

IC / CIC

0.44

0.44

	CIC	p-val	FDR
APC.MC_MUT 5q21-q22	0.49	<5.6e-06	0
ITGBL1_AMP 13q33	0.47	4.5e-05	0.036
RASA3_AMP 13q34	0.45	6.2e-05	0.036
MCF2L-AS1_AMP 13q34	0.45	6.8e-05	0.036
MIR2681_AMP 13	0.45	6.8e-05	0.036
MIR4705_AMP 13	0.45	6.8e-05	0.036
PCCA-AS1_AMP 13q32.3	0.45	6.8e-05	0.036
NALCN-AS1_AMP 13q33.1	0.45	6.8e-05	0.036
METTL21C_AMP 13q33.1	0.45	6.8e-05	0.036
F7_AMP 13q34	0.45	6.8e-05	0.036
F10_AMP 13q34	0.45	6.8e-05	0.036
NALCN_AMP 13q32.3	0.45	6.8e-05	0.036
FGF14-IT1_AMP 13q33.1	0.45	6.8e-05	0.036
LOC283481_AMP	0.45	6.8e-05	0.036
TPP2_AMP 13q32-q33	0.45	6.8e-05	0.036
TMTC4_AMP 13q32.3	0.45	6.8e-05	0.036
ABHD13_AMP 13q33.2	0.45	6.8e-05	0.036
A2LD1_AMP	0.45	6.8e-05	0.036
PROZ_AMP 13q34	0.45	6.8e-05	0.036
FGF14_AMP 13q34	0.44	8.5e-05	0.036
FAM106CP_AMP 17p11.2	0.41	0.00022	0.036
USP32P1_AMP 17p11.2	0.41	0.00022	0.036
CCDC144A_AMP 17p11.2	0.41	0.00022	0.036
MIR3169_AMP 13	0.4	0.00038	0.036
DIAPH3-AS1_AMP 13q21.2	0.4	0.00038	0.036
DIAPH3-AS2_AMP 13q21.2	0.4	0.00038	0.036
PCDH20_AMP 13q21	0.4	0.00038	0.036
TDRD3_AMP 13q14.3	0.4	0.00038	0.036
DIAPH3_AMP 13q21.2	0.4	0.00038	0.036
SERP2_AMP 13q14.11	0.4	0.00041	0.036

SW480 LARGE INTESTINE
C8B01 LARGE INTESTINE
DLD1 LARGE INTESTINE
LTV120 OVARY
T0121 LARGE INTESTINE
COL020S LARGE INTESTINE
HUT100 SMALL INTESTINE
H155 LARGES INTOMACH
SKCO1 LARGE INTESTINE
N6201 LARGE INTESTINE
SW437 LARGE INTESTINE
SW433 OESOPHAGUS
OES3 OESOPHAGUS
LOYO LUNG
L101 LUNG
C349 LUNG
OVISE OVARY
OVISE OVARY
OWM1A OVARY
A205 SKIN
CPAC1 PANCREAS
RKO LARGE INTESTINE
PANC1 PANCREAS
PANC2 PANCREAS
PANC3 PANCREAS
JHSC01 OESOPHAGUS
C0707A OVARY
PANC013 PANCREAS
PANC014 PANCREAS
LS411N LARGE INTESTINE
T05 OESOPHAGUS
HEC1A ENDOMETRIUM
K562 SKIN
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K1031

TARGET: BETA-CATENIN-REPORTER



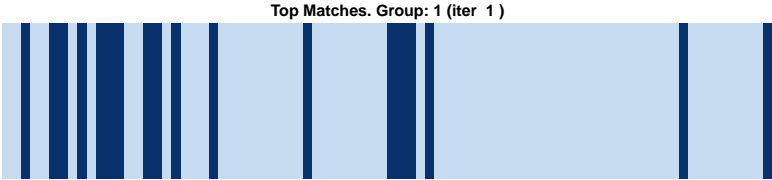
IC / CIC

SEED: CTNNB1.MC MUT
SUMMARY SEED:



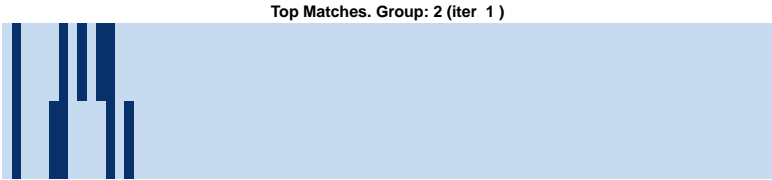
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FAM106CP_AMP 17p11.2
USP32P1_AMP 17p11.2
CCDC144A_AMP 17p11.2



