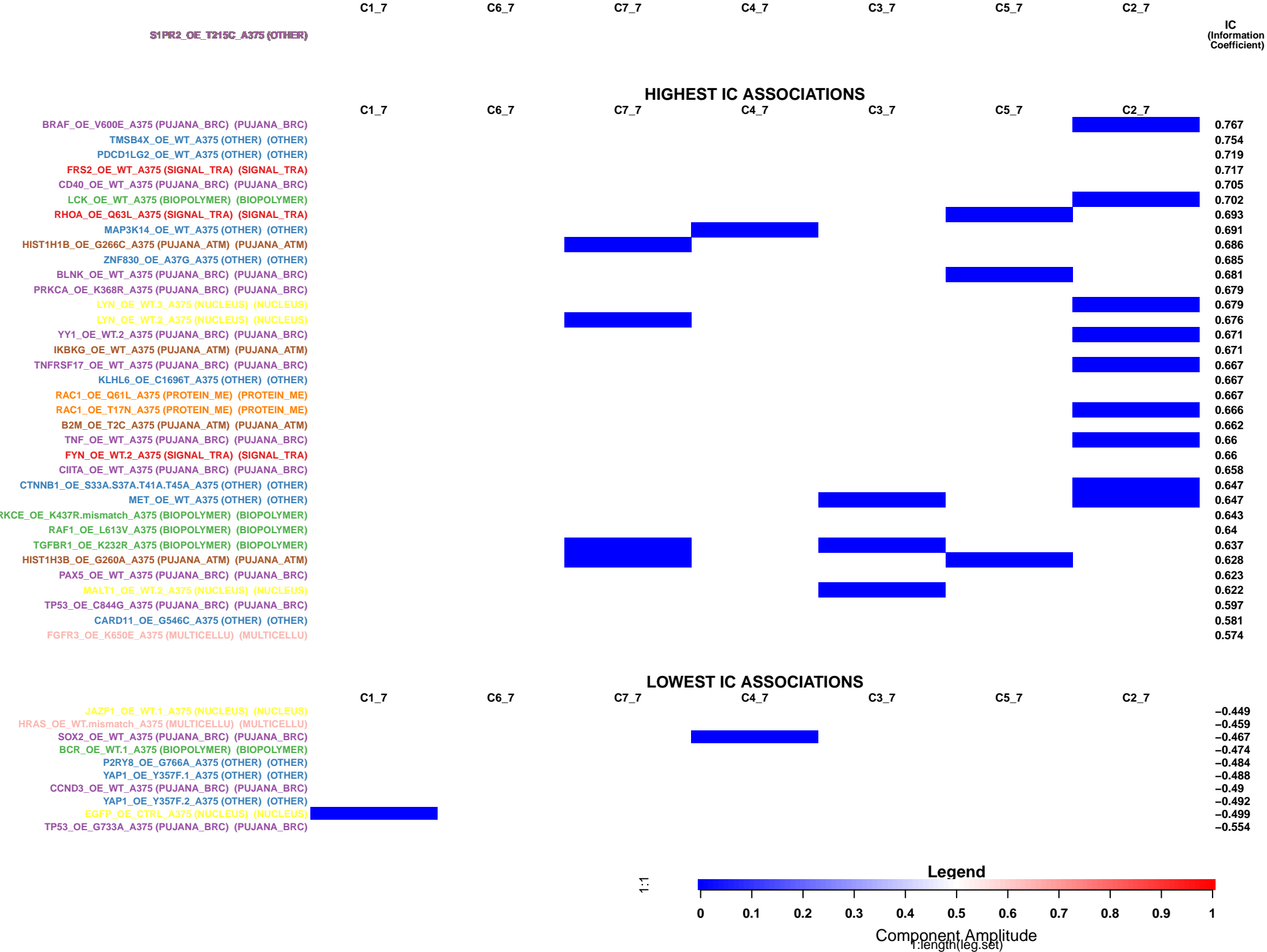
LOWEST IC ASSOCIATIONS



YY1\_OE\_WT.2\_A375 (PUJANA\_BRC)

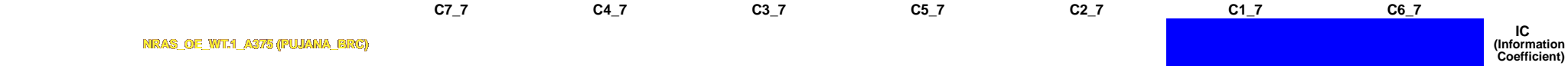


S1PR2\_OE\_T215C\_A375 (OTHER)

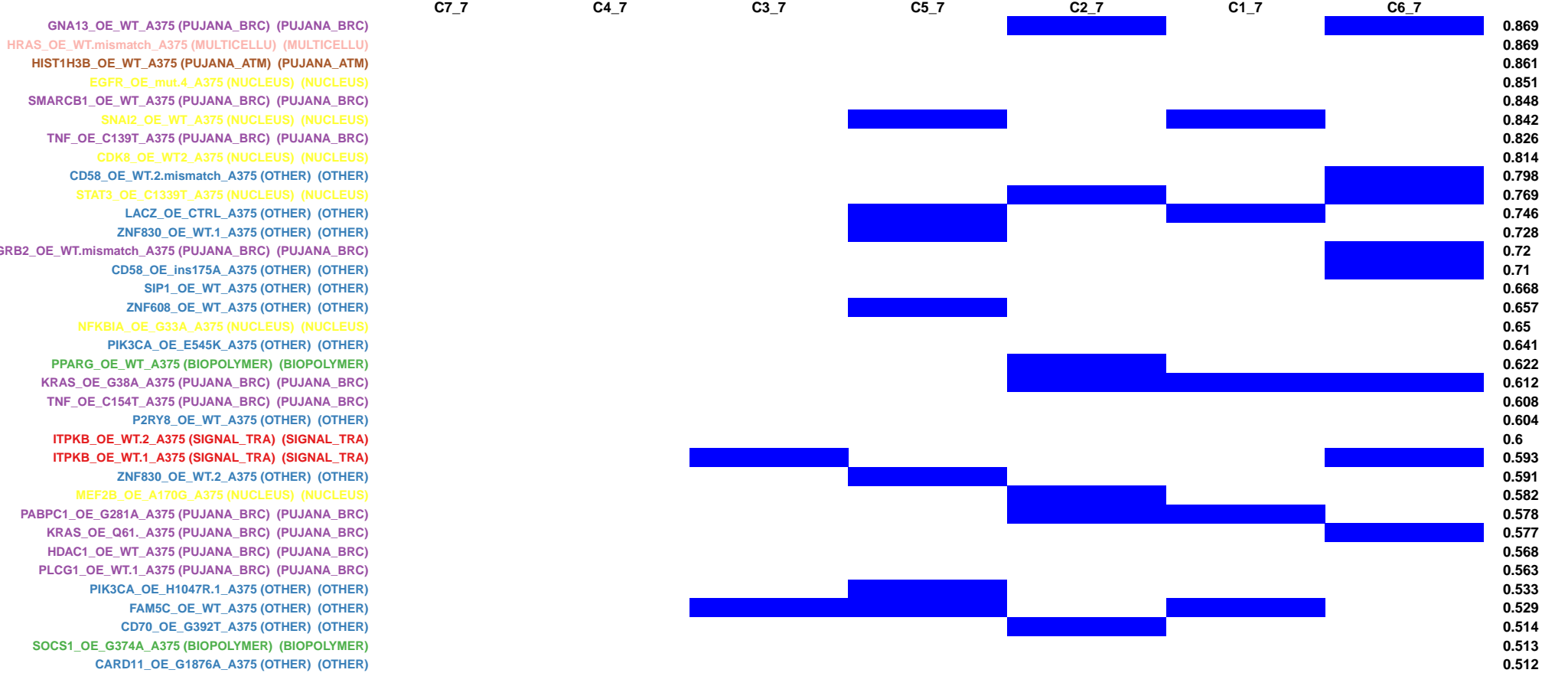




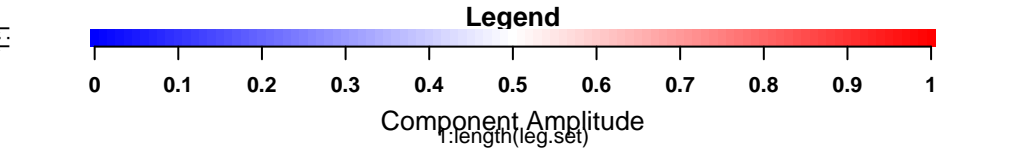
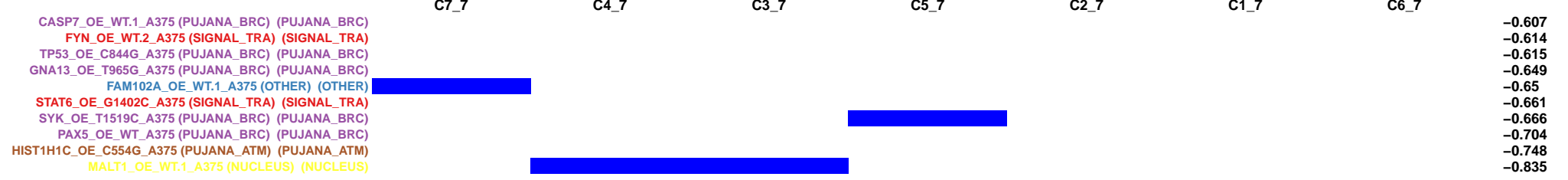
NRAS\_OE\_WT.1\_A375 (PUJANA\_BRC)



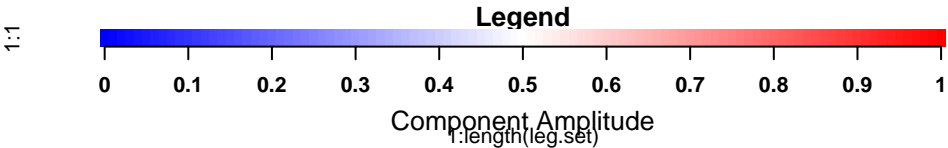
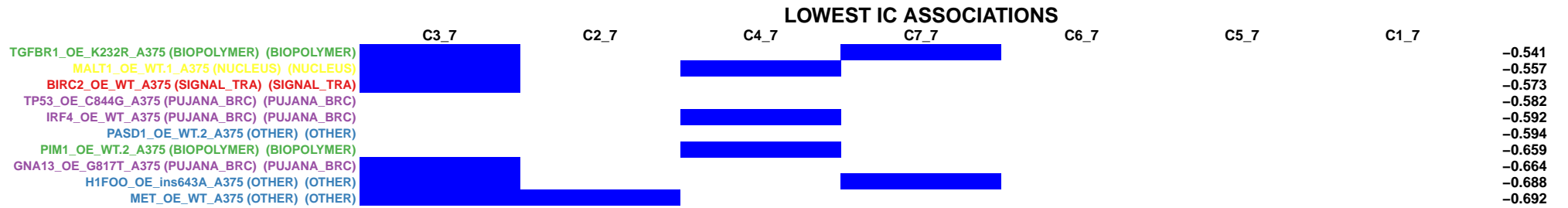
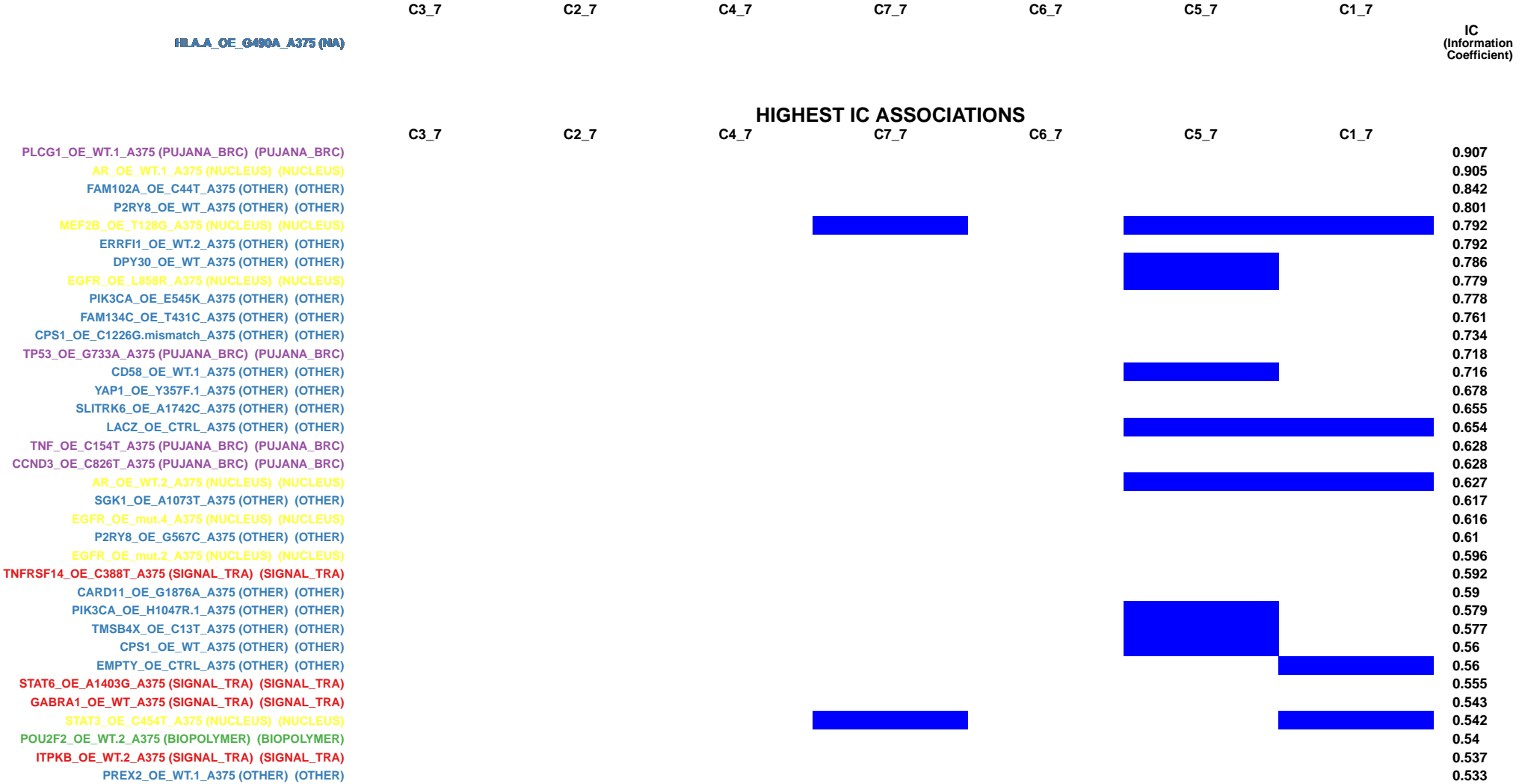
HIGHEST IC ASSOCIATIONS



