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**A yeast-based assay shows that long chain n-3
polyunsaturated fatty acids and arachidonic acid inhibit
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A yeast-based assay shows that long chain n-3 polyunsaturated fatty acids and arachidonic acid inhibit PI3K activity *in vivo*

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Abbreviations used: AA, arachidonic acid; DHA, docosahexaenoic acid; EPA, eicosapentaenoic acid; PDK1, 3'-phosphoinositide-dependent kinase 1; PH, pleckstrin homology; PI3K, phosphatidyl inositol 3' kinase; PI, phosphatidyl inositol; PIP, phosphatidyl inositol 3-phosphate; PIP₂, phosphatidyl inositol 4,5-bisphosphate; PIP₃, phosphatidyl inositol 3,4,5-trisphosphate; PKB, protein kinase B; PLCδ1, phospholipase δ1; PUFA, polyunsaturated fatty acids; PTEN, phosphatidylinositol 3-phosphatase; SH2, Src homology

ABSTRACT

The PI3K (phosphatidyl inositol 3' kinase) pathway controls the regulation of cell growth, proliferation, migration and apoptosis. In many tumors, the PI3K gene is mutated or overexpressed, and/or the PI3K pathway is hyperactive. PI3K is therefore a potential pharmacological target for the development of anti-tumor drugs. Some polyunsaturated fatty acids (PUFA), when given in the diet, may lead to a decrease in PI3K activity. We used a yeast-based model to reconstitute the PI3K/PTEN/Akt pathway to study the effects of long chain n-3 fatty acids on PI3K kinase and found that various PUFA were able to alleviate toxicity induced by overexpression of PI3K. The various PUFA had no significant effect on the steady-state level of PI3K catalytic subunit proteins in yeast. However, depletion of PIP₂ due to overexpression of the p110 α subunit was significantly reduced by treating the yeast cells with the various PUFA. The inhibition of mammalian PI3K, expressed in an exogenous cellular context in yeast, is likely to be a direct effect of these PUFA on PI3K rather than on other mammalian endogenous or environmental factors. These results are particularly promising given the abundance of active PUFA in marine foodstuffs and especially fish oils.

1 INTRODUCTION

The phosphatidylinositol 3'-kinase (PI3K) signaling pathway controls the regulation of cell growth, proliferation, migration and apoptosis [1, 2]. Type 1A PI3K is composed of a regulatory subunit, p85, which is responsible for its plasma membrane localization and a catalytic subunit, p110, responsible for its kinase activity. The activation of the catalytic subunit, p110, targets PI3K to its substrate phosphatidyl inositol 4,5-bisphosphate (PIP₂) localized at the plasma membrane. The activity of p110 generates phosphatidyl inositol 3,4,5-trisphosphate (PIP₃) from PIP₂ in the plasma membrane. PIP₃ in turn generates recognition sites for proteins containing a pleckstrin homology (PH) domain; these proteins include 3'-phosphoinositide-dependent kinase 1 (PDK1) and protein kinase B (PKB/Akt) which both translocate to the inner surface of the plasma membrane in response to the generation PIP₃. Once at the surface of the plasma membrane, PKB/Akt is phosphorylated on Thr308 by PDK1 and fully activated by phosphorylation on Ser473 by PDK2

kinases (mainly mTOR complex 2 and DNA-PKcs). Once activated, PKB/Akt phosphorylates a myriad of proteins which control cell growth, cell proliferation and apoptosis, all steps involved in tumor formation and malignant cell dissemination [2]. The PI3K pathway is negatively regulated by two phosphatases: SHIP (SH2 domain-containing inositol phosphatase) and PTEN (phosphatase and tensin homolog deleted in chromosome ten). In many tumors, the PI3K gene is mutated or overexpressed, and PTEN function is attenuated or the protein underexpressed; consequently, the PI3K/Akt pathway is overactive [3]. Thus, PI3K is involved in the development of many types of cancers [4, 5]. It also seems to be involved in the radioresistance mechanism of head and neck cancers [6]. Indeed, generation of PIP₃ at the plasma membrane is pivotal for the regulation of mammalian cell proliferation and survival by growth-promoting factors [1]. Therefore, PI3K is an attractive pharmacological target for screening for anti-cancer drugs [3].

