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2 **PP2A** inhibitor **PME-1** suppresses anoikis, and is associated with

3 therapy relapse of PTEN-deficient prostate cancers

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32 Abstract

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Identification of novel mechanisms of apoptosis resistance of prostate cancer (PCa) 34 35 cells has translational importance. Here, we discover that inhibition of tumor suppressor phosphatase PP2A by PME-1 inhibits anoikis (apoptosis in anchorage-36 37 independent conditions) in PTEN-deficient PCa cells. PME-1 physically associated with the nuclear lamina and regulated its deformability in PCa cells. In addition, PME-38 39 1 deficient cells, with highly deformable nuclear lamina, were particularly vulnerable to 40 anoikis following cell detachment. As a molecular explanation for increased nuclear 41 lamina deformability, PME-1 depletion induced dephosphorylation of nuclear lamina 42 constituents, Lamin-A/C, Lamin-B1, Lamin-B2, LAP2A, LAP2B, and NUP98, PME-1 inhibition increased apoptosis also in an in ovo tumor model, and attenuated cell 43 survival in zebrafish circulation. Clinically, PCa patients with inhibition of both PP2A 44 45 and PTEN tumor suppressor phosphatases (PME-1^{high}/PTEN^{loss}), have less than 50% 5-year secondary-therapy free patient survival, which is significantly shorter than 46 47 survival of patients with only PTEN-deficient tumors.

In summary, we discover that PME-1 overexpression supports anoikis resistance in PTEN-deficient PCa cells. Further, increased nuclear lamina deformability was identified as plausible target mechanism sensitizing PME-1depleted cells to anoikis. Clinically, the results identify PME-1 as a novel candidate biomarker for particularly aggressive PTEN-deficient PCa.

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54 Keywords: LMNA, FDPS, AR, ERG, mechanotransduction, ECM

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58 Clinical relevance

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60 While organ-confined PCa is mostly manageable, the local and distant metastatic 61 progression of PCa remains a clinical challenge. Resistance to anoikis is critical for PCa progression towards aggressive CRPC. Our data show that PME-1 expression in 62 63 human PCa cells protects the cells from apoptosis induction in anchorageindependent conditions both in vitro and in vivo. Clinically, our results identify PME-1 64 65 as a novel putative biomarker for extremely poor prognosis in PTEN-deficient PCa. Taken together, our results demonstrate novel post-translational regulation of key 66 67 cancer progression mechanisms, with clear translational implications.

69

70 Introduction

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72 Prostate cancer (PCa) is often detected early, and can remain non-aggressive, and 73 non-metastatic, for years. One of the hallmarks of PCa progression towards an 74 aggressive castration-resistant prostate cancer (CRPC) is that PCa cells acquire 75 resistance towards specific type of programmed cell death, anoikis (1). Anoikis is induced in adherent cells by their detachment from other cells, or from surrounding 76 77 extra-cellular matrix (ECM). Anoikis suppression is not only relevant for indolent PCa 78 cells to acquire anchorage-independence, but also for survival of prostate cancer cells 79 with metastatic potential in circulation (2, 3). Thereby, characterization of mechanisms 80 supporting anoikis resistance of PCa cells could provide novel therapy opportunities 81 for clinical management of PCa by preventing the progression to aggressive and 82 metastatic disease.

83

84 Mechanistically anoikis resistance in PCa has been linked to changes in cell adhesion, cytoskeleton, as well as deregulated intracellular survival pathways (1). Tumor 85 86 suppressor phosphatase PTEN is inactivated in a large fraction of high grade PCas 87 (4), and prostate-specific PTEN deletion in a mouse model leads to metastatic PCa 88 (5). Importantly, both PTEN deletion, and hyperactivity of PTEN downstream target 89 AKT increases anoikis resistance (6-8). On the other hand, recent studies in other 90 cancer types indicate for a very delicate balance between nuclear lamina stiffness and 91 anchorage-independence, anoikis resistance, and cell migration. Whereas cell 92 migration in three dimensional contexts requires a sufficiently deformable nuclear 93 lamina to allow passage through physical restrictions (9-11), too deformable nuclei

94 may limit cancer cell survival especially in anchorage-independent conditions (9, 12-15). Of nuclear lamina proteins, particularly inhibition of both Lamin-A/C and Lamin-95 96 B1 sensitizes cells to DNA damage and apoptosis (16, 17). Further, Lamin A/C 97 deficiency reduces circulating tumor cell resistance to fluid shear stress (12), and to 98 sensitize cells to anoikis (14). Together, this evidence highlights the emerging 99 importance of nuclear lamina deformability in defining anoikis resistance. However, 100 the post-translational regulation of nuclear lamina deformability, or its importance for 101 PCa progression is poorly understood.

