

Table 1. Expression of PcG in follicular lymphoma

Gene	Mean GC	Mean FL	T-stat	p value
<b><i>PRC1 members</i></b>				
CBX7	0.091	0.776	-6.015	<b>3.8x10<sup>-5</sup></b>
RING1	0.68	0.749	-0.479	0.644
RNF2	-0.888	-1.037	0.259	0.807
PHC1	0.339	-0.265	5.685	<b>8.6x10<sup>-5</sup></b>
PHC2	0.525	0.411	0.544	0.595
PHC3	-1.456	-1.697	0.555	0.596
BMI1	-0.517	-0.452	-0.189	0.854
RNF134	-0.215	-0.972	3.219	<b>0.012</b>
RNF3	-1.121	-0.598	-0.974	0.36
<b><i>PRC2 members</i></b>				
EZH1	-0.208	-0.723	1.858	0.09
EZH2	0.421	-0.841	4.397	<b>0.001</b>
SUZ12	0.743	-0.072	5.844	<b>6.1x10<sup>-5</sup></b>
EED	0.489	-0.165	2.659	<b>0.02</b>

Gene expression values in germinal center (GC) and follicular lymphoma (FL) as reported in Ramamswamy *et al.* (1). Values were obtained from ONCOMINE (<http://www.oncomine.org>). When the p value is shown in red/green, it means the increase/decrease of the expression of that PcG gene in FL vs. GC is significant (p value <0.05).

1. Ramaswamy, S., Tamayo, P., Rifkin, R., Mukherjee, S., Yeang, C. H., Angelo, M., Ladd, C., Reich, M., Latulippe, E., Mesirov, J. P., Poggio, T., Gerald, W., Loda, M., Lander, E. S. & Golub, T. R. (2001) *Proc Natl Acad Sci U S A* **98**, 15149-54.