Most known inhibitors of PI3K are synthetic molecules [3], but there is evidence that long chain n-3 fatty acids (i.e. EPA and DHA) found in marine foodstuffs, especially fish oils, may inhibit PI3K activity *in vivo*. Indeed, long chain n-3 fatty acids act as protein kinase inhibitors in the central nervous system [7]. In rats consuming a small amount of fish oils (2.2% of calories) in a normolipidic diet, PI3K activity in both liver and muscle is 54% lower than that in rats on a control diet without fish oil, as judged by determining *in vitro* the activity of PI3K after immuno-precipitation from cell extracts [8]. Also, EPA restores tamoxifen sensitivity to breast cancer cells with high Akt activity [9]. Here, we confirm that long chain n-3 fatty acids are inhibitors of PI3K activity. We used, as a model system, the reconstitution of the mammalian PI3K/PTEN/Akt pathway in yeast, as described by Rodriguez-Escudero et al [10]. In this model, the catalytic subunit of mammalian PI3K, p110 α , is overproduced and artificially targeted to the plasma membrane, where it converts PIP₂ to PIP₃: this inhibits cell growth by depleting the cellular pool of PIP₂ which is essential for yeast. This effect is completely prevented by the co-expression of catalytically active mammalian PTEN which dephosphorylates PIP₃. The inhibitory effect on yeast growth is partially relieved by LY294002, a PI3K inhibitor, and similar overproduction of a catalytically inactive mutant of p110 α (p110 α^{K802R}) has no effect on growth and viability or on PIP₂ depletion: these two observations confirm that the inhibitory effect depends on PI3K kinase activity. The production of PIP₃ in yeast by induction of the p110 α gene induces the translocation of PKB/Akt to the plasma membrane and enhances its

phosphorylation, as observed in mammals *in vivo*. Therefore, this model can be used to determine whether EPA and DHA in the culture medium of yeast inhibit the production or catalytic activity or both of the p110 α subunit and this, by freeing itself from any other mammalian endogenous or environmental factors which may interfere *in vivo* with the effects of long chain n-3 Poly Unsaturated Fatty Acids (PUFA) on PI3K activity. In addition, and contrary to the rat model, in the yeast-based model PI3K activity is directly correlated to *in vivo* measurable phenotypes.

2 MATERIALS AND METHODS

Yeast strains, culture media and plasmids

The *Saccharomyces cerevisiae* strains used were the Wild Type (WT) K699: *Mata*, *ade2-1*, *trp1-1*, *can1-100*, *leu2-3,112*, *his3-11,15*, *ura3*, *GAL+*, *psi+*, *ssd1-d2* (W303 genetic background) and its *hof1 Δ* derivative: *Mata*, *ade2-1*, *trp1-1*, *can1-100*, *leu2-3,112*, *his3-11,15*, *ura3*, *GAL+*, *psi+*, *ssd1-d2*, *hof1::KANMX* [11]. YPD (1% (w/v) yeast extract, 2% (w/v) peptone and 2% (w/v) glucose) was the general non selective medium used for yeast cell growth. Synthetic minimal medium contained 0.17% yeast nitrogen base without amino acids, 0.5% ammonium sulfate, 2% glucose, 2% raffinose or 2% raffinose/ 2% galactose and lacked appropriate amino acids and nucleic acid bases to maintain selection for plasmids. Solid media contained 2% (w/v) agar. Yeast transformations were performed by the lithium acetate procedure [12]. The plasmids YCpLG-PI3K and YCpLG-PI3K^{K802R}, allowing expression of p110 α and p110 α ^{K802R}, the catalytic subunit of a membrane-targeted mammalian PI3K, under the control of the galactose-inducible *GAL1* promoter were described in [10]. The plasmids pBM328, allowing expression of *hof1 Δ PEST* (under the control of the inducible *GAL1,10* promoter), and the empty vector, pRD53, were described previously [11]. To visualize PIP₂ *in vivo*, the plasmid pRS426-GFP-2xPH (PLC δ) which allows expression of the GFP protein fused to two tandem copies of the PLC δ 1 PH domain was used [10, 13].