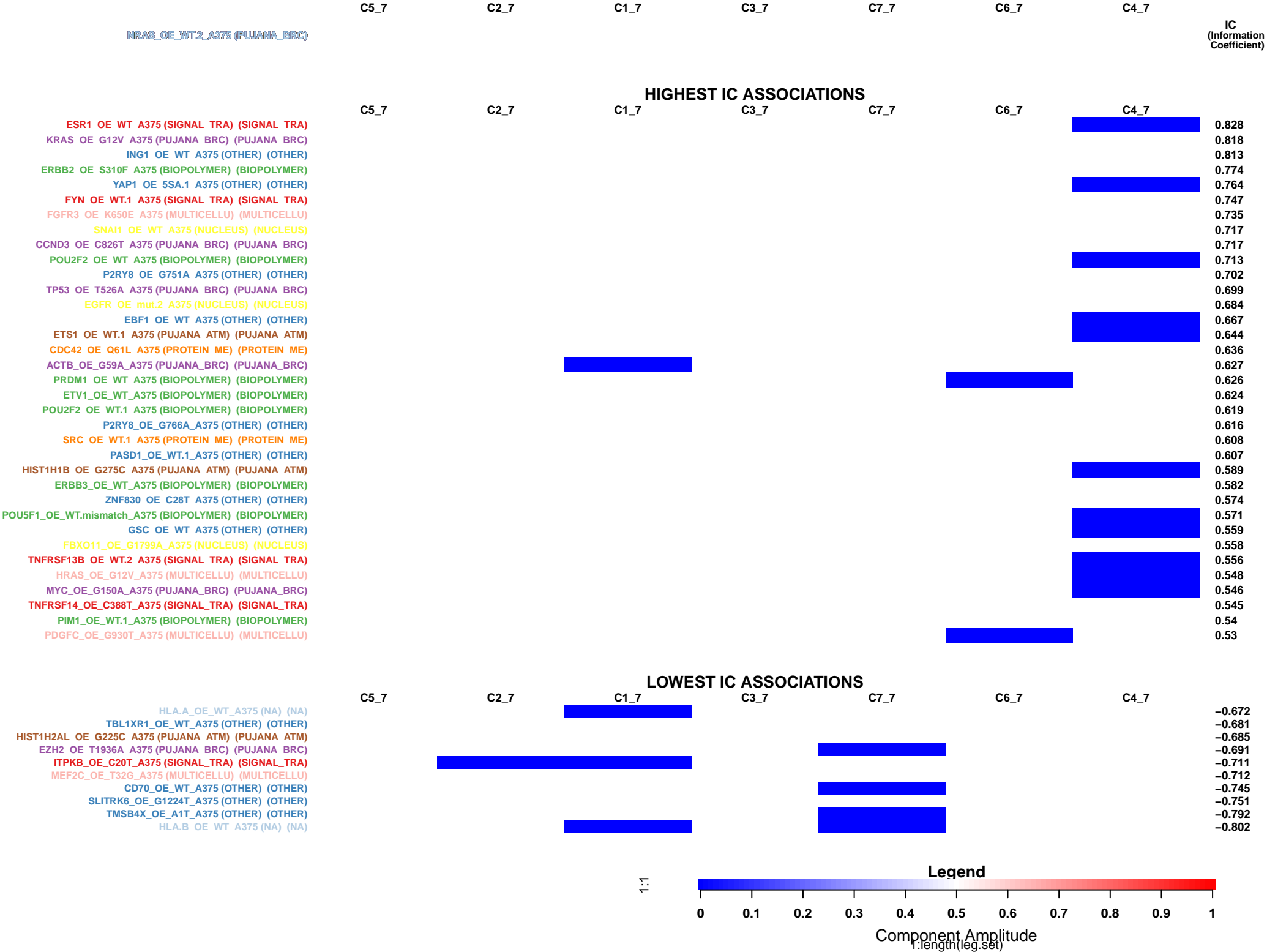
LOWEST IC ASSOCIATIONS



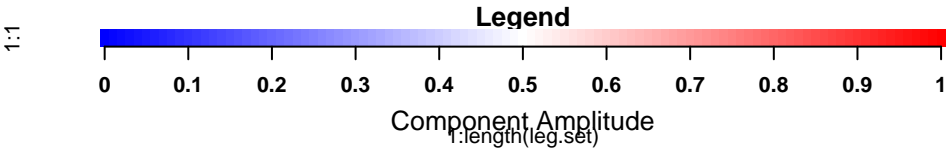
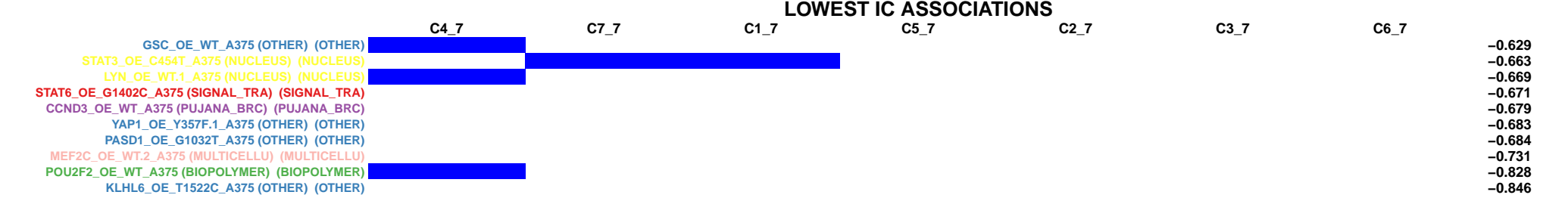
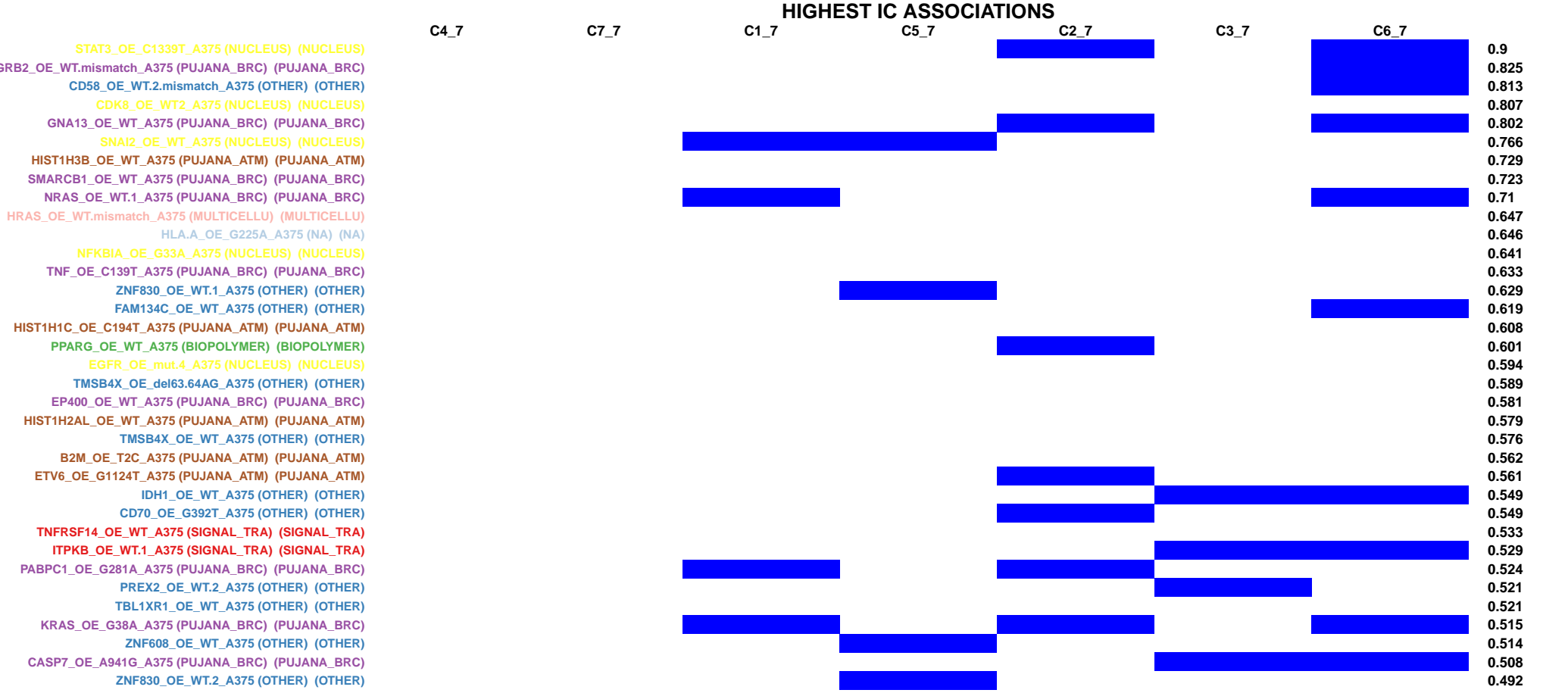
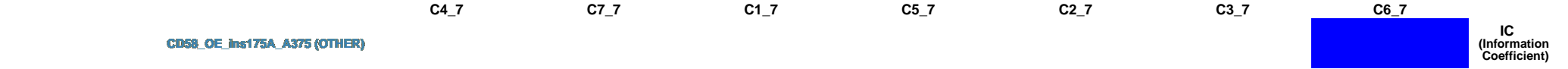
HLA.A\_OE\_G490A\_A375 (NA)



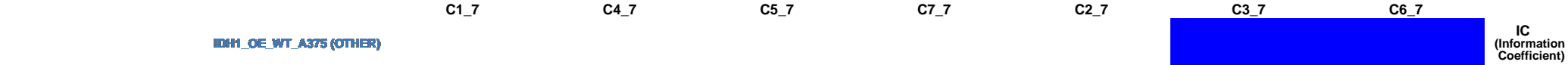
NRAS\_OE\_WT.2\_A375 (PUJANA\_BRC)



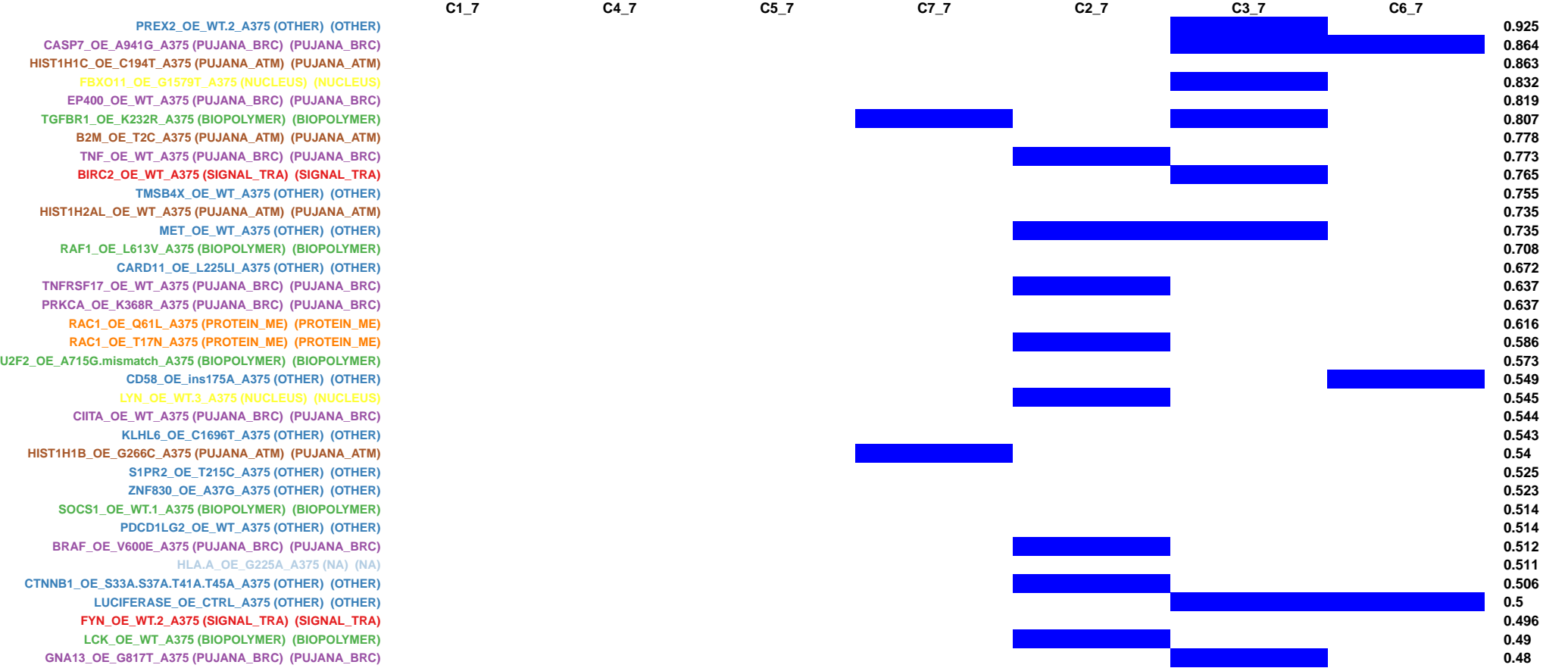
CD58\_OE\_ins175A\_A375 (OTHER)



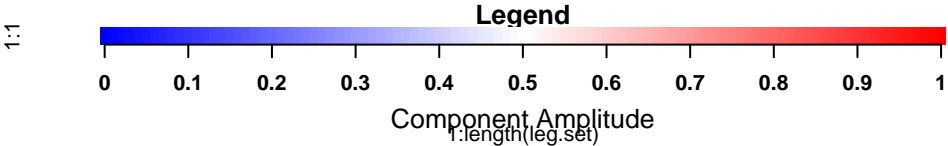
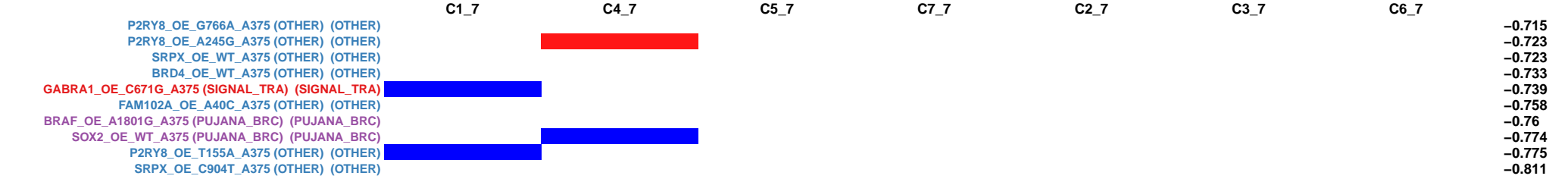
IDH1\_OE\_WT\_A375 (OTHER)



