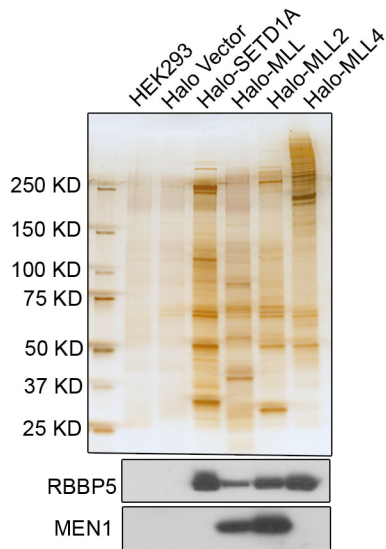


A



B

MudPIT analysis of Halo-MLL Complexes

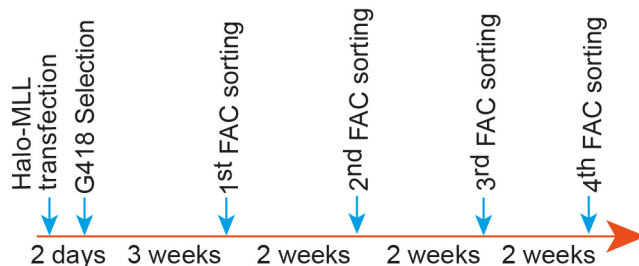
dNSAF	Vector	Halo-MLL1
MLL1	0	0.00749
RBBP5	0	0.000113
ASH2L	0	0.0000969
DPY30	0	0.00276
WDR5	0	0.00009114
HCFC1	0	0.0000374
Menin	0	0.000973
CXXC1	0	0
WDR82	0	0

E Enriched pathways in shRNA screening for MLL degradation

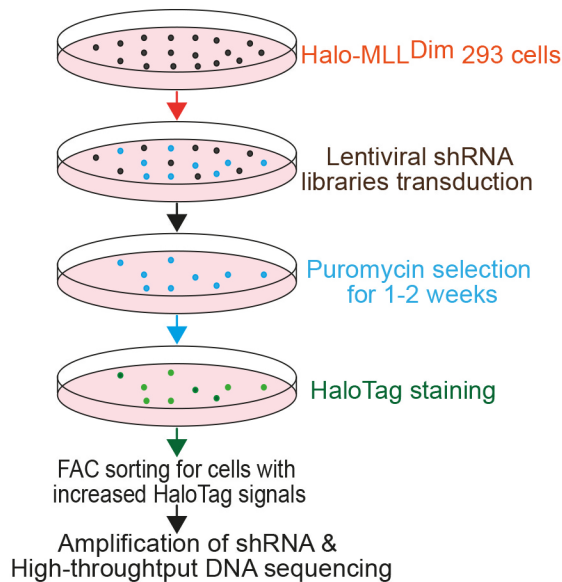
Molecular Function	Fold Enrichment	P value
Interleukin-1 receptor activity	> 5	7.50E-03
Cytokine receptor activity	> 5	1.53E-04

Biological Process	Fold Enrichment	P value
Regulation of immune response	2.46	3.59E-02
Immune response	2.43	1.38E-04

C



D



F