102

103 Protein phosphatase 2A (PP2A) is another tumor suppressor phosphatase, in addition 104 to PTEN, that is commonly inactivated in PCa. PP2A functions as a trimeric protein 105 complex composed of the scaffolding A-subunit, catalytic C-subunit, and a number of 106 regulatory B-subunits (18, 19). In most human cancers, PP2A is inactivated by 107 overexpression of PP2A inhibitor proteins such as CIP2A, SET or PME-1 (20-22), 108 whereas in a small percentage of cancers it is inhibited by either loss of specific B-109 subunits or by inactivating mutations in A-subunits (18). Overexpression of PP2A 110 inhibitor protein CIP2A clinically associates with CRPC (26), and inhibition of both 111 CIP2A and SET inhibits malignant growth of PCa cells, including PTEN-deficient 112 CRPC cells (26, 27). In contrast, role for PME-1 in PCa is currently unknown. 113 Interestingly, PP2A, CIP2A and PME-1 all physically associate with Lamin-A/C (28-114 30), and PP2A B-subunit PPP2R2A promotes nuclear lamina reformation after mitosis 115 (31). However, the specific phosphorylation sites regulated by PP2A on nuclear lamina constituents, and the oncogenic relevance of PP2A's potential role on nuclear lamina 116 117 regulation, remain to be identified.

118

119 In this study, we demonstrate that PP2A inhibitor protein PME-1 (21), that has not 120 been previously implicated in PCa, has a critical role in anoikis suppression of PTEN-121 deficient PCa cells both in vitro and in vivo. PME-1 regulates phosphorylation of 122 several nuclear lamina proteins. Consequently, PME-1-depleted PCa cells have 123 significantly more deformable nuclei, and those cells are hypersensitized to cell death 124 under anoikis-inducing conditions. Clinically, PCa tumors with inhibition of both tumor and 125 suppressor phosphatases PTEN (genetic deletion), PP2A (PME-1 126 overexpression), had significantly shorter relapse-free survival. Finally, PME-1 127 overexpression suppressed anoikis and promoted Lamin-A/C phosphorylation in 128 genetically defined PTEN-deficient cell model.

130 Result

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132 **PME-1** promotes anchorage-independent growth of prostate cancer cells.

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Based on recent reports of potential clinical relevance of PP2A inhibition in human 134 PCa (23-27, 32, 33), but lack of studies related to PME-1, we tested the impact of 135 136 PME-1 depletion on colony growth of two PTEN-deficient PCa cells lines PC-3 (PTEN null) and DU-145 (PTEN heterozygous). PME-1 silencing by two independent siRNA 137 138 sequences did not affect colony growth of either of the cell lines in 2D adherent culture 139 conditions (Figure 1A). However, PME-1 silencing did decrease anchorageindependent growth on soft agar assay in both PC-3 and DU-145 cells (Figure 1B). To 140 141 further assess whether PME-1 is particularly relevant for growth under low-attachment 142 conditions, we performed spheroid growth assays on low attachment plates. PME-1 inhibition prevented the growth of floating PC-3 spheroids (Figure S1A), and reduced 143 144 cell viability in these conditions (Figure 1C). Furthermore, we observed a strong synergy between PME-1 inhibition and low-attachment culture conditions in PARP 145 cleavage (Figure 1D), indicating that apoptotic cell death contributes to decreased 146 147 anchorage-independent growth of these cells. In order to rule out the possibility of 148 siRNA off-target effects, we created a PC-3 PME-1 knock-out (KO) cell line by 149 CRISPR/Cas9. Comparison of cleaved PARP expression between Cas9 150 overexpressing control cells with and without non-targeting gRNA (lanes 1-3), a pool 151 of PME-1 gRNA transfected cells (lane 4), and a single cell subclone of PME-1 152 targeted cells (lane 5), demonstrated dose-dependent apoptosis induction upon PME-153 1 loss in cells following detachment (Figure 1E). PME-1 KO cells also displayed 154 significantly reduced cell viability on low attachment plates (Figure S1B).

