modERN analysis call

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Re-cap

- In fly, most dataset can recover less than 10% peaks of the standard pipeline (STD) when using mockIP as input (MOCK)
- The possible explanation could be a mixture of:
 - open chromatin regions were enriched in the mock-IP experiment? For each particular factor, some of these regions might have real enrichment while some are not.
 - The Goat-V antibody has significant non-specificity issue? WB/IF shows good results. Should also see how the worm *MOCK* peaks look like.
- We want to have a *revised-STD* pipeline
 - Still follow the *STD* pipeline, but adjust the cutoff for calling a peak in order to maximize the number of peaks that overlaps with the *MOCK* pipeline.
 - We hope the cutoff could be uniform across datasets so that the tagging strategy and the *STD* pipeline can be widely applied.

Most of modERN data recovered less than 10% of *STD* peaks from the *MOCK* pipeline

Recovered peaks from mockIP



Ratio (# of MOCK / # of STD)

Picked up ten datasets representing different peak recovery rate to test

				Dataset	# STD	# MOCK	#(MOCK_in_STD)	Ratio(MOCK/STD)	Ratio(MOCK_in_ST D/MOCK)
	Recovered peaks from mockIP			tio-GFP_tio_E4-24	3733	21	7	0.006	0.333
				disco-GFP_disco_E0-24	2250	14	7	0.006	0.500
				ems-GFP_ems_E0-24	3963	33	6	0.008	0.182
۲ 40	Π			ab-GFP_ab_E0-24	1546	70	25	0.045	0.357
				Att-2-GFP_Att-2_E0-24	3067	34	11	0.011	0.324
g –				en-GFP_en_E0-16	2259	32	10	0.014	0.313
Prequency 3				pho-GFP_pho_E0-24	4204	78	38	0.019	0.487
				pnt-MIMIC_pnt_E0-24	3543	95	25	0.027	0.263
				gro-GFP_gro_E0-24	4533	484	244	0.107	0.504
				Fer1-GFP_Fer1_E6-24	7477	2179	1864	0.291	0.855
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	0.0	0.2 0.4	0.6	0.8 1.0					

Ratio (# of MOCK / # of STD)

Picked up three candidates with modENCODE matched data to double check the *MOCK* pipeline

Recovered peaks from mockIP



Ratio (# of MOCK / # of STD)

Overlap between modENCODE, *STD* and *MOCK* peaks varied.

modENCODE	modERN	modENCO DE uniq	modENCO DE common	modERN uniq	modERN common	# modENCO DE	% modENCO DE common	# modERN	% B modERN
SuH	Suh-GFP-MOCK	9529	18	62	22	9547	0.2%	84	21.4%
SuH	Suh-GFP-STD	8172	1375	2080	1486	9547	16.8%	3566	38.6%
SuHw	SuHw-GFP- MOCK	3330	1662	235	2046	4992	49.9%	2281	72.9%
SuHw	Suhw-GFP-STD	3202	1790	3115	2179	4992	55.9%	5294	33.8%
Trl	Trl-MIMIC-MOCK	11627	81	159	84	11708	0.7%	243	33.3%
Trl	Trl-MIMIC-STD	10325	1383	2919	1700	11708	13.4%	4619	29.9%

Trl: *MOCK* removed most of modENCODE and *STD* peaks



Su(Hw): Good consistency



Su(H): *MOCK* removed most of modENCODE and *STD* peaks



Open question

• Should we just go ahead with the *STD* pipeline?