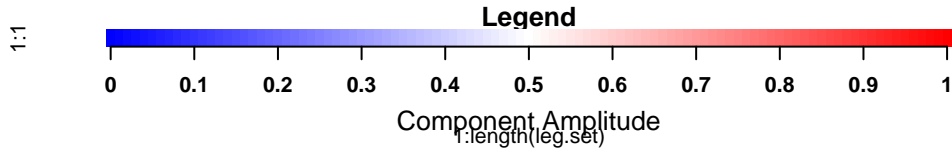
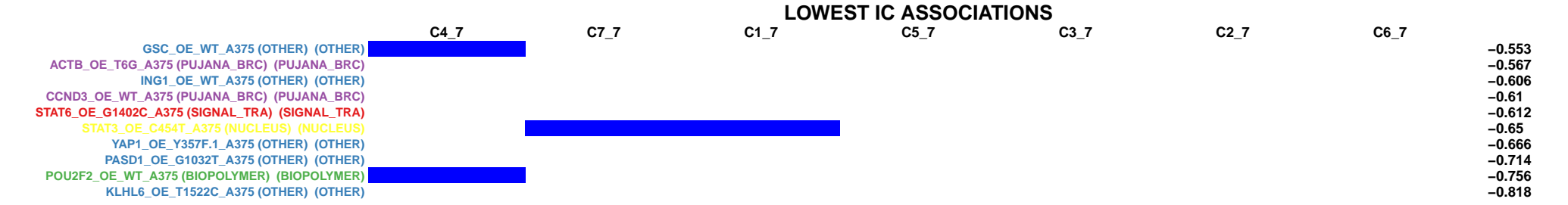
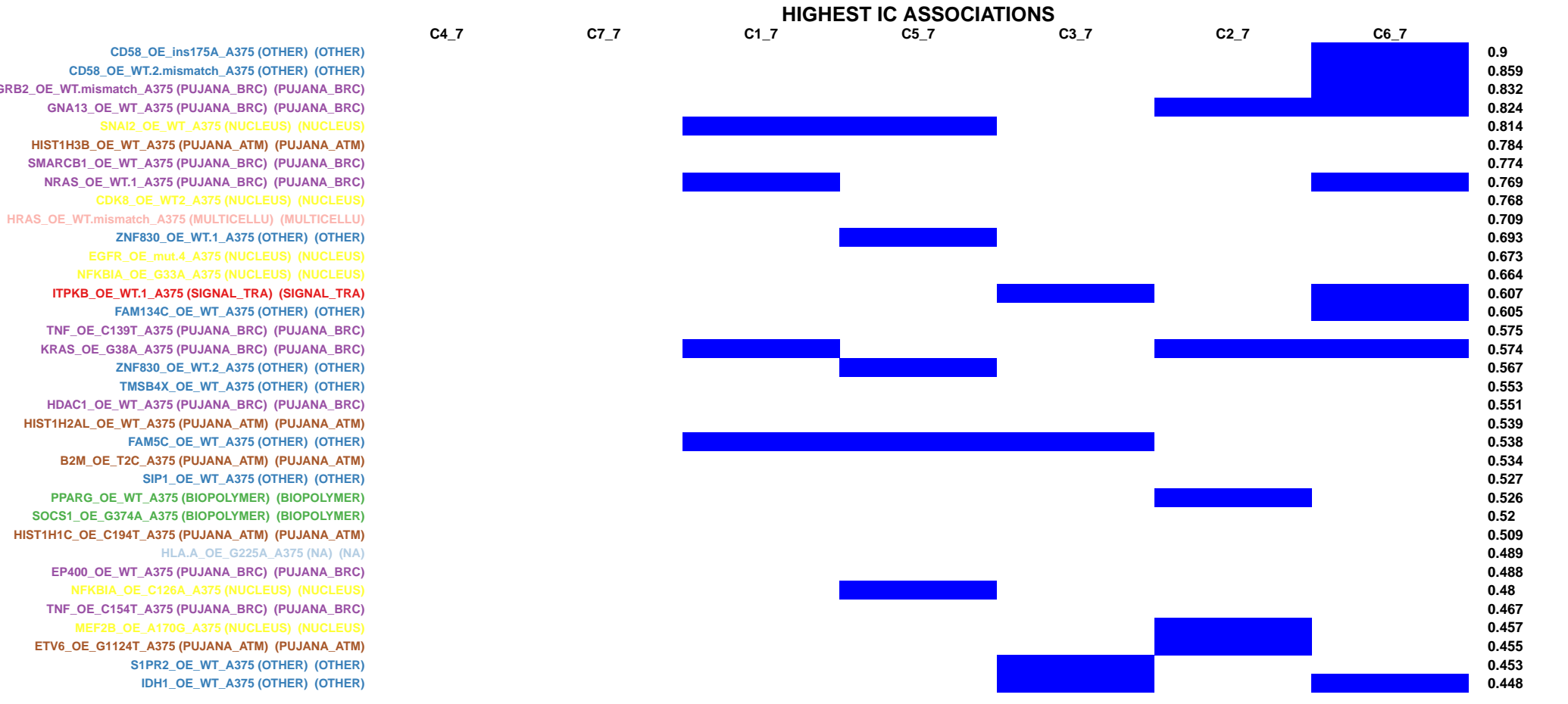
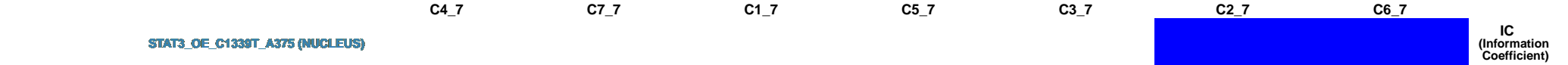
HIGHEST IC ASSOCIATIONS



LOWEST IC ASSOCIATIONS



STAT3\_OE\_C1339T\_A375 (NUCLEUS)



MYC\_OE\_WT\_A375 (PUJANA\_BRC)

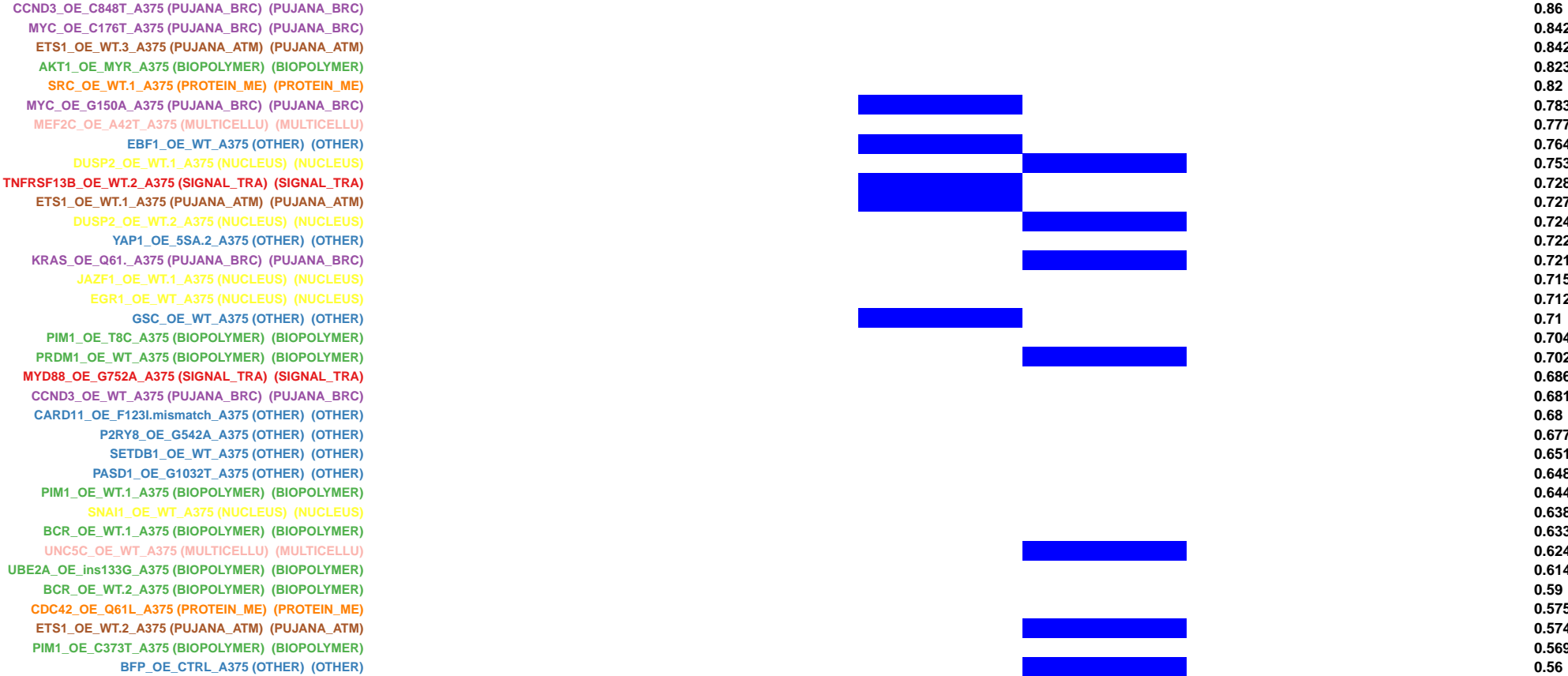
C5\_7 C3\_7 C7\_7 C4\_7 C6\_7 C1\_7 C2\_7

MYC\_OE\_WT\_A375 (PUJANA\_BRC)

IC  
(Information  
Coefficient)

