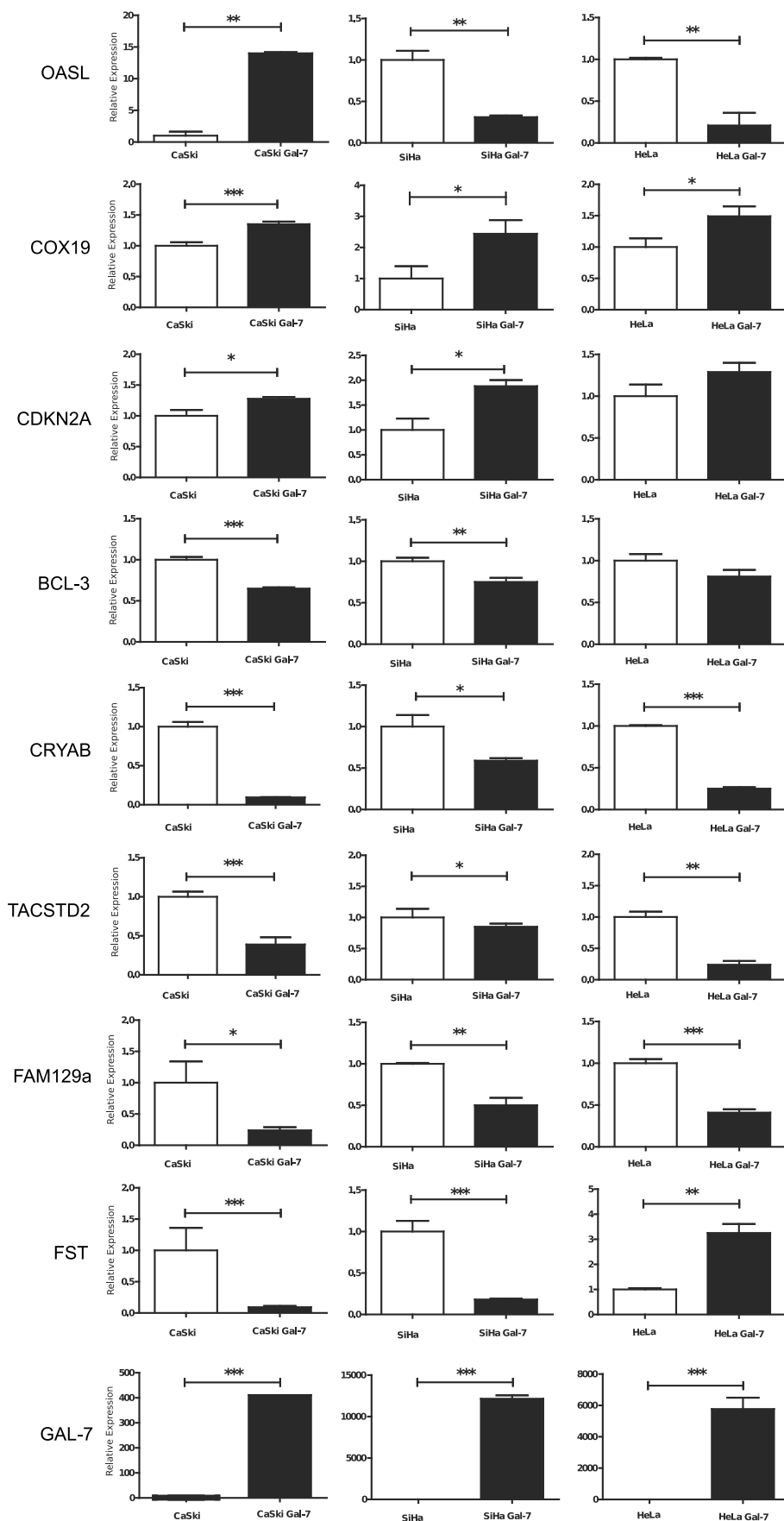




Supplementary Figure 2: transcriptional analysis of the genes identified by the integrative analysis in Gal-7+ CxCa cells. qPCR of OASL, COX19, CDKN2A, BCL-3, CRYAB, TACSTD2, FAM129a, FST and Gal-7 mRNA expression performed in Gal-7-reconstituted and negative CxCa cells (CaSki, SiHa, HeLa). Actin and GAPDH were used as normalization control. (T-Test, with a value of $P < 0.05$. ***= highly significant, **= moderately significant, or *= significant).