155

Previously, it was shown that non-transformed cells succumb to anoikis-type cell death 156 157 when plated on soft substrates, whereas cancer cells do not die under same conditions 158 (34). In order to test whether PME-1 expression is part of the apoptosis resistance 159 mechanism also for cancer cells on low stiffness matrix, either control or PME-1 siRNA transfected PC-3 cells were plated on low- (0.5 kPa) or high (50 kPa)-stiffness 160 161 hydrogels functionalized with ECM components (fibronectin and collagen I). Consistently with published results with other cancer cells (34), control PC-3 cells 162 163 showed a very small increase in PARP cleavage on low stiffness (Figure 1F). Importantly, PME-1 depletion sensitized the cells to apoptosis induction selectively in 164 low-stiffness conditions (Figure 1F). 165

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167 Together these data demonstrate that PME-1 protects human PCa cells from 168 apoptosis when cells are exposed to anchorage independence, or to low tissue 169 stiffness.

170

171 PME-1 supports *in vivo* anoikis resistance and survival of prostate cancer cells
172 in circulation

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To test the *in vivo* relevance of PME-1-mediated inhibition of PCa cell anoikis, we used either scrambled or PME-1 siRNA transfected PC-3 cells in a chicken embryo chorioallantoic membrane (CAM) assay (Figure 2A). Tumors formed by PME-1 depleted cells were overall more translucent, suggesting decreased tumor growth (Figure S1C). In accordance with anoikis suppressing activity of PME-1, histological analyses of dissected tumors revealed increased TUNEL positivity in tumors derived

from PME-1 siRNA transfected cells (Figure 2B,C). Tumor suppression was further
 confirmed by reduced number of Ki-67 positive cells in tumors derived from PME-1
 siRNA transfected cells (Figure 2B,C).

183

To test whether high PME-1 expression would suppress cell death also in response to 184 flow-shear stress, and in disseminated PCa cells, we examined survival of control and 185 186 PME-1 depleted PC-3 cells in zebrafish circulation using recently described 187 experimental setting (35). In short, cell suspensions of scrambled and PME-1 siRNA 188 transfected cells were microinjected into the common cardinal vein of the embryo 189 using a glass microinjector and successfully transplanted embryos were selected to 190 the experiment. After overnight incubation the embryos were imaged by fluorescence 191 stereomicroscopy (Figure 2D). Quantitation of fluorescent tumor cells per zebrafish 192 embryo demonstrated that PME-1 significantly supported survival of circulating PC-3 cells (Figure 2E). 193

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Together, these results validate the *in vivo* relevance of PME-1-mediated anoikis
resistance in PCa cells.

197

198 PME-1 associates with the nuclear lamina and regulates nuclear deformability199

200 PME-1-mediated support of cancer cell survival in other cell types has been mostly 201 attributed to increased AKT activity (36). However, we could not detect any consistent 202 effects on AKT phosphorylation by PME-1 silencing in PC-3 cells either *in vitro* or *in* 203 *vivo* (Figure S2A,B). As a reason for these discrepant results, it is possible that PTEN 204 loss in PC-3 is alone sufficient for activation of AKT, and PME-1 has no further effect.

Also, the levels of MYC, which is another oncogenic PP2A target relevant for PCa, were not affected by PME-1 silencing (Figure S2A,B).

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208 To seek for alternative targets for PME-1-mediated anoikis suppression in prostate 209 cancer cells, we focused on recently identified association of PME-1 with nuclear 210 lamina. PME-1 was shown to interact with both Lamin-A/C, and with nuclear lamina-211 associated farnesylase FDPS (30, 37). Further, siRNA-mediated inhibition of either 212 PME-1, or Lamin-A/C, or FDPS, resulted in comparable loss of viability phenotype in 213 PC-3 cells (30). The rationale of nuclear lamina regulation being a potential 214 mechanism protecting PME-1 expressing cells from anoikis is based on recent studies 215 demonstrating that increased nuclear lamina deformability sensitizes cancer cells to 216 apoptosis and anoikis (9, 12, 13). More specifically, inhibition of both Lamin-A/C and 217 Lamin-B1 sensitizes cells to DNA damage and apoptosis (16, 17).