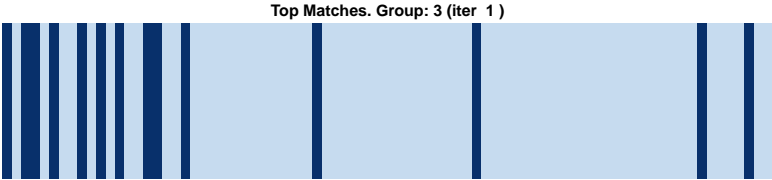
CIC	p-val	FDR
0.41	0.00022	0.036
0.41	0.00022	0.036
0.41	0.00022	0.036

RASA3_AMP 13q34
SERP2_AMP 13q14.11



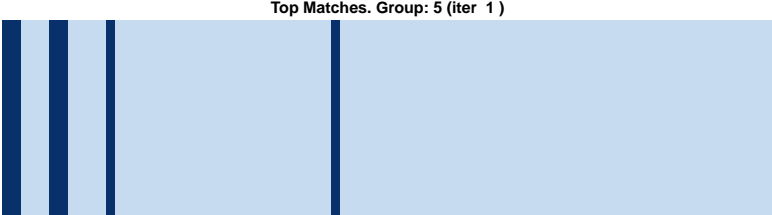
CIC	p-val	FDR
0.45	6.2e-05	0.036
0.4	0.00041	0.036

APC.MC_MUT 5q21-q22



CIC	p-val	FDR
0.49	<5.6e-06	0

MIR3169_AMP 13
DIAPH3-AS1_AMP 13q21.2
DIAPH3-AS2_AMP 13q21.2
PCDH20_AMP 13q21
TDRD3_AMP 13q14.3
DIAPH3_AMP 13q21.2



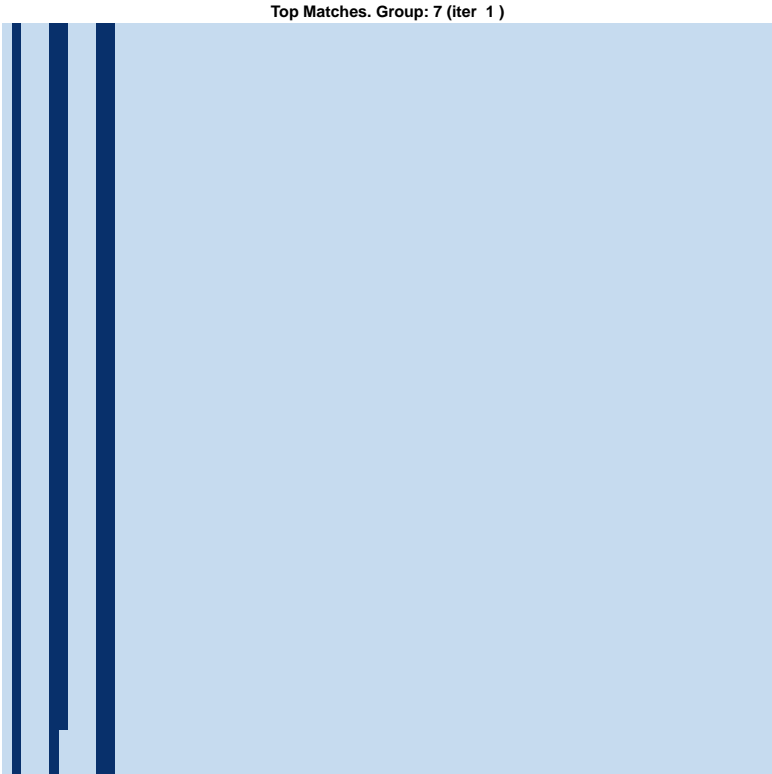
CIC	p-val	FDR
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0.4	0.00038	0.036
0.4	0.00038	0.036
0.4	0.00038	0.036
0.4	0.00038	0.036
0.4	0.00038	0.036

ITGBL1_AMP 13q33



CIC	p-val	FDR
0.47	4.5e-05	0.036

MCF2L-AS1_AMP 13q34
MIR2681_AMP 13
MIR4705_AMP 13
PCCA-AS1_AMP 13q32.3
NALCN-AS1_AMP 13q33.1
METTL21C_AMP 13q33.1
F7_AMP 13q34
F10_AMP 13q34
NALCN_AMP 13q32.3
FGF14-IT1_AMP 13q33.1
LOC283481_AMP
TPP2_AMP 13q32-q33
TMTC4_AMP 13q32.3
ABHD13_AMP 13q33.2
A2LD1_AMP
PROZ_AMP 13q34
FGF14_AMP 13q34



CIC	p-val	FDR
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.45	6.8e-05	0.036
0.44	8.5e-05	0.036