HIGHEST IC ASSOCIATIONS

C5\_7 C3\_7 C7\_7 C4\_7 C6\_7 C1\_7 C2\_7



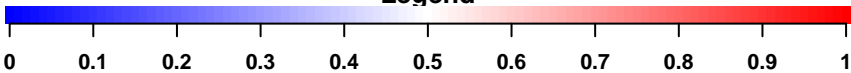
LOWEST IC ASSOCIATIONS

C5\_7 C3\_7 C7\_7 C4\_7 C6\_7 C1\_7 C2\_7



1:1

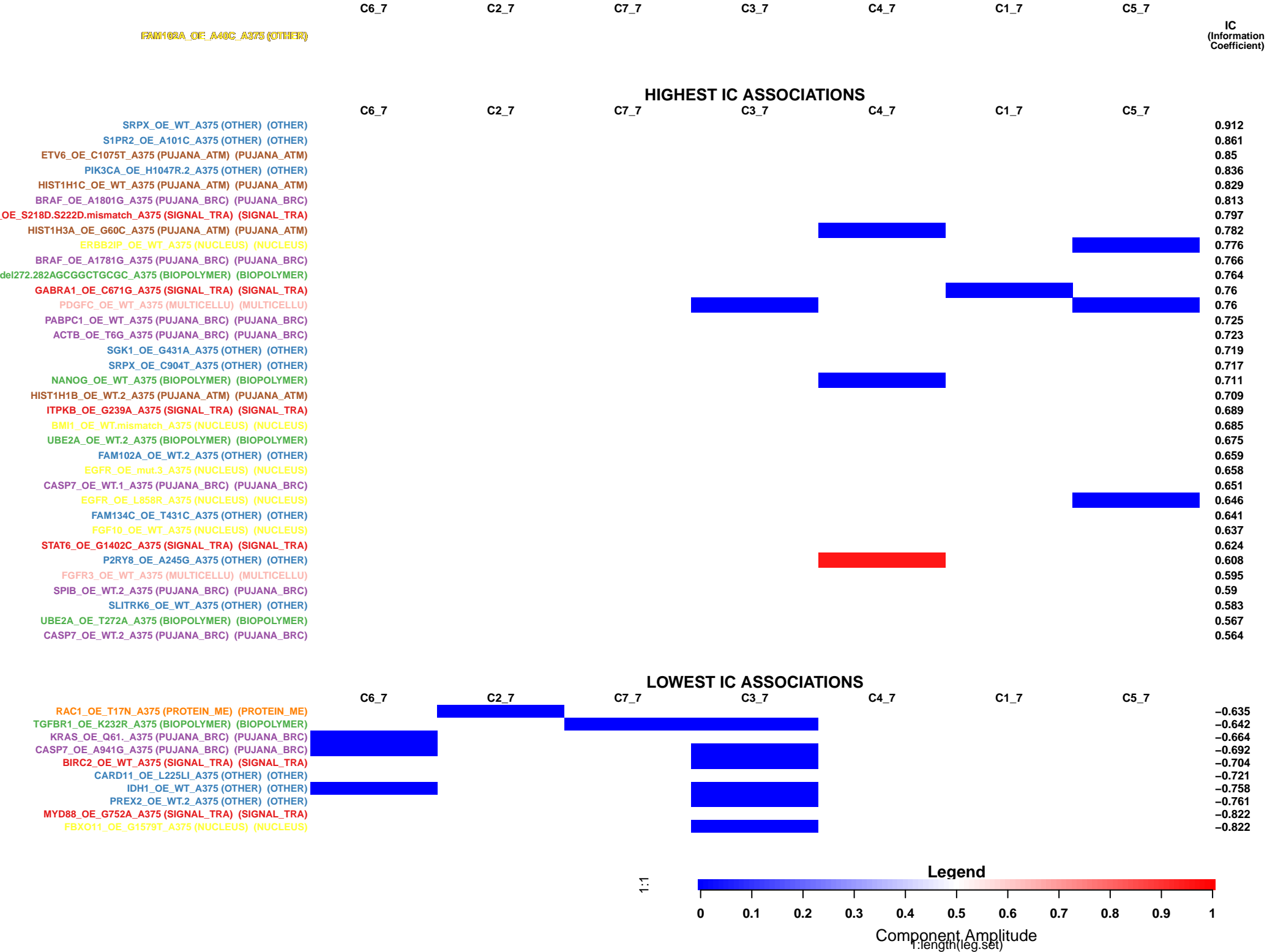
Legend



Component Amplitude

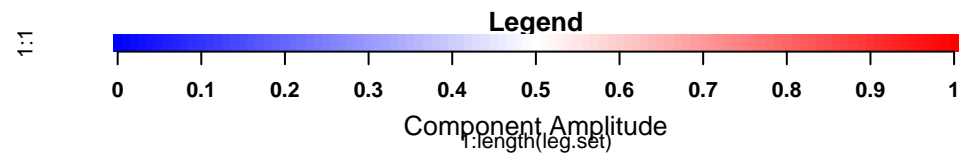
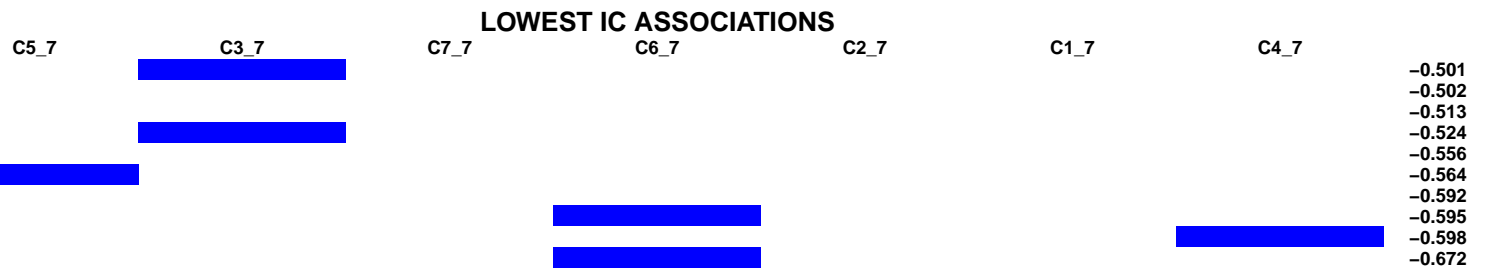
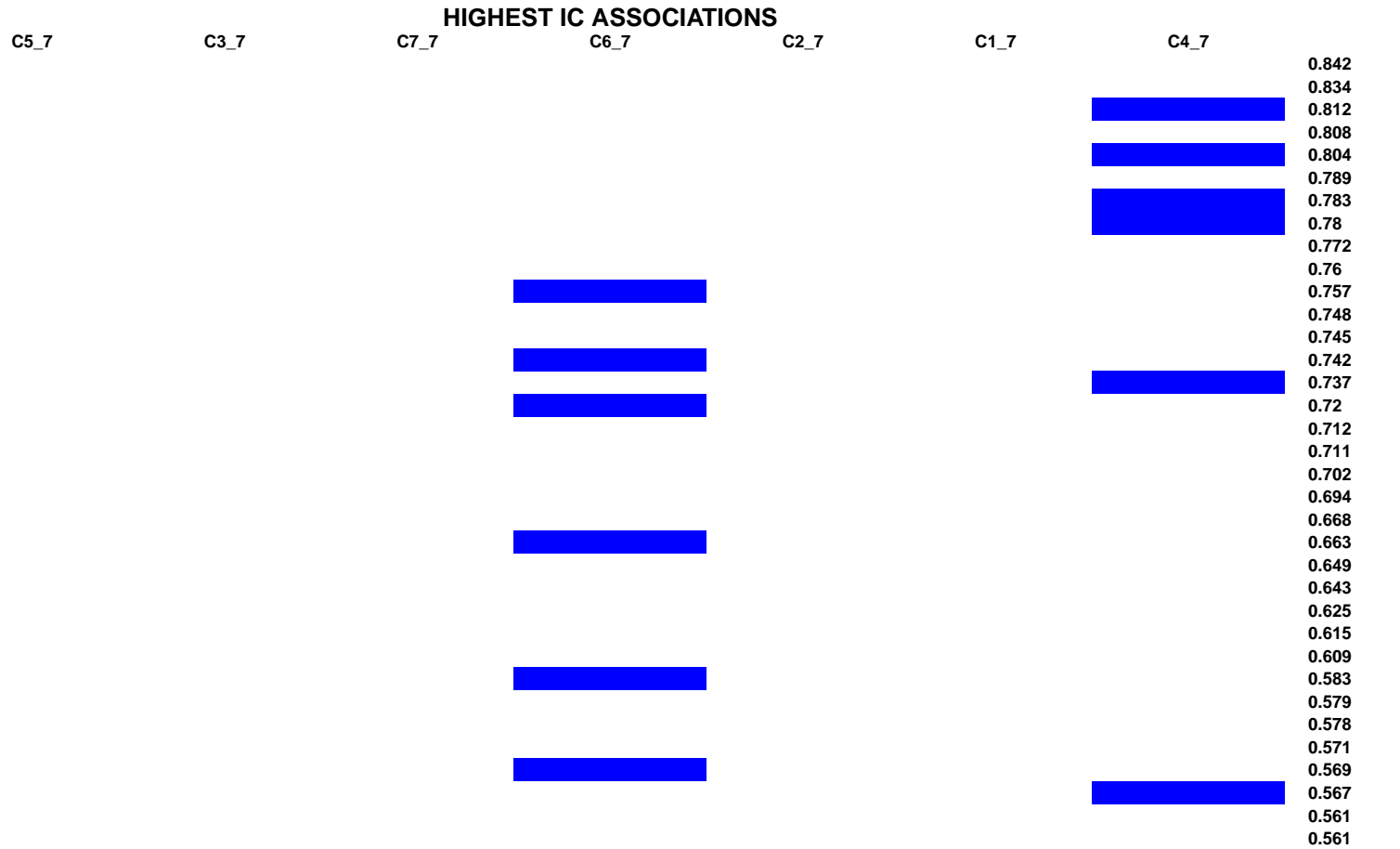
1:length(leg.set)

FAM102A\_OE\_A40C\_A375 (OTHER)

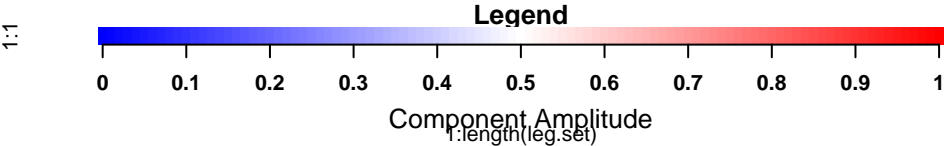
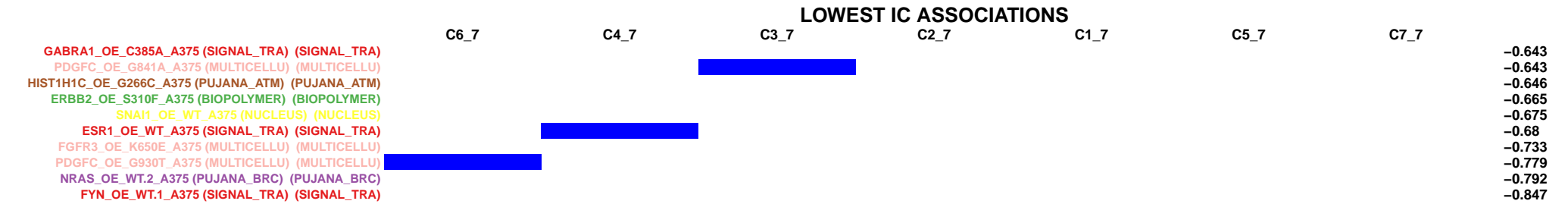
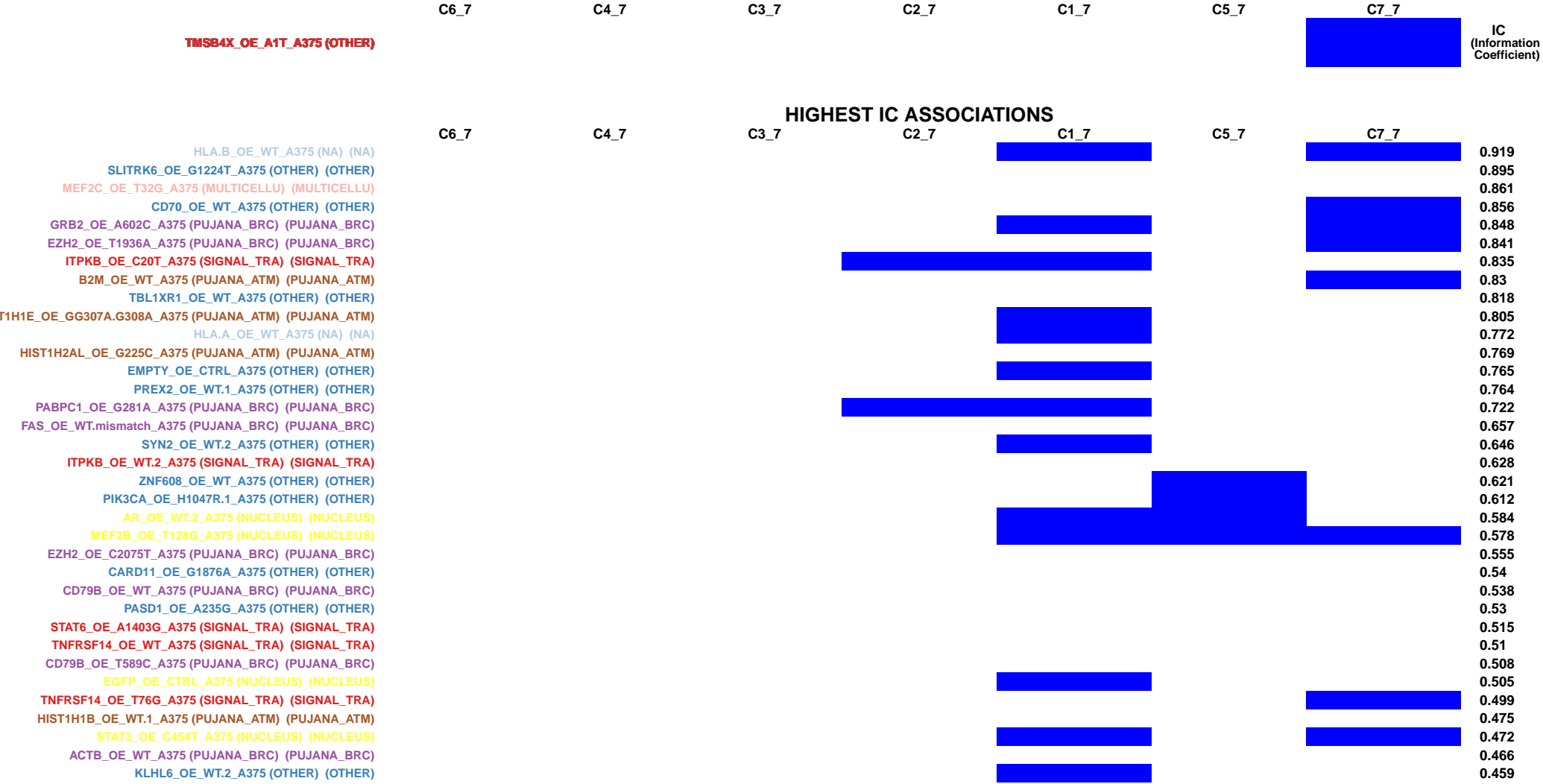




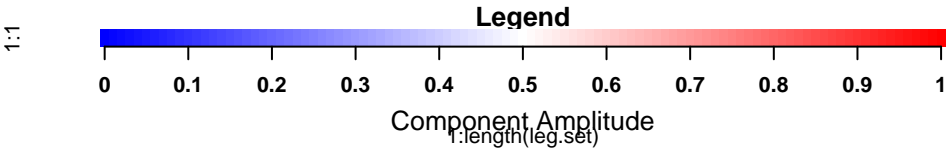
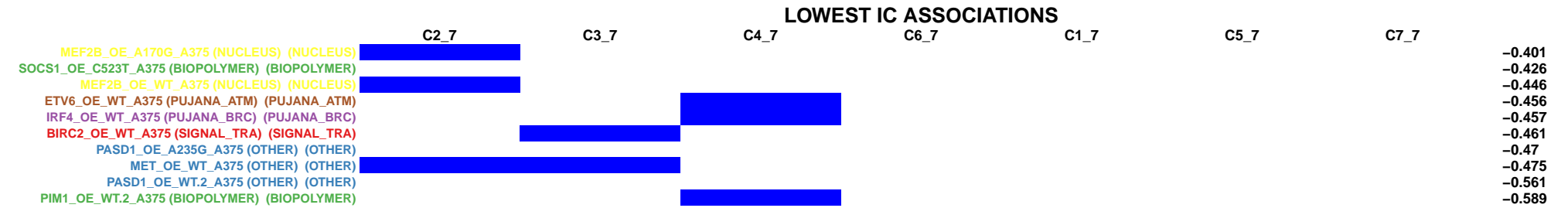
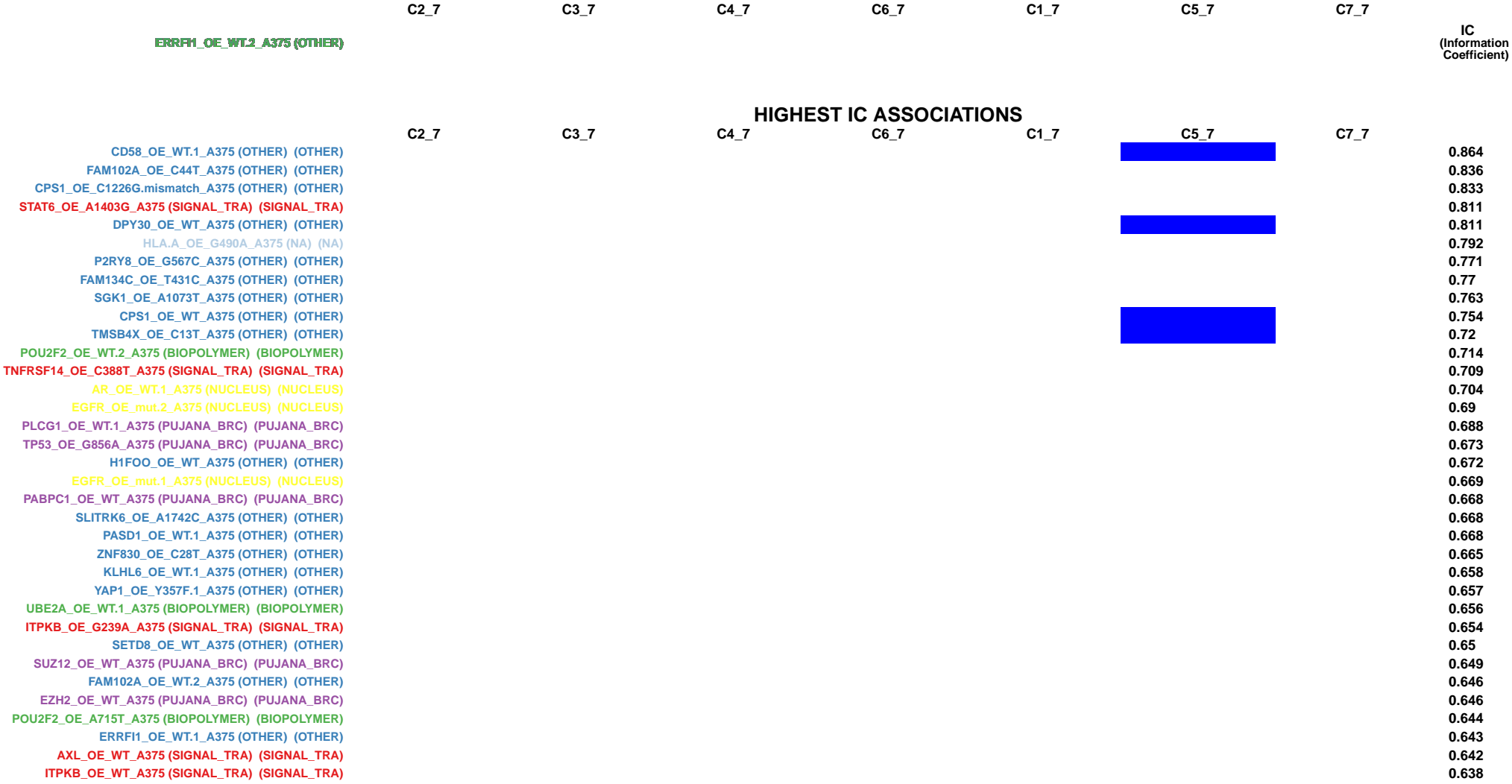
C5_7	C3_7	C7_7	C6_7	C2_7	C1_7	C4_7	IC (Information Coefficient)
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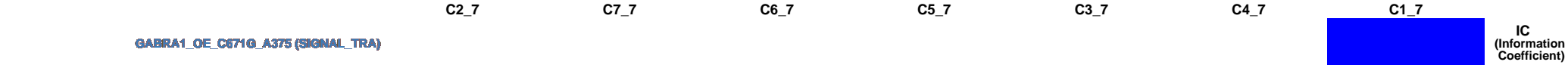
TMSB4X\_OE\_A1T\_A375 (OTHER)