218

219 We confirmed the physical association of PME-1 with Lamin-A/C in PC-3 cells by 220 proximity ligation assay (PLA) (Figure 3A). The specificity of PLA signal was confirmed 221 by clear loss of signal in PME-1 depleted cells. To identify potential PME-1-regulated PP2A target phosphorylation sites on nuclear lamina, we re-analyzed recently 222 223 reported LC-MS/MS phosphoproteome analysis data from PME-1 depleted cells (28). 224 Notably, PME-1 inhibition was found to result in dephosphorylation of several nuclear 225 lamina constituents. Specifically, PME-1 inhibition resulted in dephosphorylation of 226 Lamin-A/C, Lamin-B1 and B2, NUP98, and of chromatin anchoring proteins LAP2A 227 and LAP2B (Figure 3B), without altering cell cycle distribution (data not shown). As 228 many of the regulated proteins impact nuclear deformability, the effects of PME-1

depletion on nuclear deformability was directly assessed by subjecting the control and
PME-1 knock-out cells to a micropipette aspiration experiment as described previously
(38). Indeed, PME-1 KO cells had significantly more deformable nuclei based on
micropipette aspiration experiment (Figure 3C,D). Interestingly, both cell lines show a
similar slope for the deformation over longer time periods and mostly vary in their initial
deformation (Figure 3D), indicating differencies in their elastic properties.

235

236 Although the increased nuclear deformability in PME-1 negative cells potentially 237 involves dephosphorylation of all identified nuclear lamina components (Figure 3B), 238 Lamin-A was the only protein for which there were phosphoantibodies available for the 239 PME-1-regulated sites. From the original PP2A LC-MS/MS phosphoproteome data 240 (28), we identified 8 Lamin-A/C phosphopeptides, with 10 phosphosites, that were 241 regulated negatively by PME-1 depletion, and positively by inhibition of the PP2A 242 scaffold subunit PPP2R1A (Figure S3A). Out of these sites, currently there are phosphoantibodies available for Serine 22 (S22) and S392 and these were used to 243 244 validate nuclear lamina phosphorylation regulation by PME-1 under anoikis-inducing 245 conditions.

246

Recent studies using hematopoetic stem cells have demonstrated increase in S22 phosphorylation of Lamin-A/C in response to loss of cell adhesion (39). We could confirm this phenotype also in scrambled siRNA transfected PC-3 cells by trypsininduced detachment (Figure 3E). Importantly, the cells in which PME-1 expression was inhibited, either by siRNA, or by knockout, showed clearly lower levels of Lamin-A/C serine 22 phosphorylation up to 210 minutes of follow-up after detachment (Figure 3E and S3B). Importantly, similar results were observed with serine 392

254 inhibition of Lamin-A/C phosphorylation (Figure 3E). Further, serine 22 255 phosphorylation was sustained for up to 24 hours in PME-1 depleted cells that were 256 detached and re-plated on low stiffness hydrogel matrix (Figure 3F and S3C). To 257 further strengthen the PP2A-dependence of Lamin-A/C phosphorylation regulation by PME-1, we carried out a rescue experiment by co-depletion of the PP2A B-subunit 258 259 PPP2R2A. PPP2R2A mediates PME-1 effects in human glioma cell therapy resistance 260 (40), and it is deleted in 8% of human prostate cancers (24, 25) (Figure S3D). 261 Reassuringly, co-depletion of PPP2R2A prevented inhibition of Lamin-A/C 262 phosphorylation in detached PME-1 depleted cells (Figure S3E,F).

