# BABRAHAM INSTITUTE

# PIP3 project

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# 1 Role of PTEN/SHIP2 in PI(3,4)P2 metabolism

Let us consider a small part of phosphoinositide signaling pathway consisting of PIP3 and PI(3,4)P2 species:

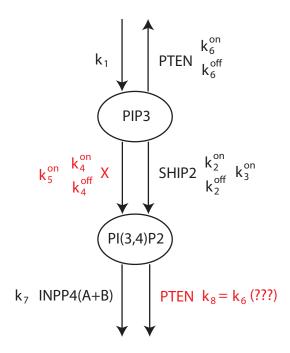


Figure 1: Diagram of PIP3 and PI(3,4)P2 production and degradation with corresponding reaction constant (see next paragraph). X is an unknown phosphatase that can act together with SHIP2. Role of PTEN in PI(3,4)P2 degradation is validated in the next paragraphs.

#### 1.1 Chemical basis

Diagram in Fig. 1 can be described by the following system of chemical reactions:

$$\xrightarrow{k_1} PIP_3 \tag{1}$$

$$PIP_{3} + SHIP_{2} \xrightarrow[k_{2}^{of}]{k_{2}^{of}} PIP_{3} - SHIP_{2}$$
(2)

$$PIP_3-SHIP_2 \xrightarrow{k_3^{on}} PI_{(3,4)}P_2 + SHIP_2$$
 (3)

$$PIP_3 + X \stackrel{k_4^{on}}{\longleftrightarrow} PIP_3 - X \tag{4}$$

$$PIP_3 - X \xrightarrow{k_5^{on}} PI_{(3,4)}P_2 + X$$
 (5)

$$PIP_{3} + PTEN \xrightarrow{k_{6}^{on}} PIP_{3} - PTEN$$
(6)

$$PI_{(3,4)}P_2 + INPP_4AB \xrightarrow{k_7} PI_{(3,4)}P_2 INPP_4AB$$
 (7)

#### 1.2 PI3K inhibition

In the presence of 1uMPI-103 PI3K inhibitor  $k_1 = 0$ . PIP3 degradation is performed by PTEN, SHIP2 and some other phosphatase X. If additionally assumed that  $k_6^{on} >> k_6^{off}$ ,  $k_2^{on} >> k_2^{off}$  and  $k_4^{on} >> k_4^{off}$  then changes of PIP3 concentration can be written as:

$$\frac{d[PIP3]}{dt} = -[PIP3](k_6^{on}[PTEN] + k_2^{on}[SHIP2] + k_4^{on}[X])$$

Solution of this equation:

$$\frac{d[PIP3]}{[PIP3]} = -dt(k_6^{on}[PTEN] + k_2^{on}[SHIP2] + k_4^{on}[X])$$

$$ln([PIP3]) = -(k_6^{on}[PTEN] + k_2^{on}[SHIP2] + k_4^{on}[X])t + const$$

$$[PIP3] \sim e^{-(k_6^{on}[PTEN] + k_2^{on}[SHIP2] + k_4^{on}[X])t}$$

Assuming that concentrations of [PTEN], [SHIP2] and [X] are constant:

$$[PIP3] \sim e^{-(PTEN_{act} + SHIP2_{act} + X_{act})t}$$
 (8)

### 1.3 Exponential regression

 $PTEN_{act}$ ,  $SHIP2_{act}$  and  $X_{act}$  in equation (8) can be defined from time courses of PIP3 concentration data (in the presence of 1uMPI-103 PI3K inhibitor). We performed exponential regression of PIP3 data and results are shown in Fig. 2.

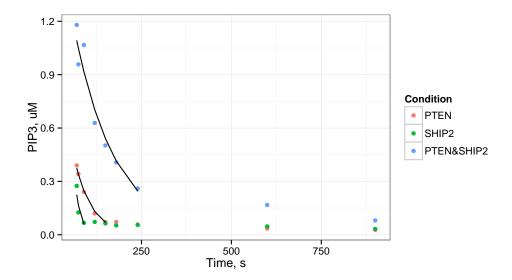


Figure 2: PIP3 time courses in the presence of 1uMPI-103 PI3K inhibitor. Black lines - regression of the first few time points based on equation (8).

Note that time courses of PIP3 concentration have a biphasic behavior and it was not possible to fit all of the data points to a single exponent, thus regression analysis was performed only on a few first time points of the time courses. Regression analysis provides:

$$SHIP2_{act} + X_{act} = 0.0212[\mu M \cdot s]^{-1}$$
 
$$PTEN_{act} + X_{act} + \alpha \cdot SHIP2_{act} = 0.0654[\mu M \cdot s]^{-1}$$
 
$$X_{act} + \alpha \cdot SHIP2_{act} = 0.0088[\mu M \cdot s]^{-1}$$

 $\alpha$  is a percentage of active SHIP2 left in the cells after removing it using siRNA (90% of SHIP2 RNAs are removed). If  $X_{act} = 0$  then  $\alpha \simeq 41.5\%$ . Taking into account that there is a non-linear dependence between active SHIP2 left in the cells and the amount of silenced SHIP2 RNAs (90%) one could suggest that there

is no other phosphatase X in the pathway and there is still 41.5% of active SHIP2 left in the cells.