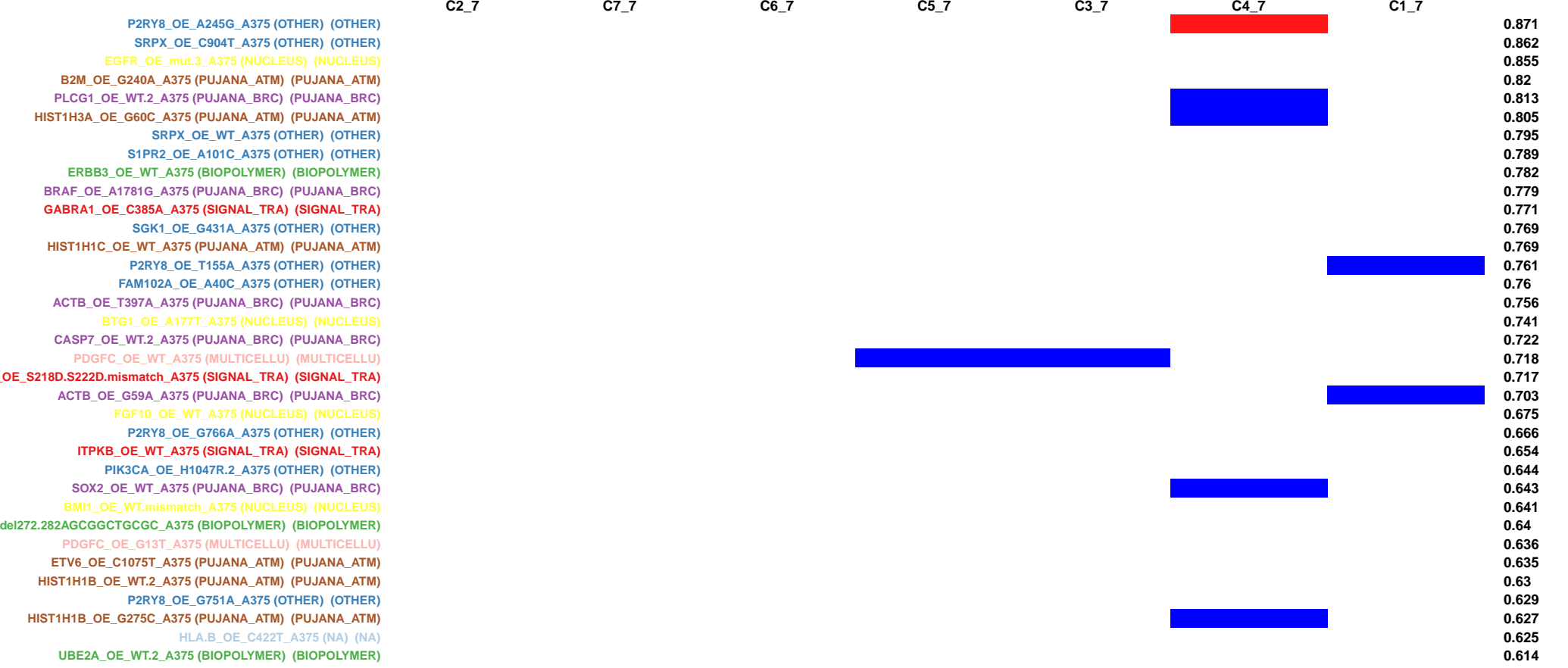
ERRFI1\_OE\_WT.2\_A375 (OTHER)



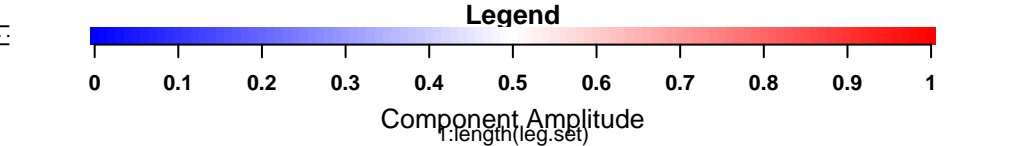
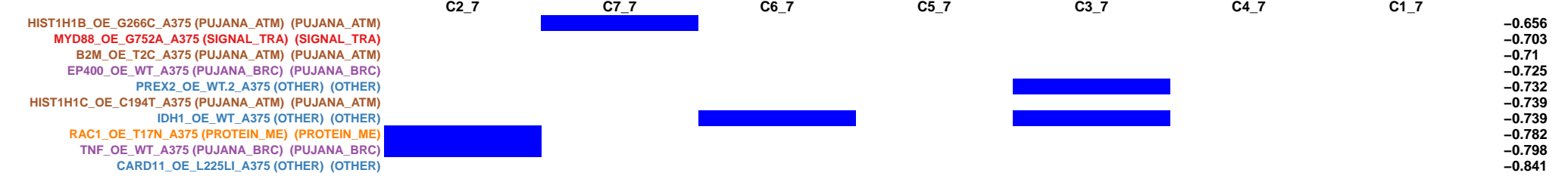
GABRA1\_OE\_C671G\_A375 (SIGNAL\_TRA)



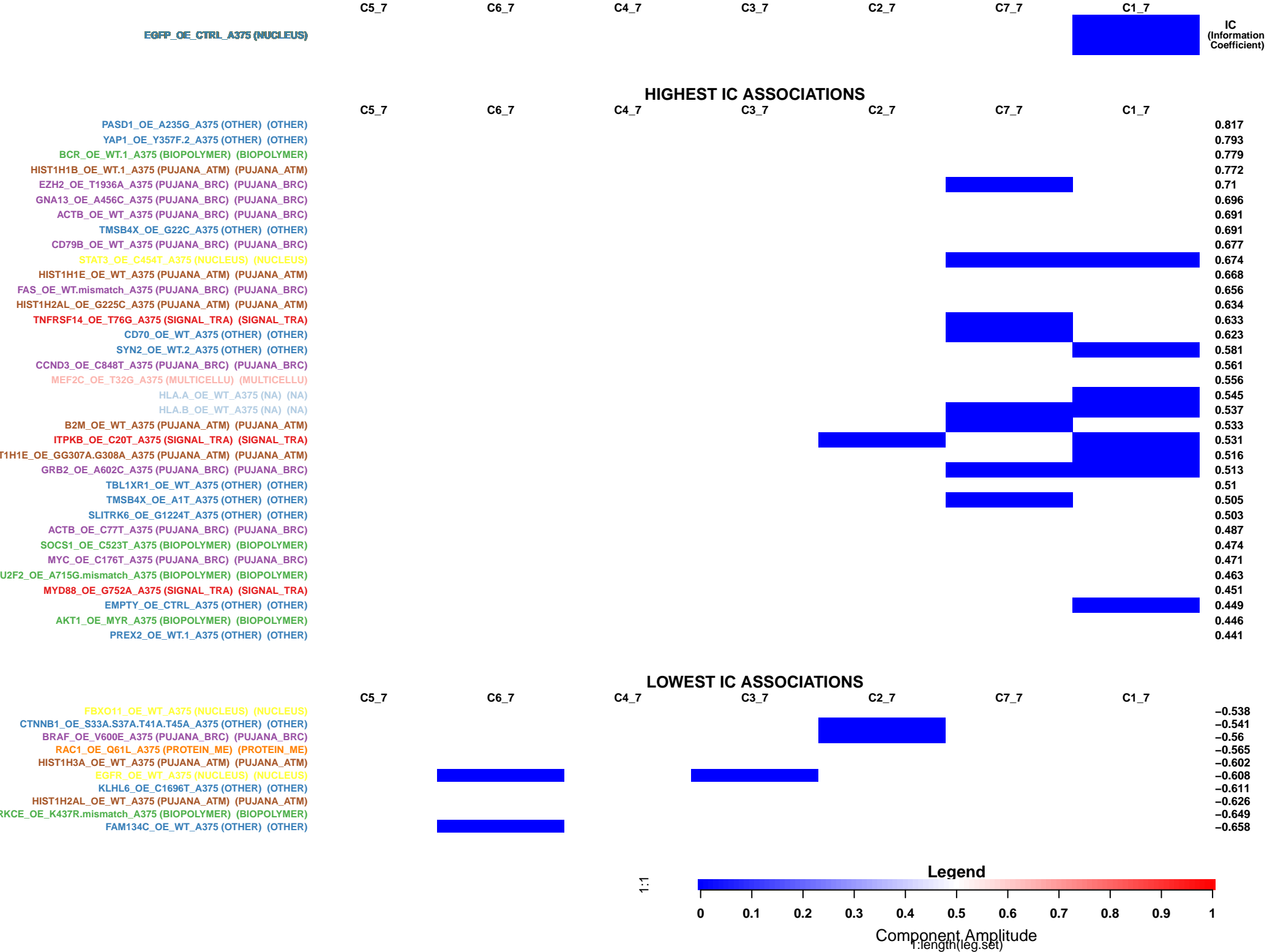
HIGHEST IC ASSOCIATIONS



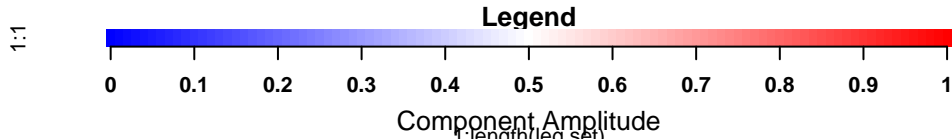
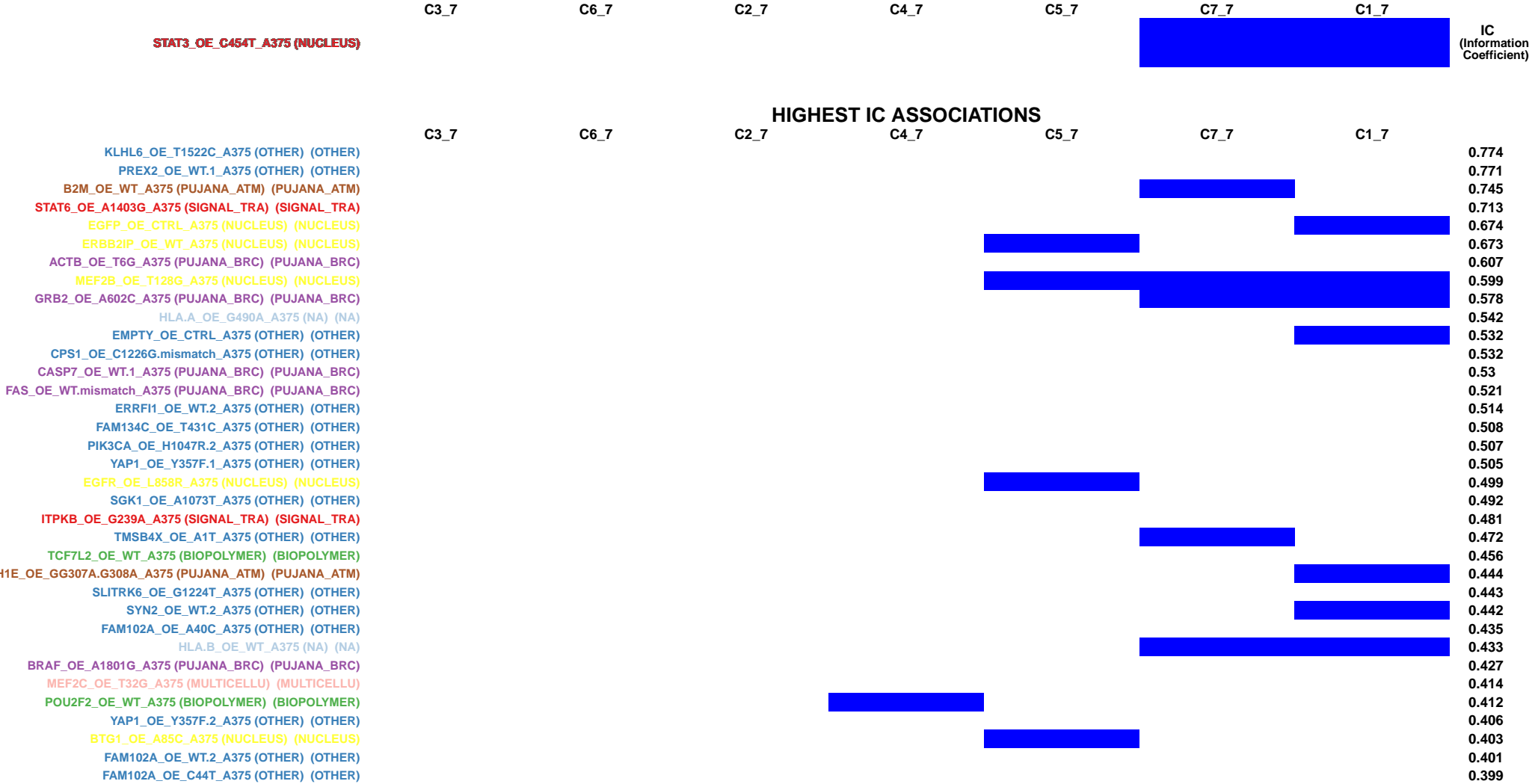
LOWEST IC ASSOCIATIONS