263

Given that Lamin-A/C is the major determinant of nuclear lamina elasticity (14), and 264 265 that PME-1 regulated several Lamin-A/C phosphorylation sites, we used Lamin-A/C 266 phosphorylation mutants to study the functional relevance of this phosphorylation 267 regulation for anoikis resistance. To this end we used PC-3 cells in which endogenous Lamin-A/C expression was suppressed by shRNA targeting the 3'UTR of Lamin-A/C 268 269 gene (Figure 3G), and the cells were transiently transfected with either wild-type Lamin-A/C or with a Lamin-A/C mutant with both serine 22 and 392 mutated to alanine 270 271 (Figure S3G). Supportive of the importance of Lamin-A/C S22 and S392 272 phosphorylation for survival in low attachment conditions, re-expression of phospho-273 deficient Lamin-A/C mutants reduced the survival of cells with low endogenous Lamin-274 A/C levels, whereas overexpression of the WT Lamin-A/C resulted in a modest 275 increase in survival (Figure 3H).

276

These results identify nuclear deformability as a novel PP2A-regulated phenotype in cancer cells. Together with data linking nuclear lamina deregulation to increased

apoptosis susceptibility (9, 12, 13, 16, 17), PME-1-mediated regulation of nuclear
lamina is a plausible mechanism by which PME-1 negative prostate cancer cells are
protected from anoikis (Fig. 4H).

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PME-1 overexpression in prostate cancer associates with PTEN loss and with therapy relapse

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To evaluate the clinical relevance of PME-1 in human PCas, PME-1 protein expression, and its clinicopathological associations, were evaluated in PCa tissue microarray (TMA) material consisting of 358 patients treated primarily with radical prostatectomy in the Helsinki University Hospital between 1983 and 1998. The clinical cohort has been previously described in detail (41, 42) (see also Table S1 for demographics).

292

293 The specificity of the PME-1 antibody for immunohistochemical (IHC) stainings has 294 been validated previously (43). PME-1 expression was scored using 4-tier scale of 295 negative, low, intermediate and strong expression (Figure 4A). As each patient had 296 three cancerous cores in the TMA, the maximum values of PME-1 scores were used 297 and each patient was dichotomized as either high or low (strong vs. negative to 298 intermediate staining). In the correlation analysis with clinical variables, strong PME-1 299 protein expression correlated with higher Grade group and advanced stage (Table 2). 300 We also correlated PME-1 status with previously assessed PTEN, ERG and AR 301 status. Notably, high PME-1 expression statistically significantly associated with 302 complete PTEN loss, but also with ERG positivity and high AR expression status 303 (Figure S4A). Further assessment of patient populations based on this data, and

regarding predicted activity status of PTEN and PP2A, defined a subcohort of
 approximately 8% of patients with impaired activities of both tumor suppressors
 (Figure 4B).

307

308 Survival analysis showed that the patients with high PME-1 expression have 309 shortened disease-specific survival, although this was not statistically significant 310 (Figure S4B). However, patients with high PME-1 expression had statistically 311 significantly shorter time to secondary therapies after primary treatment (i.e. relapse 312 free survival) (Figure 4C). Most importantly, the patients with both complete loss of 313 PTEN and high PME-1 expression defined a remarkably aggressive patient population 314 with less than 50% 5-year secondary-therapy free survival (Figure 4D). Similar, albeit 315 less prominent cooperative effects were observed with PME-1 overexpression and 316 with either ERG or AR expression (Figure 4E and S4C).

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318 Together with functional data these observations indicate clinical relevance for PME-319 1-mediated anoikis suppression in PTEN-deficient PCa. To confirm this functional 320 association of two tumor suppressor phosphatases in a more genetically defined 321 model system, PME-1 cDNA construct was stably transfected to PTEN/p53 knock-out 322 mouse embryonic fibroblasts (MEFs). In accordance with siRNA and Crispr/Cas 323 results, overexpression of PME-1 in PTEN/p53 knock-out MEFs prevented PARP 324 cleavage upon 24h incubation in low attachment conditions (Figure 4F). Moreover, 325 PME-1 overexpression increased Lamin-A/C phosphorylation in PTEN/p53 knock-out 326 MEFs consistently with other results (Figure 4G).