Yeast-based assay

This assay was adapted from an existing test [14, 15]. Briefly, 240 μ l of a culture of exponentially growing cells, adjusted to an OD₆₀₀ of 0.25, were spread homogeneously with sterile glass beads ($\approx$ 3mm diameter)

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Inhibition of PI3K by fatty acids

on a square Petri dish (12 cm x 12cm) containing synthetic minimal medium and 2% glucose or 2% raffinose and 2% galactose. Sterile filters (similar to those used for antibiograms) were placed on the agar surface. A volume of 1µl of a pure solution of the fatty acids (DHA, EPA, AA or oleic acid, as indicated) was spotted onto the filters as indicated in the left panel of **Figures 2 and 3**. Aliquots of 1µl of bromopalmitate or LYS294002 (PI3K inhibitor) in solution in DMSO (at 200µM and 100µM, respectively) were similarly tested on the same Petri plates. As a negative control, a filter was loaded with 1µl of DMSO, the vehicle. Plates were then incubated at 30°C for three to five days and scanned using a Snap Scan1212 (Agfa).

Growth curves and protein detection

Exponentially growing cells were cultured in a liquid minimal selective medium containing 2% raffinose overnight at 30°C, diluted and allowed a few hours of recovery in the same medium. Then, 2% galactose with, where indicated, 0.005% of DHA, EPA, AA (from a 5% solution in DMSO) or DMSO was added to the yeast cultures. For growth curves, OD_{600nm} was determined at the times indicated. For protein detection, 3 to 5 ml of yeast cultures were incubated for six hours (OD₆₀₀ around 0.5) then rapidly harvested by centrifugation. The cell pellets were cooled on ice, resuspended in 150µl of cold lysis buffer (1.85M NaOH, 13% β-mercaptoethanol) and 150µl of a cold TCA solution at 50% was added. The samples were centrifuged at 15,000 g for 2 minutes at 4°C: the precipitated proteins were washed with 1ml of cold acetone and centrifuged again. The acetone was discarded, Laemmli buffer added and the samples were heated at 95°C for 3 min. These protein extracts were then analyzed by SDS-PAGE and transferred to a nitrocellulose membrane (Schleicher & Schuell). Membranes were incubated with primary antibody (at 1:1,000 for anti-PI3K p110α, Cell Signaling Technology or 1:10,000 for anti-actin, Calbiochem JLA20) in TBST buffer (50mM Tris, pH 7.4, 150mM NaCl, 0.1% Tween 20) containing 5% non-fat dry milk for 1 hour. The membrane were then washed with TBST, incubated with secondary antibody (1:3,000 goat anti-rabbit antibody conjugated to horseradish peroxidase, Biorad, for anti-PI3K p110α and 1:2,000 goat anti-mouse IgM, Calbiochem, for anti-actin). Enhanced Chemiluminescence (ECL, Amersham) and a Vilbert-Lourmat Photodocumentation Chemistart 5000 imager were used to reveal bound antibody.

Microscopy

Cells co-transformed with the plasmids YCpLG-PI3K or YCpLG-PI3K^{K802R} and the pRS426-GFP-2xPH(PLC δ 1) vector (which allows constitutive expression of GFP-PH) were grown overnight in a liquid minimal selective medium containing raffinose. Cultures were diluted and allowed to recover for a few hours in the same medium, and 2% galactose (to induce PI3K overexpression) or 2% glucose (to repress PI3K) was added with 0.005% of DHA, EPA, AA or DMSO (from a 5% solution in DMSO). The GFP signal was visualized after 8 hours using a Chroma GFP4 filter (excitation 455-495 nm) on an Olympus BX61 microscope, photographed with a Spot RT cooled CCD camera and analyzed with Photoshop 7.0 software (Adobe). To obtain reliable and statistically significant data, ≥ 250 cells were examined for each condition. Values reported are means of three independent experiments.

3 RESULTS

Ability of fatty acids EPA, DHA and AA to alleviate PI3K-driven toxicity in yeast

The mammalian PI3K/PTEN/Akt pathway has been reconstituted in the yeast *S. cerevisiae* in which PI3K expression and activity inhibits cell growth [10]. Expression of the p110 α catalytic subunit of PI3K under the control of the strong galactose-inducible *GAL1* promoter led to the conversion of the essential PIP₂ pool into PIP₃. This PI3K overexpression-dependent depletion of PIP₂ impaired yeast growth by altering morphogenesis and vesicular trafficking (Figure 1, upper panel and [10]). An analogous construct was obtained with an allele of PI3K carrying a point mutation (Lys⁸⁰² $\rightarrow$ Arg) which encodes a catalytically inactive subunit (p110 α ^{K802R}) (Figure 1, lower panel): there was no growth inhibition confirming that the effect was due to the kinase function of PI3K [10]. In both the p110 α and p110 α ^{K802R} constructs, the coding regions were fused in frame at their 3' end with the coding sequence of the CAAX box of H-ras to target the subunits to the plasma membrane. This was to compensate for the absence of p85, the regulatory subunit of PI3K, which is responsible for the plasma membrane localization of the enzyme in mammals. Therefore the expressed p110 α can be considered as a constitutively active form of PI3K.