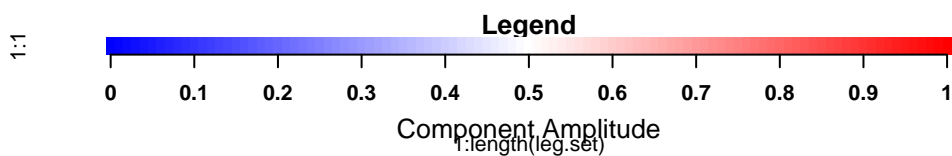
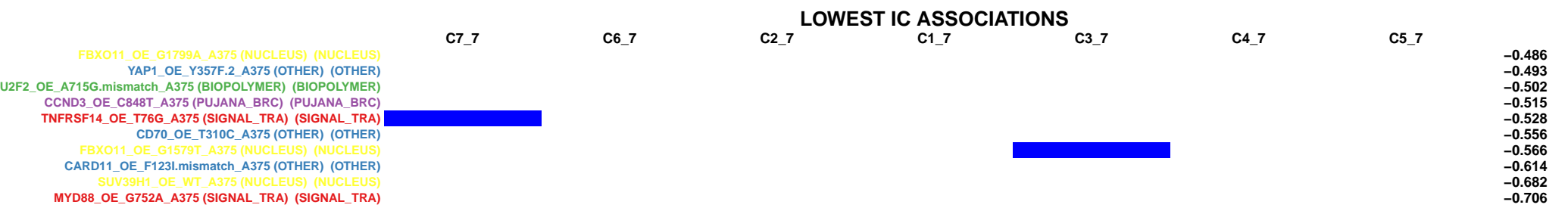
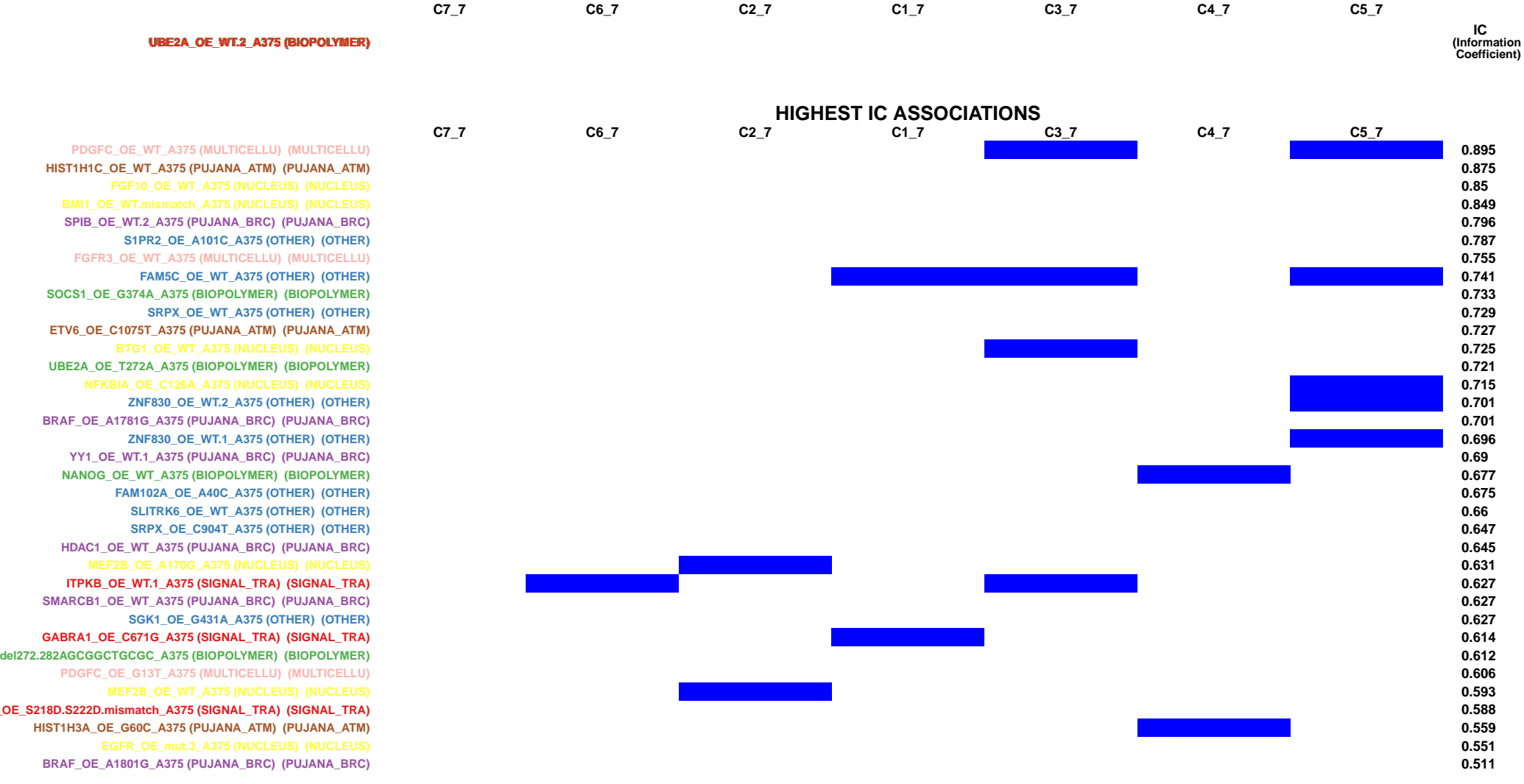
EGFP\_OE\_CTRL\_A375 (NUCLEUS)



STAT3\_OE\_C454T\_A375 (NUCLEUS)



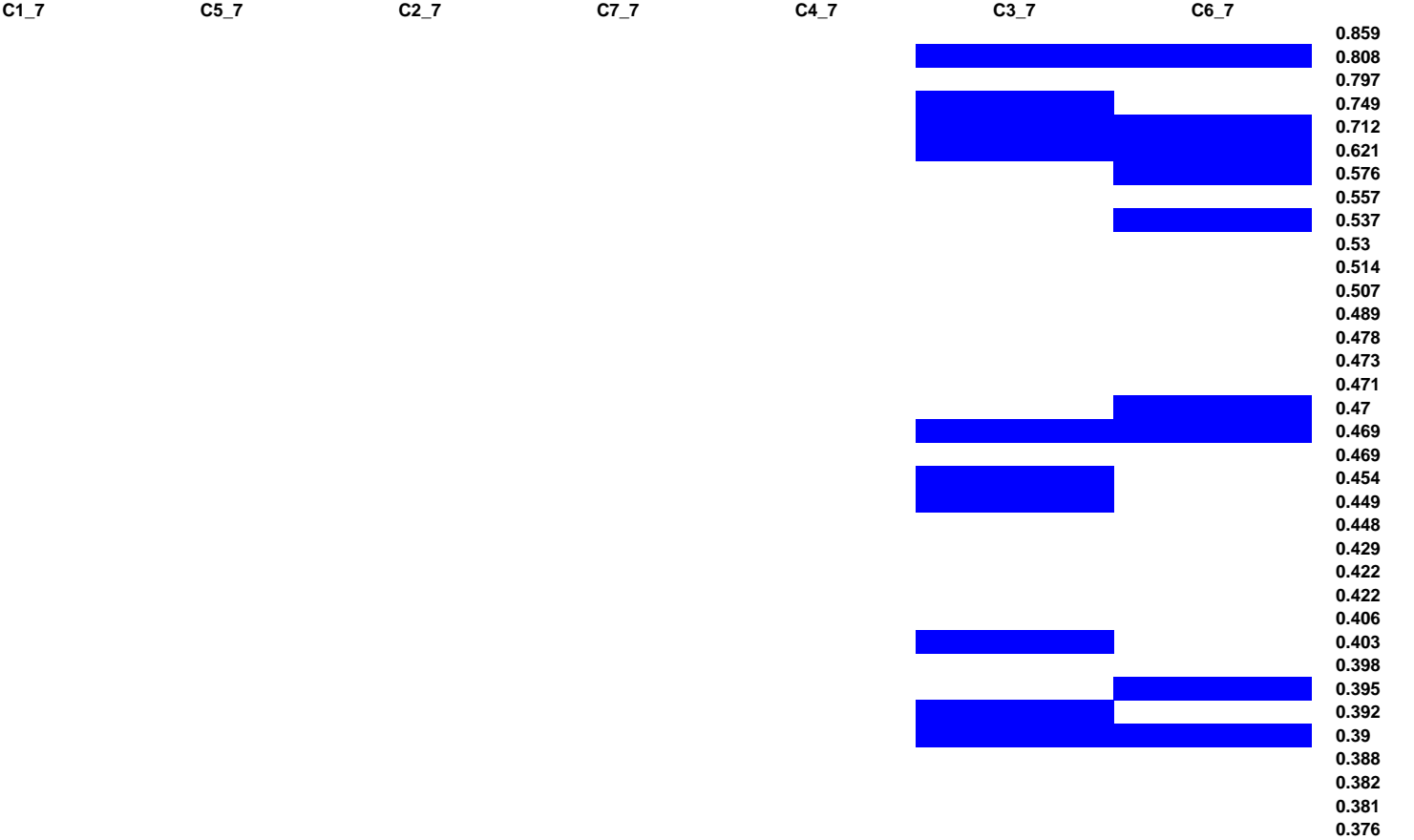
UBE2A\_OE\_WT.2\_A375 (BIOPOLYMER)



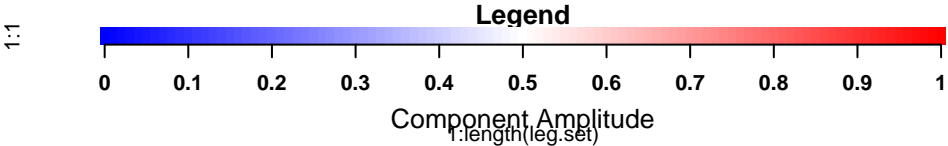
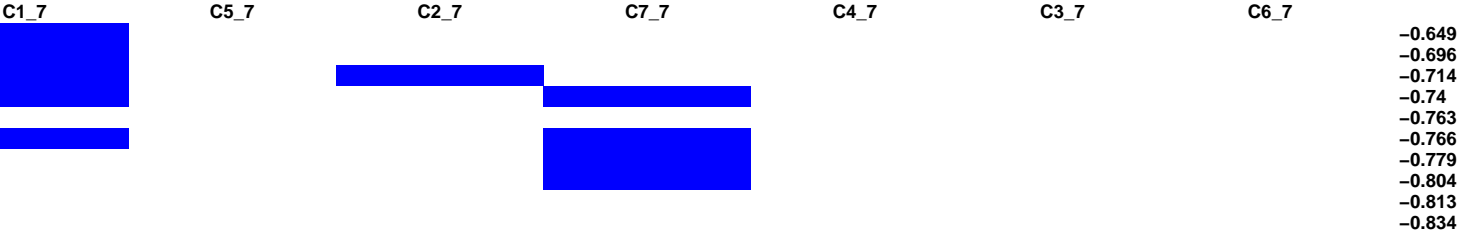
PDGFC\_OE\_G930T\_A375 (MULTICELLU)