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328

329 **Discussion**

330 During tumor growth, and particularly during invasion and metastatic spread, cancer 331 cells are exposed to varying levels of stress. Particularly detachment of cells from 332 other cells, and from ECM, makes cells vulnerable to anoikis induction. Thereby, mechanisms that inhibit cancer cell anoikis facilitate local tumor spreading, and 333 334 eventually metastasis. Here we report high expression of PP2A inhibitor PME-1 as a 335 anoikis resistance mechanism in PTEN-deficient PCa cells. The clinical relevance of 336 anoikis suppression by PME-1 is supported by association of patient population with 337 PTEN^{loss}/PME-1^{high} tumors with remarkably short time to secondary therapies.

338

339 Although anoikis suppression is a generally relevant mechanisms for tumor 340 progression, it may have particular clinical importance in slowly progressing cancers 341 such as PCa where cancers can be diagnosed in indolent phase, and there is a strong 342 need to be able to both predict and inhibit the likelihood of disease progression (4). 343 Previously anoikis resistance in PCa cancer has been linked mechanistically to 344 regulation of cell adhesion, cytoskeleton (1), PTEN inhibition (6), and AR activity (44). 345 In addition to remarkable clinical connection between PTEN and PME-1 (Figure 4D), we found that low PME-1 and AR expression tumor expression was associated with 346 347 significantly prolonged therapy-free survival (Figure 4E). These results are particularly 348 interesting given that pharmacological PP2A reactivation leads to AR degradation, and 349 to significant PCa tumor regression (32), corroborating situation seen in the less 350 aggressive tumors with high PP2A activity (PME-1^{low}) and low AR expression (Figure 351 4E).

352

PTEN and PP2A have both been identified independently as PCa tumor suppressor 353 354 phosphatases (4, 23, 27, 33), but the clinical relevance of their co-operation has not 355 been studied thus far. Here, we demonstrate an important role for PME-1-mediated 356 PP2A inhibition in anoikis suppression in various PTEN-deficient models. Clinically, 357 simultaneous inhibition of both tumor suppressor phosphatases was associated with 358 very aggressive disease with high propensity for early disease relapse (Figure 4D). As 359 PTEN deficiency has also been shown to promote anoikis resistance of PCa cells (6), the tumor cells with inhibition of both PP2A and PTEN are assumed to be particularly 360 361 well protected from anoikis, and we hypothesize that this is likely to contribute to 362 observed clinical aggressiveness of these cancers. Mechanistically PTEN-mediated anoikis resistance is mediated by AKT signaling (6, 7), whereas we showed that PME-363 364 1 depletion had no effect on AKT phosphorylation (Figure S3A,B). The non-365 overlapping downstream mechanisms for PTEN and for PME-1 in anoikis suppression further explains their synergetic actions. As PTEN genetic status can be routinely 366 367 evaluated in current clinical PCa diagnostic practice (4), our results indicate diagnostic utility of assessment of PME-1 status on patients with complete PTEN loss. Although 368 369 further studies are clearly needed to validate these conclusions, our results indicate that patients with PTEN^{loss}/PME-1^{high} tumors might benefit from more intensive follow-370 371 up, and/or from more aggressive therapies as the first, and second line treatments.

372

PME-1 expression protected PCa cells from anoikis in various anchorage-independent conditions (soft agar, trypsin-detachment), or under conditions with low ECM-cell mechanotransduction (low adhesion plates, low stiffness). We also demonstrate for the first time that PP2A modulation regulates nuclear deformability. Our results are fully consistent with the published model that different levels of nuclear lamina