Using the same expression vectors as those used by Rodriguez-Escudero et al. who developed this model, p110 α and p110 α^{K802R} were overproduced in *S. cerevisiae* under the control of the strong galactose-inducible *GAL1* promoter. We then tested the ability of various PUFA to suppress PI3K overexpression-dependent growth inhibition. Wild-type (WT) *S. cerevisiae* cells (K699) transformed with the vectors were spread on minimal synthetic medium under either repressing (glucose medium) or inducing conditions (galactose medium). Sterile filters were then placed on the agar surface and various fatty acids were deposited onto the filters as indicated in Figure 2, left panel. This simple method allows creation of a full gradient of concentration of a compound around the filter where it was deposited [14, 15]. We tested two n-3 PUFA found in fish oils, docosahexaenoic acid (DHA, C22:6 n-3) and eicosapentaenoic acid (EPA, C20:5 n-3), and arachidonic acid (AA, C20:4 n-6), an n-6 PUFA which presents the same carbon chain length as EPA. We also evaluated the effects of oleate (or oleic acid), a monounsaturated n-9 fatty acid; bromopalmitate, a non-metabolizable fatty acid (an inhibitor of a variety of enzymes and proteins associated with lipid metabolism) and DMSO, the vehicle used to dissolve some of the molecules tested. The PI3K inhibitor, LY294002, which restores cell growth to yeast overproducing PI3K [10], was used as a positive control. After three days at 30°C and as previously observed [10], WT yeast cells expressing p110 α failed to grow on galactose medium whereas WT yeast expressing p110 α^{K802R} , the catalytically inactive PI3K, grew normally (Figure 2). Both grew normally on glucose medium on which the expression of both forms of p110 α is repressed. DHA and EPA (the two n-3 PUFA), and AA (the n-6 PUFA), all relieved the growth inhibition, as judged by the appearance of a growth halo around the filters on which they were deposited. Oleate, bromopalmitate and DMSO had no such effect on yeast growth. Thus, like the PI3K kinase inhibitor LY294002, both n-3 and n-6 PUFA were able to alleviate PI3K-induced growth inhibition in yeast, suggesting that they may inhibit the kinase activity of PI3K *in vivo*. Interestingly, no clear synergistic effect was observed between the various PUFA on one hand and LY294002 on the other hand (data not shown and Figure 2), suggesting that they act via a common mechanism.

The ability of AA, EPA and DHA to alleviate PI3K-driven toxicity in yeast is not due to an effect on PI3K expression

There are at least two mechanisms by which AA, EPA and DHA may alleviate yeast growth inhibition due to overexpression of PI3K: a direct or indirect effect on PI3K kinase activity; or interference with PI3K gene expression (for example by affecting transcription from the galactose-inducible *GALI* promoter). If AA, EPA and DHA affect expression of the gene, the abundance of PI3K in yeast would be affected; if the activity of the enzyme is affected, its abundance should be unaffected.

We investigated the effects of AA, EPA and DHA on *GALI* promoter-driven expression by testing the effect of these polyunsaturated fatty acids on the toxicity of hof1 Δ PEST overexpressed from this promoter in yeast. hof1 Δ PEST is a stabilized truncated form of Hof1, a protein involved in cytokinesis in *S. cerevisiae* which has to be directed by the E3-ligase SCF^{Grr1} to ubiquitin-26S proteasome dependent degradation at the end of mitosis to allow cell separation [11]. Therefore *GALI* promoter-driven overexpression of Hof1, and especially of its stabilized form hof1 Δ PEST, inhibits yeast cell growth by preventing cell separation at the end of mitosis [11]. We used an experiment similar to that described in Figure 2 except that yeast cells (K699 *hof1 Δ*) were transformed with either a plasmid allowing strong overexpression of hof1 Δ PEST from the same *GALI* promoter or with an empty vector (negative control). None of AA, EPA or DHA (or DMSO, bromopalmitate, oleate or LY294002) was able to alleviate the growth inhibition due to hof1 Δ PEST overproduction. As the only common point between hof1 Δ PEST- and p110 α -driven toxicity is the strong overexpression from the galactose-inducible *GALI* promoter, these results suggest that none of these PUFA significantly interfered with this overexpression (Figure 3a). Next, we tested the effects of AA, EPA and DHA on the steady-state level of p110 α . Yeast cells carrying p110 α under the *GALI* promoter were grown in a galactose medium in the presence of various PUFA (as indicated in Figure 3b). Protein extracts were prepared and analyzed by Western blotting. The amount of p110 α did not differ between cells grown in the presence of AA, EPA and DHA and controls. Findings were similar for yeast cells overexpressing p110 α ^{K802R}, the catalytically inactive PI3K (data not shown). Using the same cells and under the same conditions, we found that long chain n-3 PUFA and AA also improve the growth in liquid culture of yeast cells overexpressing p110 α (Figure 3c for DHA and data not shown). Therefore, AA, EPA and DHA all appear to act by inhibiting the kinase activity of PI3K.