However, if  $X_{act} \neq 0$  and  $\alpha = 10\%$ , then:

$$SHIP2_{act} = 0.0138[\mu M \cdot s]^{-1}$$

$$X_{act} = 0.0074[\mu M \cdot s]^{-1}$$

$$PTEN_{act} = 0.0566[\mu M \cdot s]^{-1}$$

So that  $X_{act} \simeq 53.6\% SHIP2_{act}$  and  $SHIP2_{act} \simeq 24.4\% PTEN_{act}$ 

### 1.4 Inconsistency in PTEN KO and SHIP2 silencing

Experimental data (in the absence of inhibitors) show that removing either PTEN or SHIP2 from the cells increase a peak of PIP3 concentration to  $\sim 5[uM]$ . However, in previous section it is shown that activity of PTEN on PIP3 is  $\sim 5$  times higher than that of SIP2 on PIP3. Thus, theoretically it is expected that PTEN will increase the peak of PIP3 to  $\sim 1.4 - 1.5[uM]$  (checked with COPASI).

## 1.5 Is all PI(3,4)P2 produced from PIP3 through SHIP2?

Let us assume there is no X phosphatase in the pathway in Fig. 1. Then PI(3,4)P2 is produced and degraded as described by equations (3) and (7). If INPP4AB is also silenced then PI(3,4)P2 would be only produced as described by equation (3):

$$PIP_3\_SHIP_2 \xrightarrow{k_3^{on}} PI_{(3,4)}P_2 + SHIP_2$$

where production of PIP3\_SHIP2 complex is described by equation (2):

$$\mathrm{PIP}_{3} + \mathrm{SHIP}_{2} \xleftarrow[\mathrm{k_{2}^{on}}]{} \mathrm{PIP}_{3} \text{\_SHIP}_{2}$$

In these two reaction there is only information about  $k_2^{on}$  (see chapters above about  $SHIP2_{act}$ ). However, one can estimate the maximum production of PI(3,4)P2 by assuming that  $k_3^{on} \gg k_2^{on} \gg k_2^{off}$ . In this case two reactions above can be simplified into one:

$$PIP_3 + SHIP_2 \xrightarrow{k_2^{on}} PI_{(3,4)}P_2 + SHIP_2$$
 (9)

Note that equation (9) correspond to the case when the rate of PI(3,4)P2 production is maximal possible. Based on this equation changes of PI(3,4)P2 concentration can be written as:

$$\frac{d[PI(3,4)P2]}{dt} = k_2^{on}[SHIP2][PIP3]$$

Using the definition of  $SHIP2_{act}$  (see equation (8)):

$$d[PI(3,4)P2] = SHIP2_{act}[PIP3]dt$$

And after integration:

$$[PI(3,4)P2](t) = SHIP2_{act}S_{PIP3}(t) + [PI(3,4)P2](0)$$
(10)

where  $S_{PIP3}(t)$  is the area and the curve of PIP3 time course.

We then calculated the maximum concentration of PI(3,4)P2 produced from PIP3 by SHIP2 using experimental data on PIP3 concentration and equation (10) and compared it with experimental data on PI(3,4)P2 concentration. Results are shown in Fig. 3.

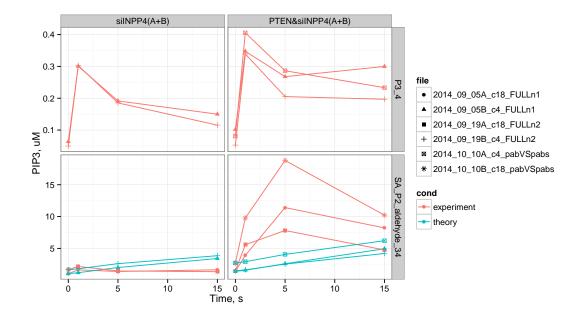


Figure 3: PI(3,4)P2 experimental and theoretical (based on equation (10)) time courses.

It is clear that just SHIP2 is not enough to be able to produce such a high peak of PI(3,4)P2 concentration as shown in Fig. 3 (PTEN&siINPP4(A+B) condition).

### 1.6 Hypothesis: PTEN dephosphorylates PI(3,4)P2 to PI4P

Transient concentrations of PI(3,4)P2 in PTEN&siINPP4(A+B) condition are very large to be controlled by only one phosphotase SHIP2 (see paragraph above). Since in siINPP4(A+B) condition concentration of PI(3,4)P2 does not significantly change, an obvious hypothesis would be that PTEN might be another phosphotase that take a role in dephosphorylation of PI(3,4)P2 to PI4P (see Fig. 1). Thus, when PTEN is knocked out PI(3,4)P2 accumulates faster. However, adding PTEN to PI(3,4)P2 regulation would in theory add an extra outflux (higher than the influx of SHIP2) from PI(3,4)P2 pool. This outflux cannot explain accumulation of PI(3,4)P2 in the absence of PTEN and would completely vanish PI(3,4)P2 in siINPP4(A+B) condition (blue line in Fig. 3). Therefore, there must be another source of PI(3,4)P2 of approximately the same strength as PTEN activity to be able to neutralize PI(3,4)P2 degradation by PTEN in siINPP4(A+B) condition and to accumulate a large amount of it in PTEN&siINPP4(A+B) condition (checked with COPASI).