378 deformability support different cellular functions (Figure 4H). Whereas a stiff lamina 379 prevents cell migration through physical restrictions, a highly deformable lamina 380 renders cells vulnerable to apoptosis induction by low tissue tension, and fluid stress 381 (9, 12, 15, 17, 45). Further, a recent study indicates that aggressive prostate cancer 382 cells such as PC-3 and DU-145 have softer nuclei than immortalized benign prostate 383 epithelial cells (46). As we demonstrate an important role for PME-1 in preventing 384 further softening of the nuclear lamina in PC-3 cells, and link this to the anoikis 385 suppression, we speculate that during cellular transformation nuclear lamina becomes 386 softened to allow cell migration, whereas PME-1 overexpression in cancerous cells 387 prevents softening of their nuclear lamina to the levels that would make them 388 sensitized to anoikis (Figure 4H). Although alternative mechanisms downstream of 389 PME-1 may contribute to its anoikis suppressing function, we emphasize that nuclear 390 lamina deformability regulation may have been overlooked as a critical mechanism 391 supporting viability of PCa cells during their progression towards full malignancy. 392 Importantly, although Lamin-A/C phosphor-antibody analysis was used to validate 393 PP2A-mediated regulation of nuclear lamin proteins, it is very likely that PP2A-394 mediated dephosphorylation of other detected nuclear lamina proteins are important 395 for both the observed nuclear deformability phenotype. In addition to phosphotargets 396 of PME-1, PME-1 also interacts with a FDPS(30) which contributes to farnesylation of 397 Lamin-A/C, B1 and B2(37). Therefore also this nuclear lamina-associated PME-1 398 protien interaction may contribute to the observed nuclear deformability regulation. 399 The molecular mechanisms by which too deformable nuclei sensitizes apoptosis is 400 currently under debate. As nuclear chromatin is tightly connected with lamina (47). 401 and DNA arrangements sensitize to apoptosis, it is possible that dephosphorylation of

402 chromatin arranging LAP2A/B proteins may contribute to anoikis induction in PME-1
403 negative cells under low attachment conditions.

404

405 Together these results identify anoikis resistance as a candidate mechanism by which 406 PME-1-mediated PP2A inhibition promotes malignant progression of prostate cancer. 407 Together with emerging orally bioavailable PP2A reactivating compounds with 408 profound antitumor activity in in vivo PCa models (32), these results clearly emphasize 409 future importance of comprehensive understanding of PP2A biology for management 410 of aggressive prostate cancer patients. The results also identify PME-1 as a potential 411 biomarker for improved stratification and better therapy response among PCa patients 412 with complete PTEN loss. Together with other findings, results further indicate that 413 diagnostic PME-1 evaluation from PTEN negative tumors may help to identify patients 414 that would benefit from more intensive follow-up after therapy.

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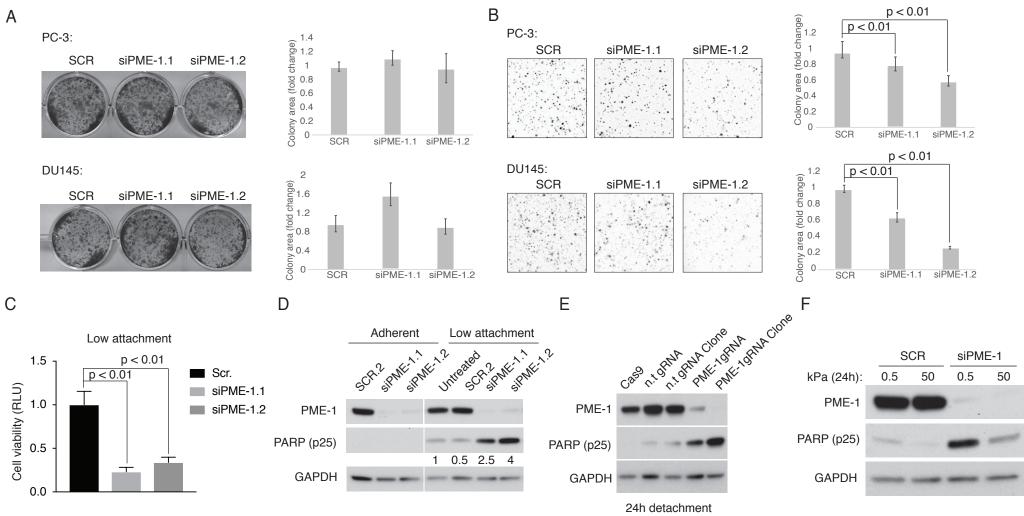
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425 Acknowlegdements