IC / CIC

0.44

0.49

0.61

	CIC	p-val	FDR
ITGBL1_AMP 13q33	0.49	6.8e-05	0.0082
RASA3_AMP 13q34	0.46	9e-05	0.0082
LINC00462_AMP 13q14.2	0.45	0.00011	0.0082
TPT1-AS1_AMP 13q14.13	0.45	0.00011	0.0082
MIR1297_AMP 13	0.45	0.00011	0.0082
MIR320D1_AMP 13q14.11	0.45	0.00011	0.0082
MIR3170_AMP 13	0.45	0.00011	0.0082
MIR4305_AMP 13	0.45	0.00011	0.0082
MIR3665_AMP 13	0.45	0.00011	0.0082
MIR3613_AMP 13	0.45	0.00011	0.0082
RBM26-AS1_AMP 13q31.1	0.45	0.00011	0.0082
OR7E37P_AMP 13q14.11	0.45	0.00011	0.0082
VWA8-AS1_AMP 13q14.11	0.45	0.00011	0.0082
CPB2-AS1_AMP 13q14.13	0.45	0.00011	0.0082
MIR4500_AMP 13q31.2	0.45	0.00011	0.0082
MIR4703_AMP 13	0.45	0.00011	0.0082
TPTE2P5_AMP 13q14.11	0.45	0.00011	0.0082
MIR5693_AMP 13	0.45	0.00011	0.0082
MIR5006_AMP 13	0.45	0.00011	0.0082
LINC00563_AMP 13q14.13	0.45	0.00011	0.0082
LINC00441_AMP 13q14.2	0.45	0.00011	0.0082
RNU6-66_AMP	0.45	0.00011	0.0082
RNU6-69_AMP	0.45	0.00011	0.0082
MED4-AS1_AMP 13q14.2	0.45	0.00011	0.0082
DLEU7-AS1_AMP 13q14.3	0.45	0.00011	0.0082
LINC00331_AMP 13q31.1	0.45	0.00011	0.0082
LINC00351_AMP 13q31.1	0.45	0.00011	0.0082
LINC00353_AMP 13q31.3	0.45	0.00011	0.0082
LINC00366_AMP 13q13.3	0.45	0.00011	0.0082
LINC00379_AMP 13q31.3	0.45	0.00011	0.0082

[illegible]

IC / CIC

0.44

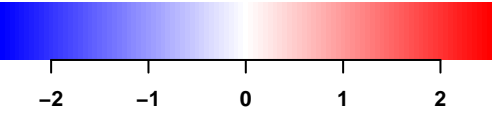
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Standardized Target Profile

REVEALER – Results

TARGET: BETA-CATENIN-REPORTER

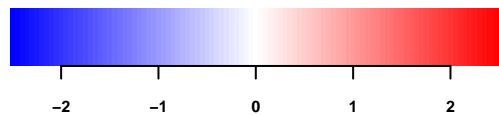
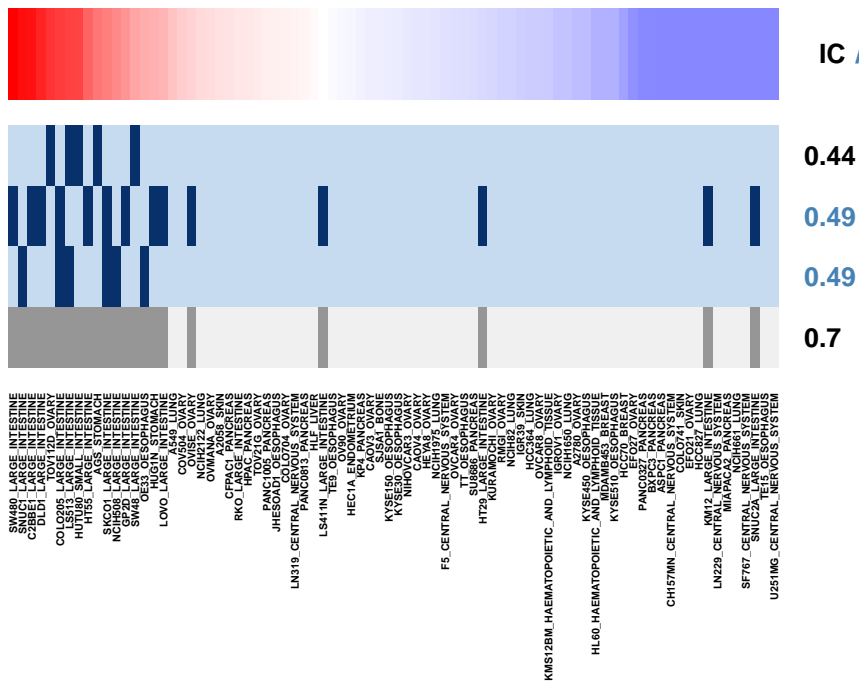
IC / CIC

SEED: CTNNB1.MC_MUT

ITERATION 1: APC.MC_MUT 5q21-q22

ITERATION 2: ITGBL1_AMP 13q33

FINAL SUMMARY



Standardized Target Profile