

Supplementary Table 3. Mass spectrometry total spectral counts for PR association with Pol III holoenzyme subunits.

Gene	UCD4				UCD65			
	IP:		IP:		IP:		IP:	
	IgG	IgG	PR	PR	IgG	IgG	PR	PR
POLR3A	0	0	70 ¹	95	0	0	22	16
POLR3B	0	0	33	49	0	0	8	8
POLR3C	0	0	22	29	0	0	4	6
POLR3D	0	0	12	13	0	0	4	9
POLR3E	0	0	20	37	0	0	5	3
POLR3F	0	0	8	8	0	0	0	0
POLR3GL	0	0	0	2	0	0	0	0
POLR3H	0	0	0	1	0	0	0	0
POLR3K	0	0	4	4	0	0	0	0
CRCP	0	0	8	14	0	0	2	0
POLR1C	0	0	13	19	0	0	3	4
POLR1D	0	0	2	2	0	0	0	0
POLR2E	0	0	4	5	0	0	0	0
POLR2H	1	0	0	1	0	0	2	0
POLR2L	0	0	0	1	0	0	0	0

Duplicate UCD4 and UCD65 tumors grown with E2 plus MPA were analyzed using ActiveMotif's (Carlsbad, CA) rapid immunoprecipitation followed by mass spectrometry (RIME) services for PR. Total spectral counts were determined using 95% probability and 2 peptide minimum using Scaffold software (Portland, OR). ¹Gray shade indicates 95% probability of sequence identity.