AA, EPA and DHA lead to a partial suppression of PIP₂ depletion

To confirm that AA, EPA and DHA inhibit the activity of mammalian PI3K kinase in yeast, we studied the effects of these fatty acids on PIP₃ production driven by the overexpression p110 α . We used PIP₂ depletion as a marker of the kinase activity of mammalian PI3K. Mammalian PI3K overproduced in yeast cells, by catalyzing the generation of PIP₃ at the expense of PIP₂ (which is essential for yeast growth), leads to PIP₂ depletion thereby inhibiting yeast growth [10]. Importantly, Vps34, a class III PI3K and the sole PI3K homolog in *S. cerevisiae*, catalyzes the conversion of phosphatidyl inositol (PI) into phosphatidyl inositol 3-phosphate (PIP) but is unable to use PIP₂ as a substrate [16, 17]. Therefore the PIP₂ depletion observed upon expression of mammalian PI3K in yeast directly reflects its kinase activity and cannot be imputed to an endogenous yeast PI3K activity. To detect and follow the fate of PIP₂ in yeast, cells were co-transformed with a plasmid expressing a fluorescent PIP₂-specific reporter consisting of GFP fused to two tandem copies of the PH domain of mammalian phospholipase δ 1. In WT yeast cells, this reporter fusion protein specifically decorates PIP₂ in the plasma membrane [13]. As previously observed [10], we found that the reporter clearly and prominently stained the plasma membrane of all cells overexpressing p110 α^{K802R} , in the presence or absence of the various PUFA (Figure 4, right rows), similar to the staining of cells in which the expression of p110 α^{K802R} was not induced. This confirms the inability of p110 α^{K802R} to catalyze conversion of PIP₂ to PIP₃. In contrast, in cells in which p110 α was overproduced, only diffuse cytoplasmic fluorescence was observed, and this in about 95% of the cells (Figure 4, left rows, second line), indicating substantial PIP₂ depletion from the plasma membrane. This effect was specific: the reporter clearly and prominently stained the plasma membrane of all control cells in which the expression of p110 α was not induced (Figure 4, left rows, first line). Cells overproducing p110 α were treated with AA, EPA or DHA and tested for staining. Interestingly, we observed normal plasma membrane localization of the GFP signal in a significant proportion of cells (23%, 30% and 28% respectively). This confirms that these various PUFA inhibited the kinase activity of mammalian PI3K in yeast, thereby counteracting the depletion of PIP₂ (Figure 4, left rows, lines 3 to 5). As controls, we checked that the three PUFA had no effect on the plasma membrane localization of the marker in cells overexpressing p110 α^{K802R} (Figure 4, right rows, lines 3 to 5).