HIGHEST IC ASSOCIATIONS

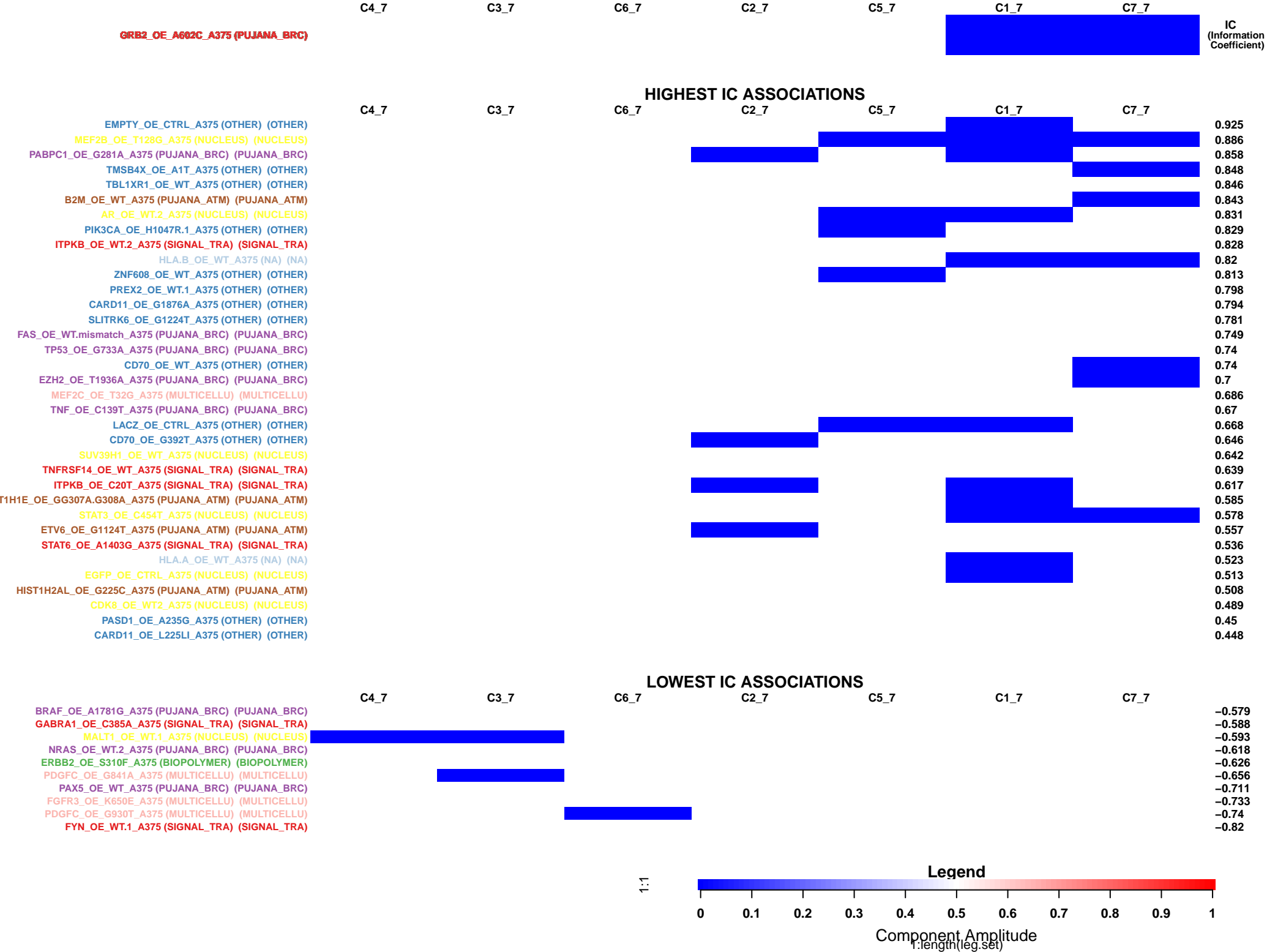


LOWEST IC ASSOCIATIONS

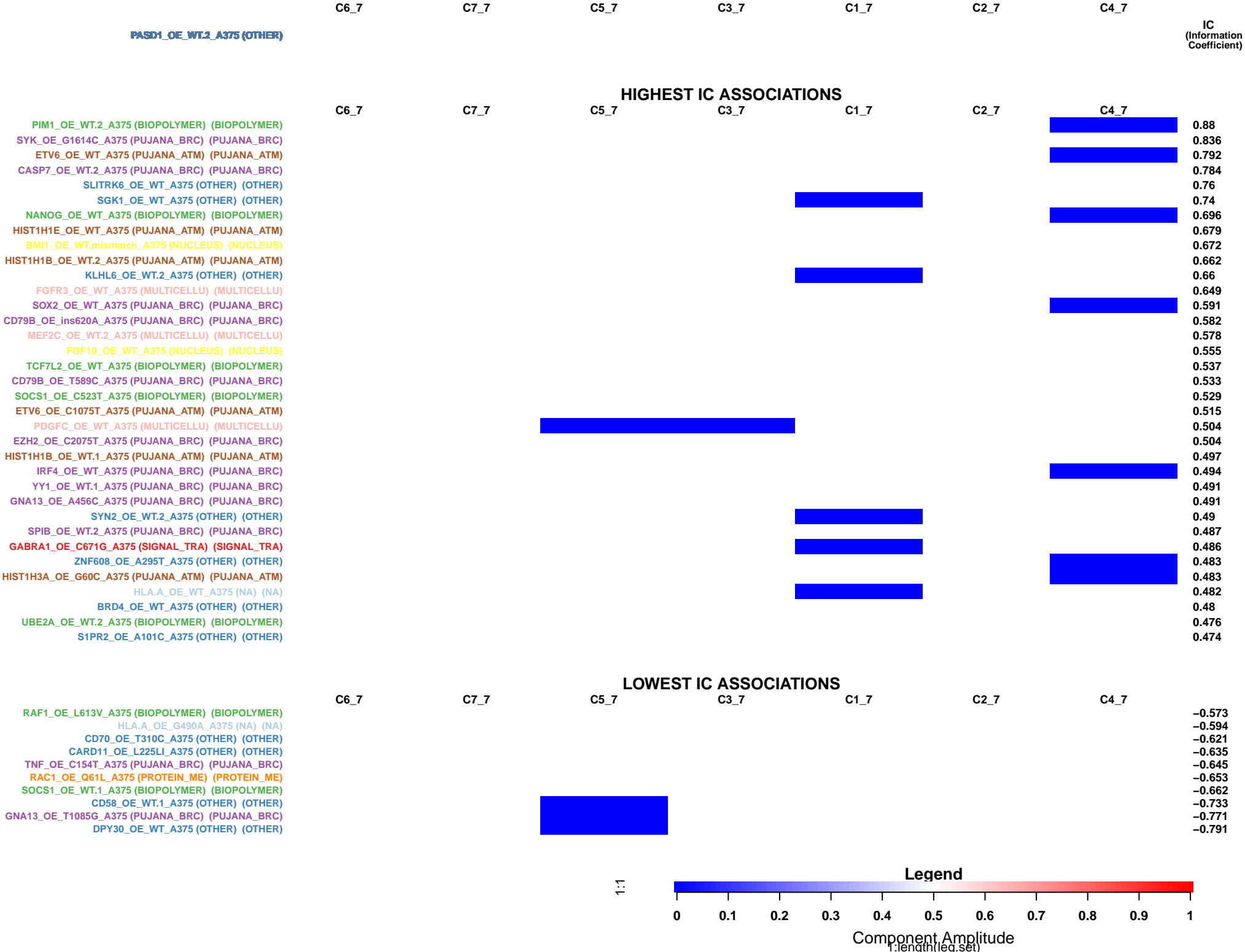




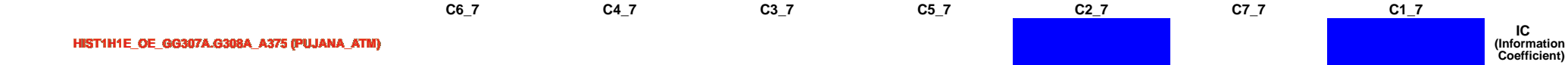
GRB2\_OE\_A602C\_A375 (PUJANA\_BRC)



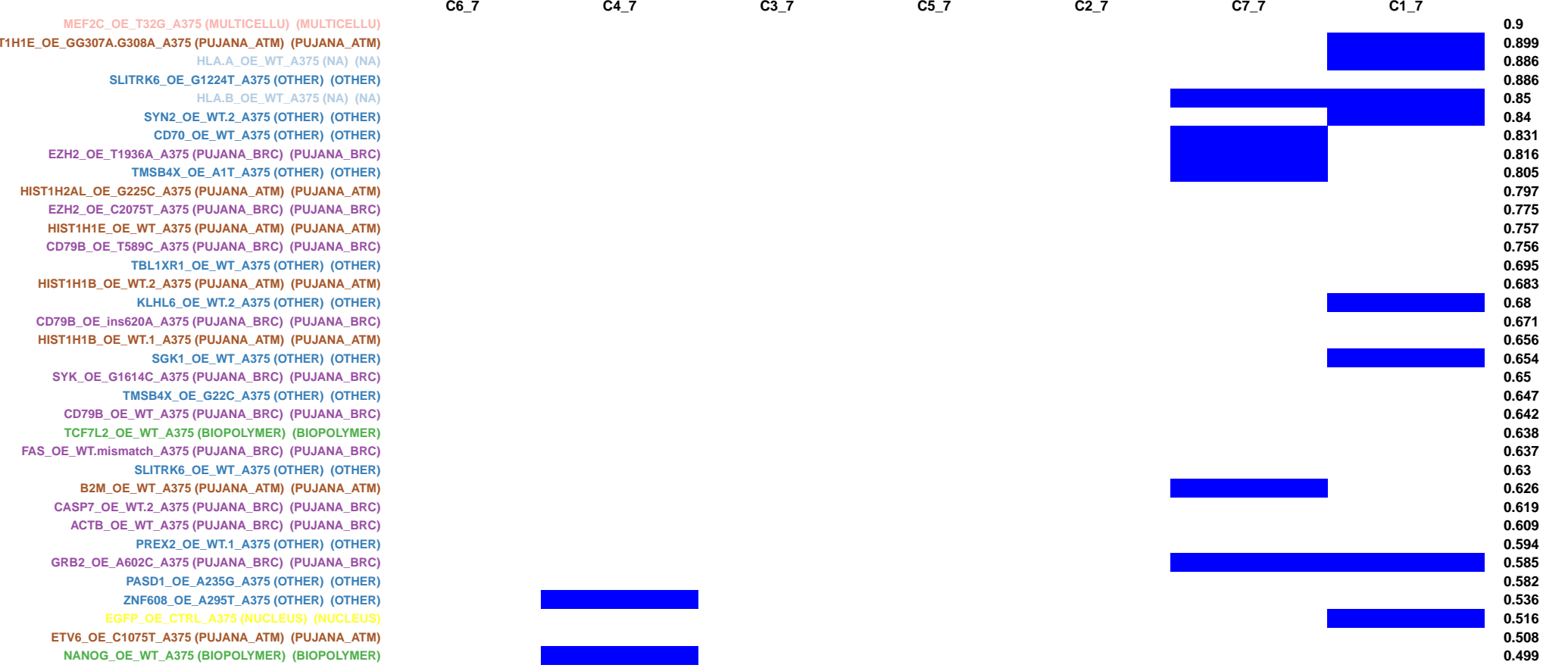
PASD1\_OE\_WT.2\_A375 (OTHER)



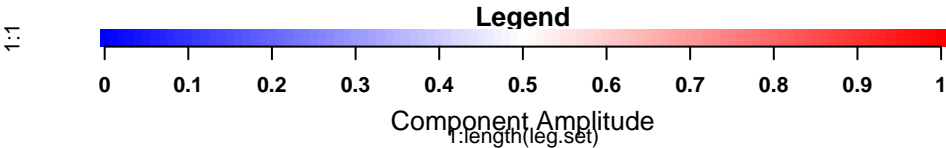
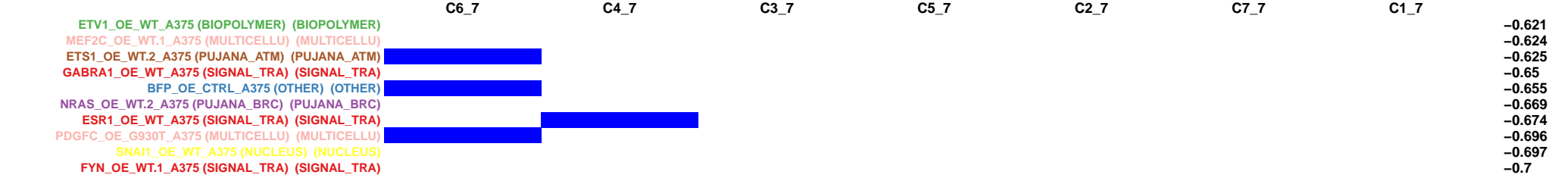
HIST1H1E\_OE\_GG307A.G308A\_A375 (PUJANA\_ATM)



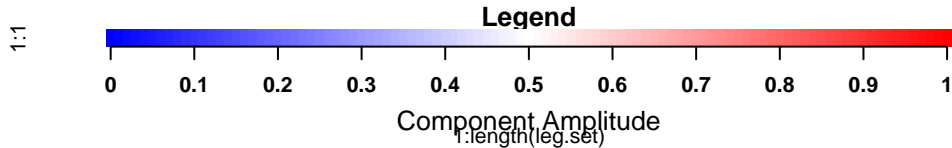
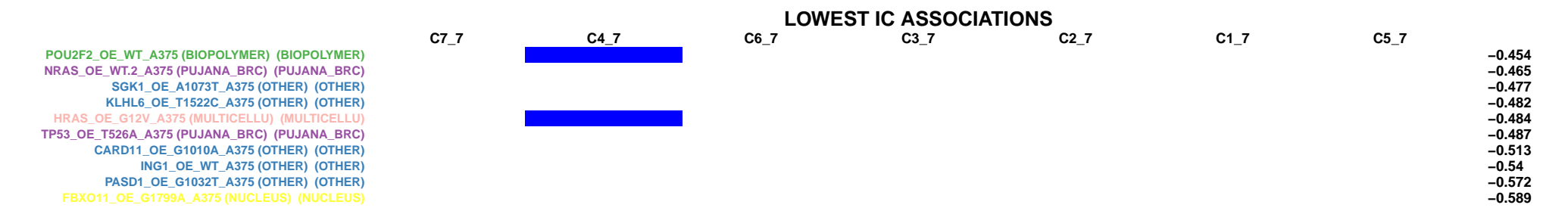
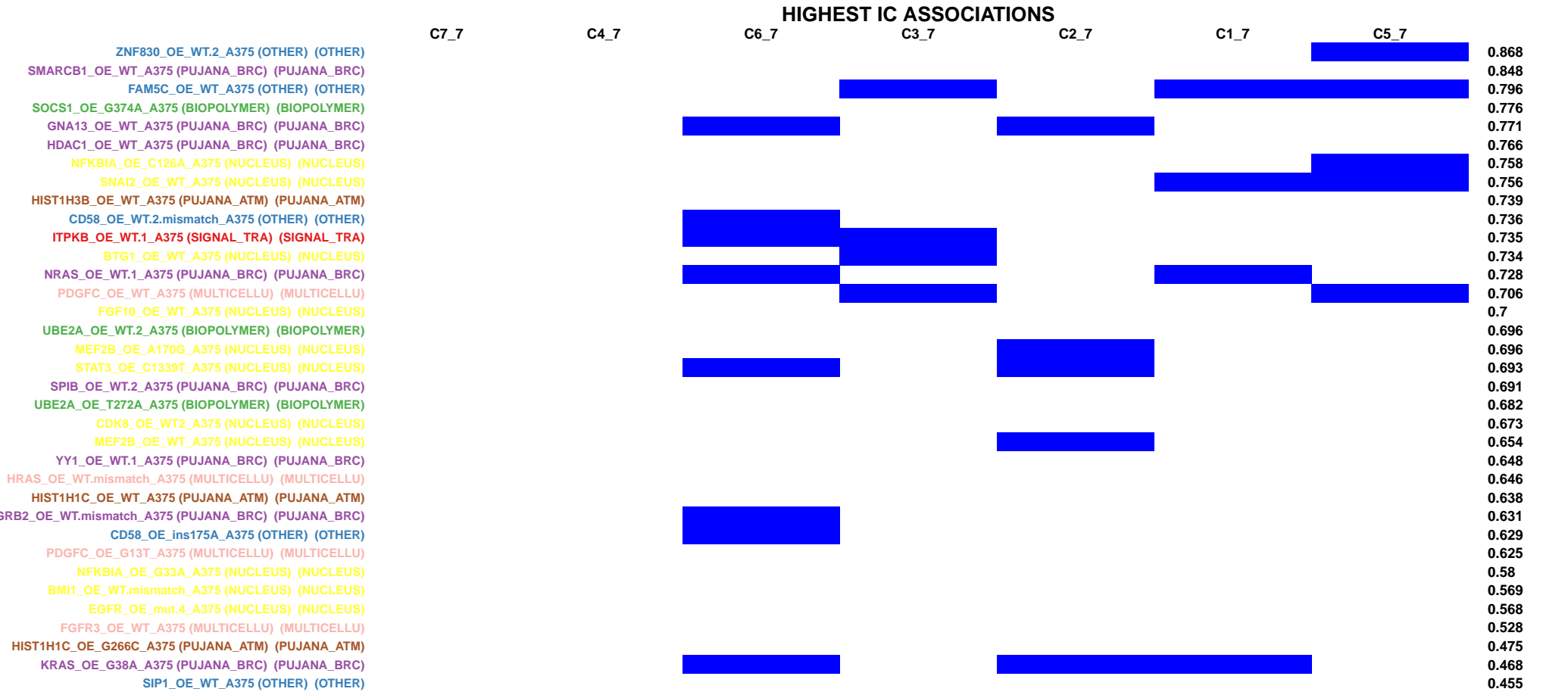
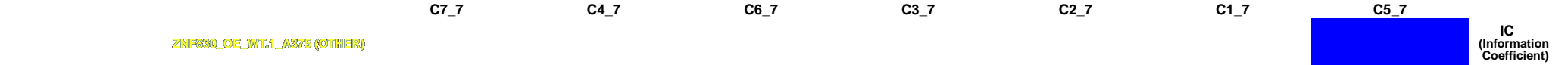
HIGHEST IC ASSOCIATIONS



