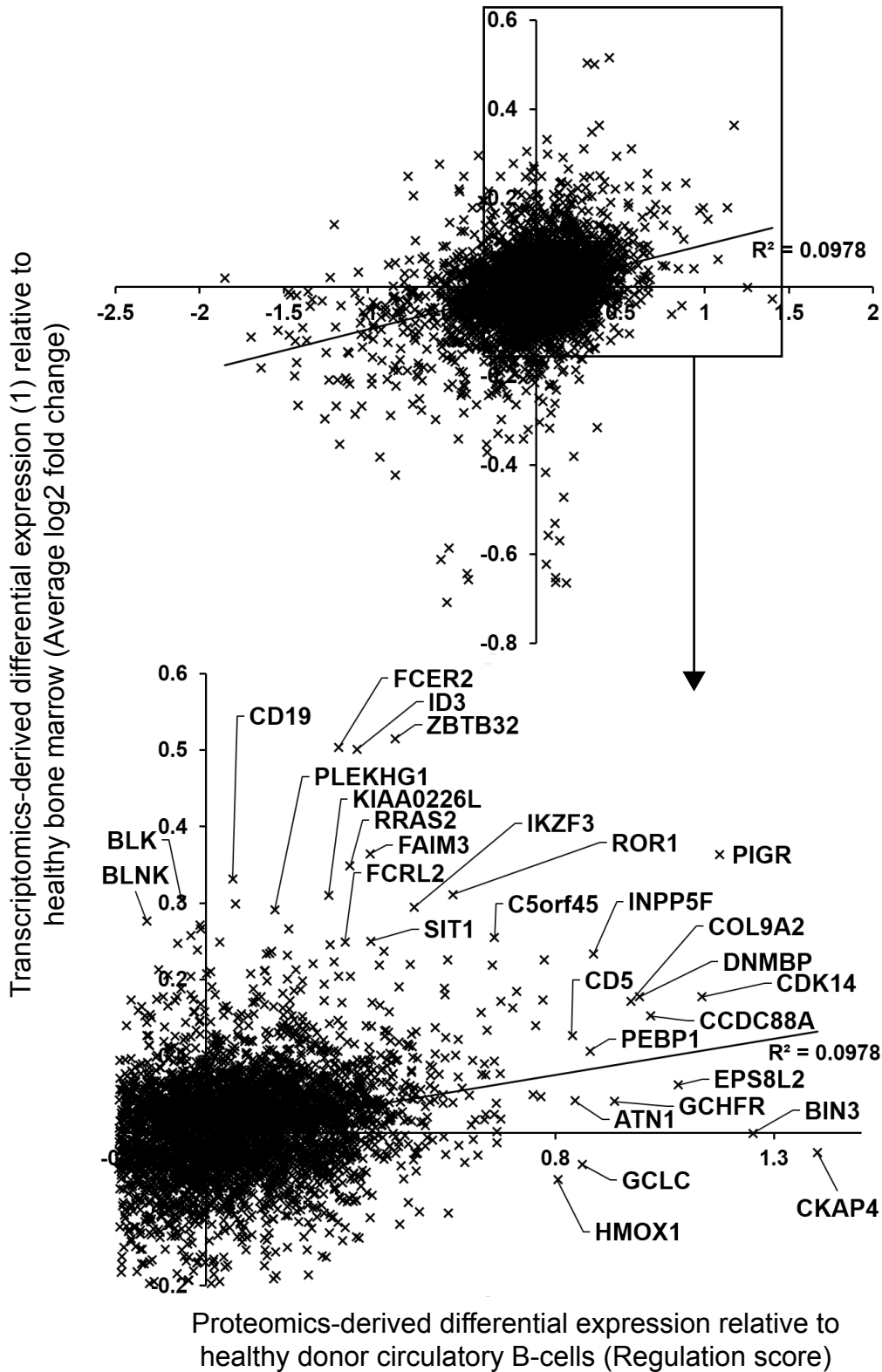


Correlation between transcriptomics (1) and proteomics results for differential gene product expression in CLL vs healthy controls



(1) Haferlach, T., Kohlmann, A., Wieczorek, L., Basso, G., Kronnie, G. T., Bene, M. C., De Vos, J., Hernandez, J. M., Hofmann, W. K., Mills, K. I., Gilkes, A., Chiaretti, S., Shurtleff, S. A., Kipps, T. J., Rassenti, L. Z., Yeoh, A. E., Papenhausen, P. R., Liu, W. M., Williams, P. M., and Foa, R. (2010) Clinical utility of microarray-based gene expression profiling in the diagnosis and subclassification of leukemia: report from the International Microarray Innovations in Leukemia Study Group. *J Clin Oncol* 28, 2529-2537