4 DISCUSSION

Our results strongly suggest that in a yeast model reconstituting the PI3K/PTEN/Akt pathway, three long chain PUFA, namely AA, EPA and DHA, inhibited PI3K activity. This is in accordance with previous observations *in vivo* in liver and muscle of rats receiving a diet enriched with a small amount of fish oils (rich in EPA and DHA) [8]. In this study, the PI3K activity was evaluated *in vitro* after immunoprecipitation of tyrosine phosphorylated proteins, thereby justifying the use, in the present study, of the yeast-based system where PI3K activity is directly correlated to *in vivo* measurable phenotypes: growth inhibition and depletion of PIP₂ from the plasma membrane. The novel finding of our study is, therefore, that the inhibition of PI3K activity by AA, EPA and DHA may be direct, like the effects of LY294002 or wortmannin. There are several arguments for a direct mechanism of inhibition. The inhibitory effect previously observed *in vivo* in rats [8] was reproduced in our exogenous (for example the plasma membrane composition differs between yeast and mammalian cells and, yeast cells have a cell wall) and simplified model, and this is a strong argument in favor of a direct inhibition. Indeed, the PI3K/PTEN/Akt pathway was reconstituted in yeast, so the inhibitory effect of AA, EPA and DHA on PI3K activity was independent of any association with most of the other molecules belonging to this pathway, such as insulin receptor substrates 1 or 2 (IRS-1 or IRS-2), two upstream factors that regulate type 1A PI3K activity *in vivo* in mammals [18]. It was also not due to an indirect effect on the level of expression of the PI3K gene. Indeed, none of the three PUFA altered the steady-state abundance of PI3K or alleviated growth inhibition due to hof1ΔPEST overexpression from the same galactose-inducible *GAL1* promoter. It is also unlikely that the inhibitory effect of the three long chain PUFA was due to one or more of their metabolites because the metabolism of long chain n-3 PUFA and of AA generate different metabolites. The inhibitory effect also cannot be explained by inhibition of palmitoylation of PI3K because 2-bromopalmitate which inhibits palmitoylation [19] did not inhibit PI3K.

AA, EPA and DHA (fatty acids with chain length ≥ 20 carbons and with 4, 5 and 6 double bonds respectively) had similar inhibitory effects on PI3K activity whereas oleic acid (chain length: 18 carbons and one double bond) had no such inhibitory effect. This implicates chain length, i.e. ≥ 20 C, and/or the high degree of unsaturation as possible determinants of the inhibitory effect of the three PUFA on PI3K activity.

However, we did not address the exact mechanisms underlying the inhibitory effect of long chain PUFA on PI3K. LY294002 and wortmannin, the most widely used PI3K inhibitors, both competitively target the ATP/substrate-binding site of the p110 catalytic subunit [3]. Further studies are needed to determine whether the inhibitory effect of AA, EPA and DHA involves this type of mechanism. Note that EPA and DHA are potent non competitive inhibitors of several protein kinases, including pKA, pKC, MAPK and CaMKII [7]. PI3K/PTEN signaling is central to mechanisms of tumorigenesis, angiogenesis, and anti-cancer drug and radio resistance [2, 6, 20-29]. Our observation that long chain n-3 and n-6 PUFA are natural PI3K inhibitors is therefore of particular interest. Indeed, there is diverse evidence suggesting that, *in vitro* or during nutritional interventions, long chain n-3 fatty acids enhance the response of tumors to the effects of anti-neoplastic agents (reviewed in [30]).

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Inhibition of PI3K by fatty acids

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FIGURE LEGENDS

Figure 1: flowchart of the mammalian p110 α overexpression-driven toxicity in yeast

A WT yeast strain was transformed with a vector allowing expression from the strong galactose-inducible *GALI* promoter, of either p110 α , the catalytic subunit of PI3K (upper panel) or p110 α^{K802R} , a catalytically inactive form of PI3K (lower panel). Both proteins are fused in frame at their C-termini with the -CAAX box of H-ras which targets them to the plasma membrane to compensate for the absence of p85, the PI3K regulatory subunit which targets p110 α to the plasma membrane in its normal context in mammals. Galactose (right diagram) but not glucose (repressing conditions, left diagram) activates transcription from the strong *GALI* promoter therefore allowing a high level of expression of p110 α which in turn converts the essential PIP₂ into PIP₃ (upper panel). As a consequence, yeast cell growth is strongly impaired due to altered morphogenesis and vesicular trafficking. p110 α^{K802R} , even when overexpressed at a similar level, has an effect neither on yeast cells growth nor on PIP₂ level (lower panel) showing that this growth inhibition effect involves the kinase activity of PI3K.