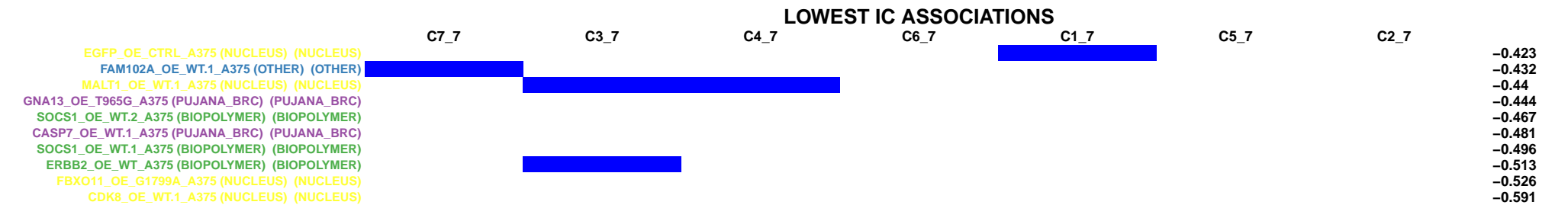
LOWEST IC ASSOCIATIONS



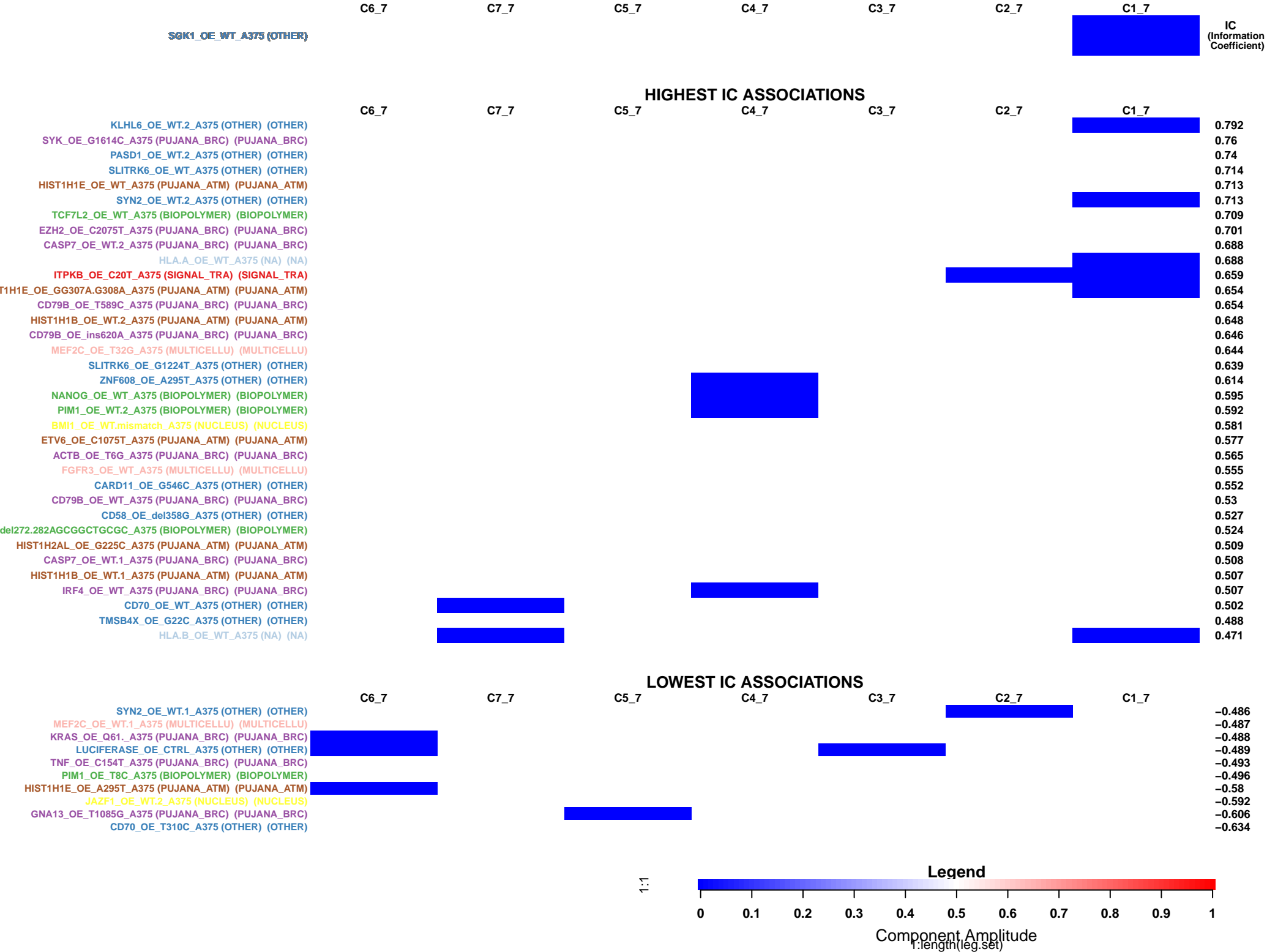
ZNF830\_OE\_WT.1\_A375 (OTHER)



C7_7	C3_7	C4_7	C6_7	C1_7	C5_7	C2_7	IC (Information Coefficient)



SGK1\_OE\_WT\_A375 (OTHER)



## 1:1

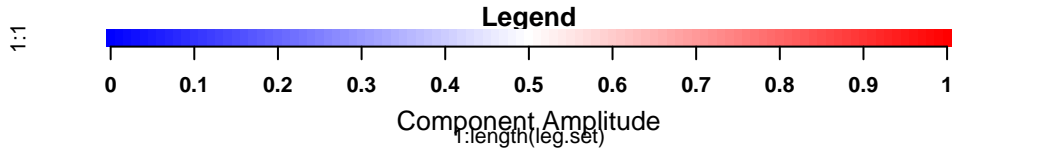
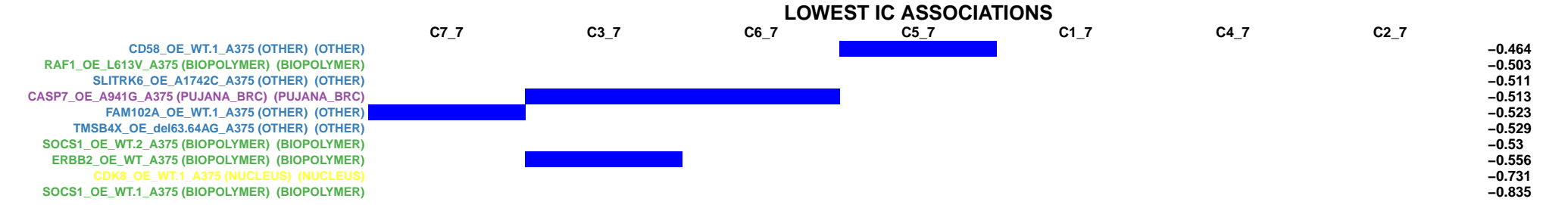
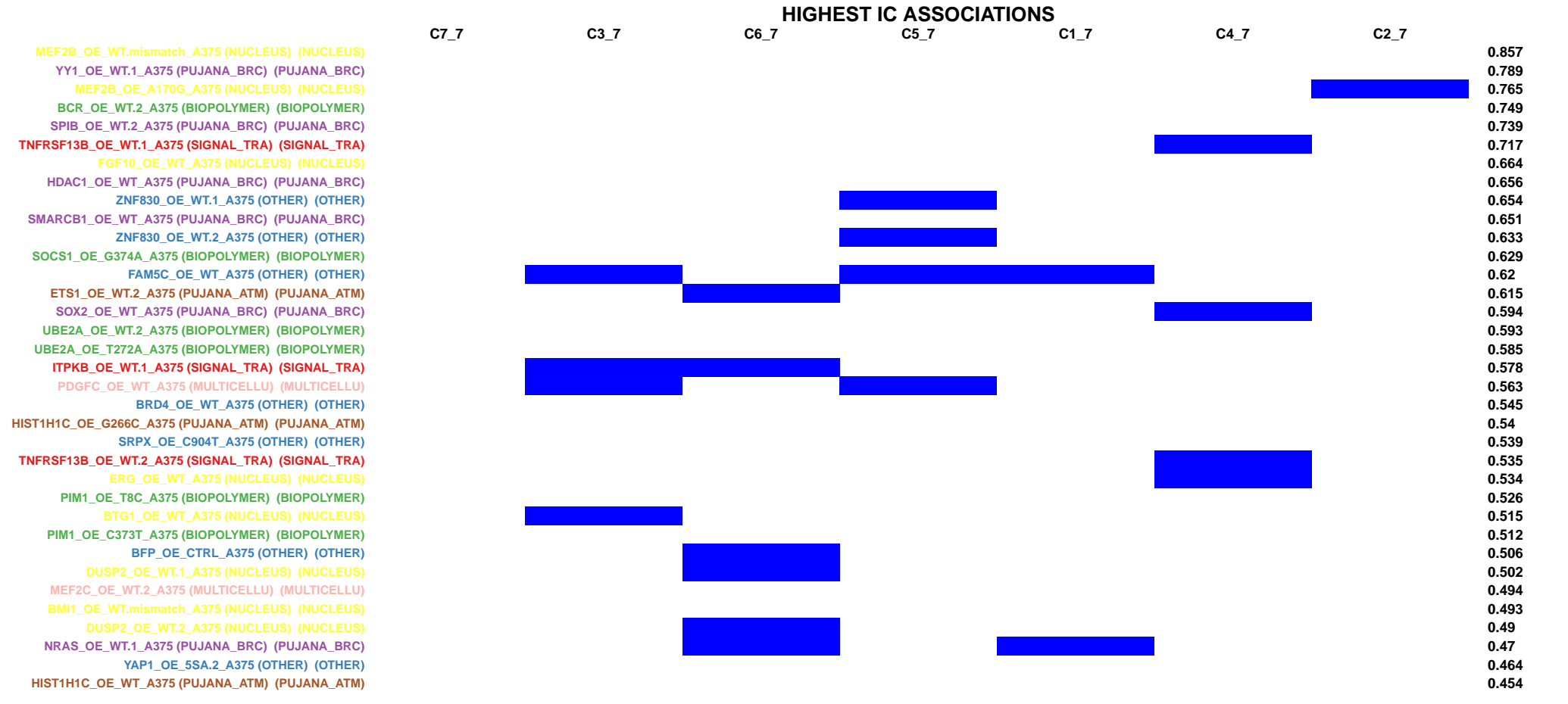
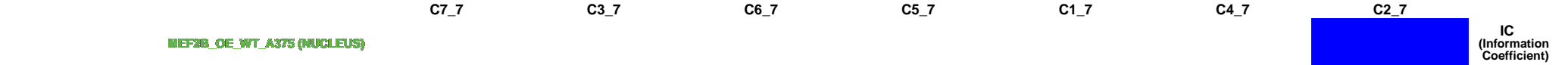


## 1:1

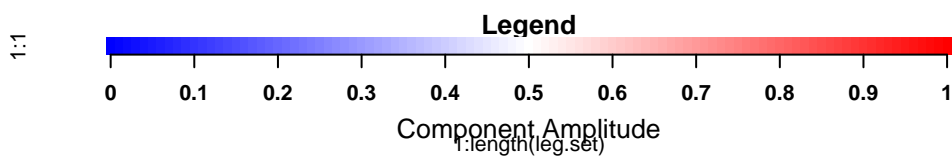
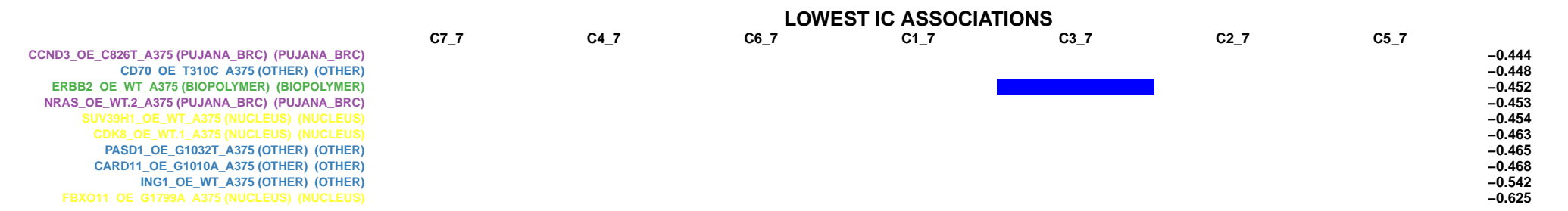
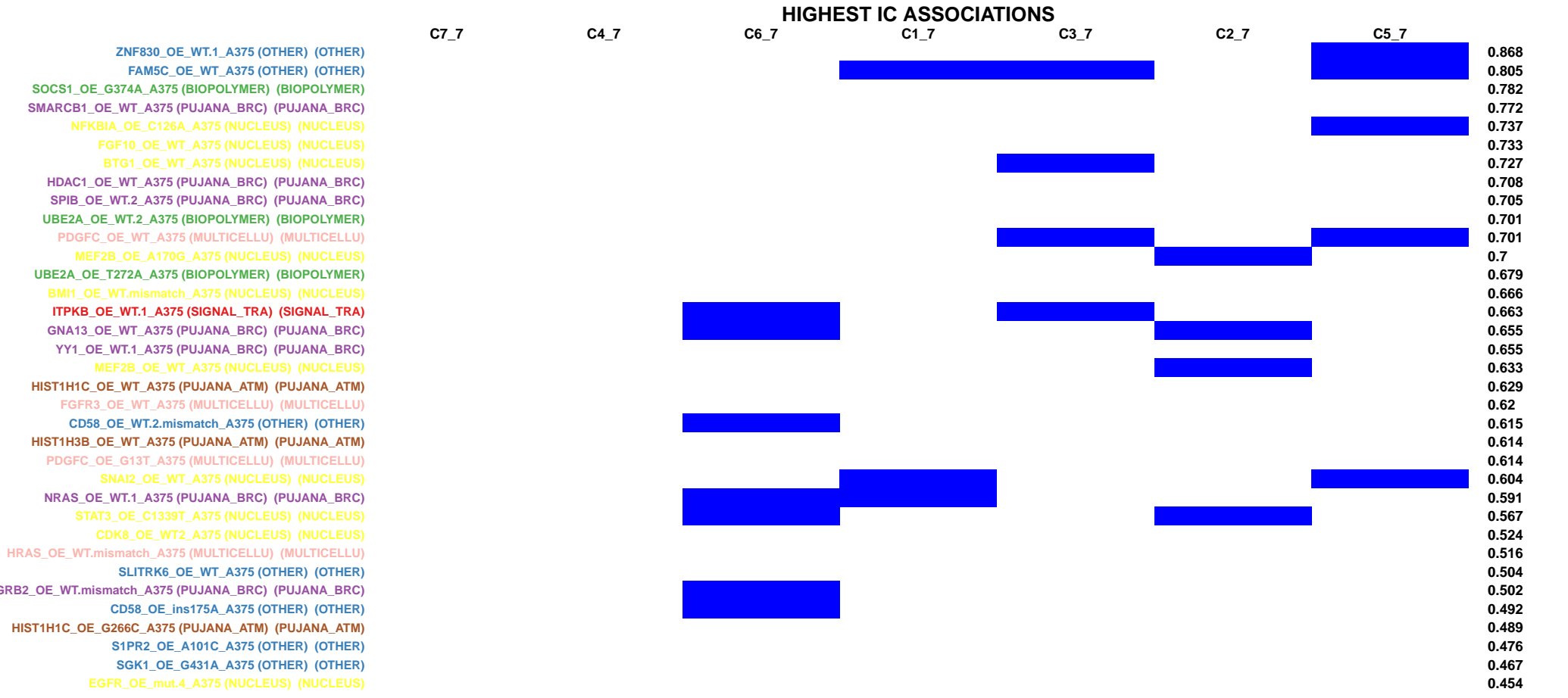
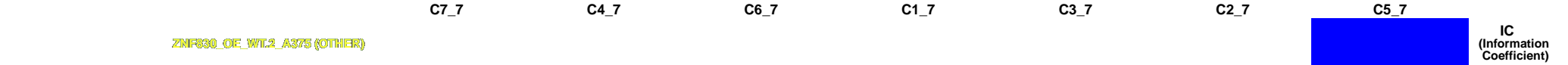




MEF2B\_OE\_WT\_A375 (NUCLEUS)



ZNF830\_OE\_WT.2\_A375 (OTHER)



NOTCH1\_OE\_ICN1.1\_A375 (NUCLEUS)

