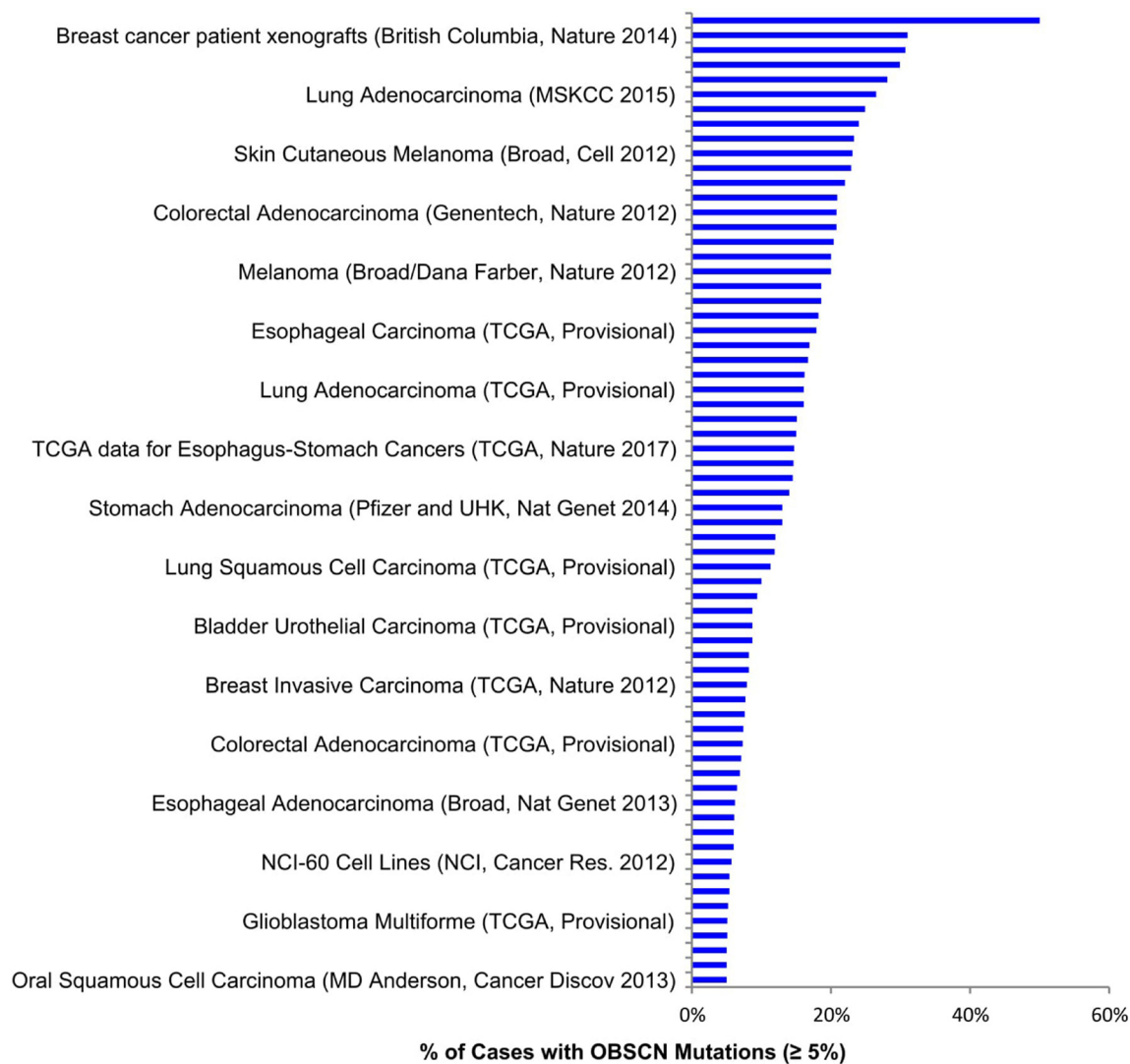


A comprehensive genomic meta-analysis identifies confirmatory role of *OBSCN* gene in breast tumorigenesis

SUPPLEMENTARY MATERIALS



Supplementary Figure 1: Percentage of cases with *OBSCN* gene mutations over various cancer types (> 5%).

Supplementary Table 1: Protein variants of *OBSCN* gene predicted using SIFT and POLYPHEN-2 with reference SNP (rs-id) identifier, gene and protein variants information with SIFT and Polyphen-2 scores. See Supplementary_Table_1