Figure 2: toxicity linked to overexpression of the mammalian p110 α catalytic subunit of PI3K in yeast is alleviated by several fatty acids

WT yeast strain (K699) expressing either p110 α , the catalytic subunit of PI3K (left Petri plates) or p110 α^{K802R} , a catalytically inactive form of PI3K (right Petri plates), was spread on selective media containing either glucose (lower plates) which inhibits expression from the galactose-inducible *GALI* promoter or galactose (upper plates) which allows a strong expression from the promoter and thus of p110 α (left row) or p110 α^{K802R} (right row). Sterile filters (similar to those used for antibiograms) were then placed on the agar surface and the various compounds to be tested were spotted onto the filters as indicated in the scheme on the left. Pictures of the Petri dishes after 4 days of incubation at 30°C are shown. Only p110 α expression led to a profound growth defect, as previously shown. AA, EPA and

DHA alleviated this growth defect to an even greater extent than LY294002, the positive control (a known chemical inhibitor of PI3K).

Figure 3: the ability of fatty acids -AA, EPA and DHA- to alleviate PI3K-driven toxicity in yeast is not due to an effect on *GALI* promoter-driven PI3K expression

- a. An experiment similar to that described in **Figure 2** was performed except that the yeast cells used were K699 *hof1Δ* transformed either with a vector allowing strong expression of *hof1ΔPEST* (left Petri plate) from the same strong galactose-inducible *GALI* promoter as used for PI3K overexpression or, with an empty vector as a control (right Petri dish). As previously reported [11] when overexpressed, *hof1ΔPEST* led to a strong growth defect (compare left plate with right plate) because it blocks cytokinesis. Note that none of the tested compounds (indicated in the schema on the left) alleviated this growth defect. Therefore, their effect of alleviating toxicity induced by overexpression of p110α is not due to their interference with the *GALI* promoter-driven overexpression.
- b. WT yeast strain (K699) transformed either with an empty vector (lane 1) or with plasmid YCpLG-PI3K allowing overexpression of p110α, the catalytic subunit of PI3K (lanes 2-6) was grown in the presence or absence of galactose and of 0.005% of the fatty acids as indicated. Cell extracts were prepared and analyzed by Western blot using an antibody raised against p110α (upper gel) or actin (lower gel) as a loading control. None of the tested fatty acids tested had a significant effect on the steady state abundance of p110α indicating that their ability to alleviate toxicity due to overexpression of p110α is not caused by an effect on the amount of this catalytic subunit of PI3K.
- c. Left curves: growth curves of WT yeast strain (K699) overexpressing either p110α (PI3K, gray filled circles) or p110α^{K802R} (PI3K-K802R, black circles). Overexpression of the two forms of PI3K was induced at time 0 by addition of galactose. Overexpression of p110α strongly inhibited yeast growth and this effect was first evident 6H after induction of overexpression. Right curves: WT yeast strain (K699) overexpressing p110α was treated either with DMSO (gray-filled triangles) or with 0.005%

DHA (black diamonds). DHA partially suppressed the ability of overexpressed p110 α to inhibit yeast growth.

Figure 4: AA, EPA and DHA lead to a partial suppression of PIP₂ depletion

WT yeast strain (K699) was transformed with a vector expressing GFP-2xPH, a fluorescent PIP₂-specific reporter together with either YCpLG-PI3K allowing overexpression of p110 α (PI3K, left rows) or YCpLG-PI3K^{K802R} allowing overexpression of p110 α ^{K802R} (PI3K^{K802R}, right rows). These strains were grown in a liquid selective medium in the absence (first line) or in the presence of galactose (lines 2 to 5) as indicated on the left (P_{GAL} OFF or ON) to repress or induce, respectively, expression of p110 α or p110 α ^{K802R}. Under repressing conditions, GFP staining and therefore PIP₂ was localized at the plasma membrane (first line). As previously observed [10], only overexpression of PI3K/ p110 α led to the depletion of PIP₂ from the plasma membrane in almost all cells (only 5% of the cells maintained a GFP signal at the membrane, left rows, second line), whereas 100% of cells overexpressing PI3K^{K802R} maintained GFP signal at the plasma membrane (right rows, second line). The effect of AA (third line), EPA (fourth line) and DHA (fifth line) on PIP₂ depletion was evaluated. The presence of each of these fatty acids led to a strong increase in the percentage of cells exhibiting a GFP signal at the plasma membrane (left rows, lines 3 to 5), indicating a reduced PIP₂ depletion (and thus reduced PI3K activity), whereas they had no effect on the GFP signal at the membrane in cells overexpressing PI3K^{K802R} (right rows, lines 3 to 5).

Figure 1

Figure 2

Figure 3

a

b

c

Figure 4

Numbers indicate the percentage of cells with GFP signal at the membrane