

Figure S3: The FAK phosphorylation data of figure 4C was quantified by densitometry using the ImageJ program. The analysis was done separately for each paired data set (+/- doxycycline) corresponding to the different transfected cell types and normalized with respect to the uninduced (-dox) state. The data was not normalized with respect to total FAK expression, as a direct effect of the induction on FAK expression cannot be ruled out.