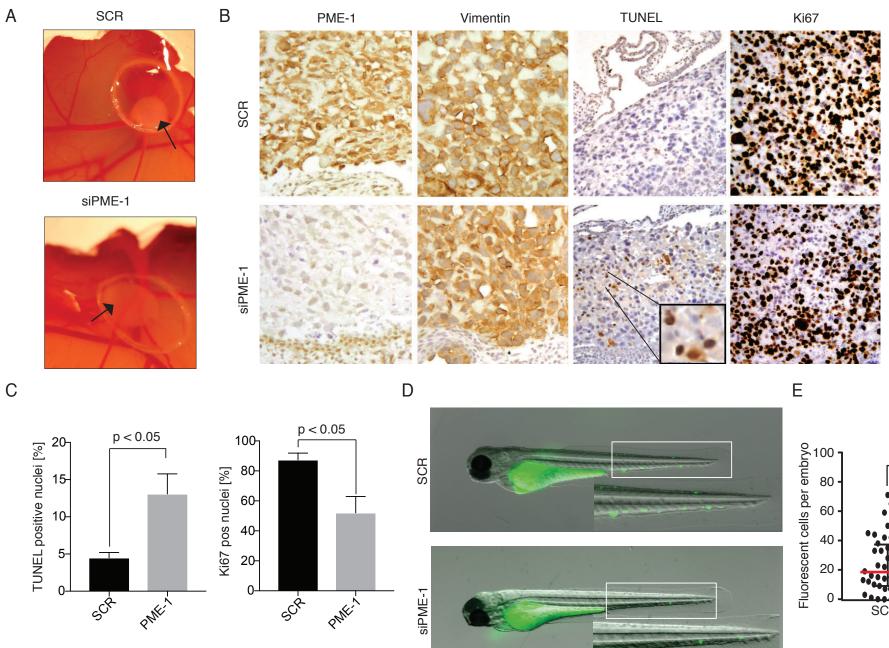
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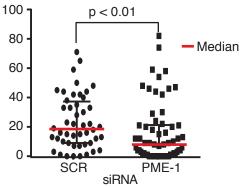
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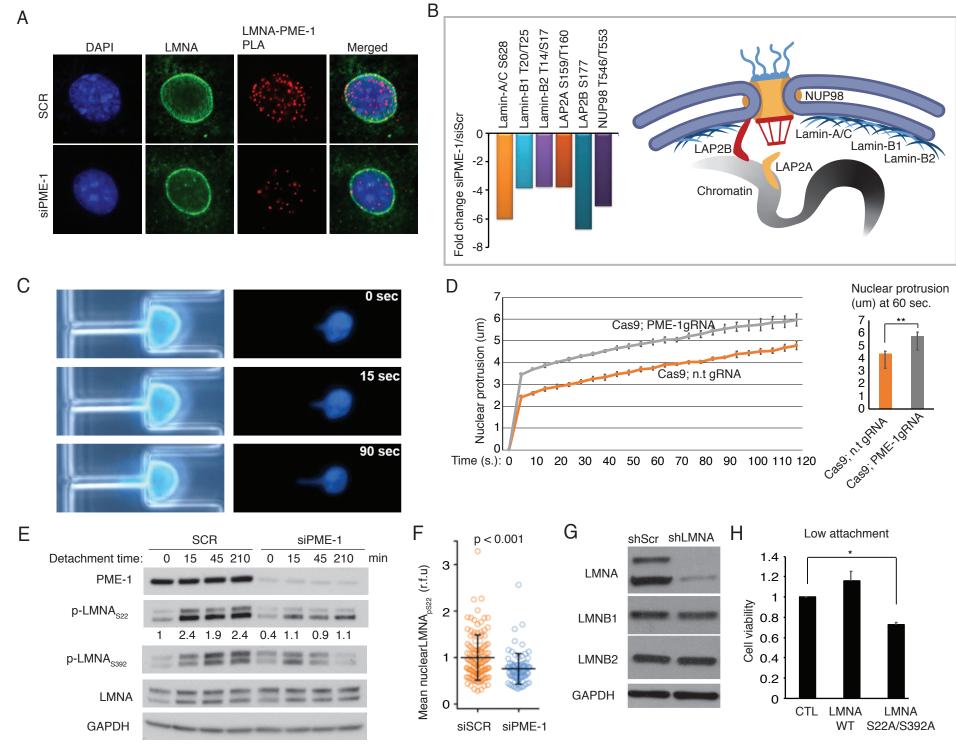


Figure 3

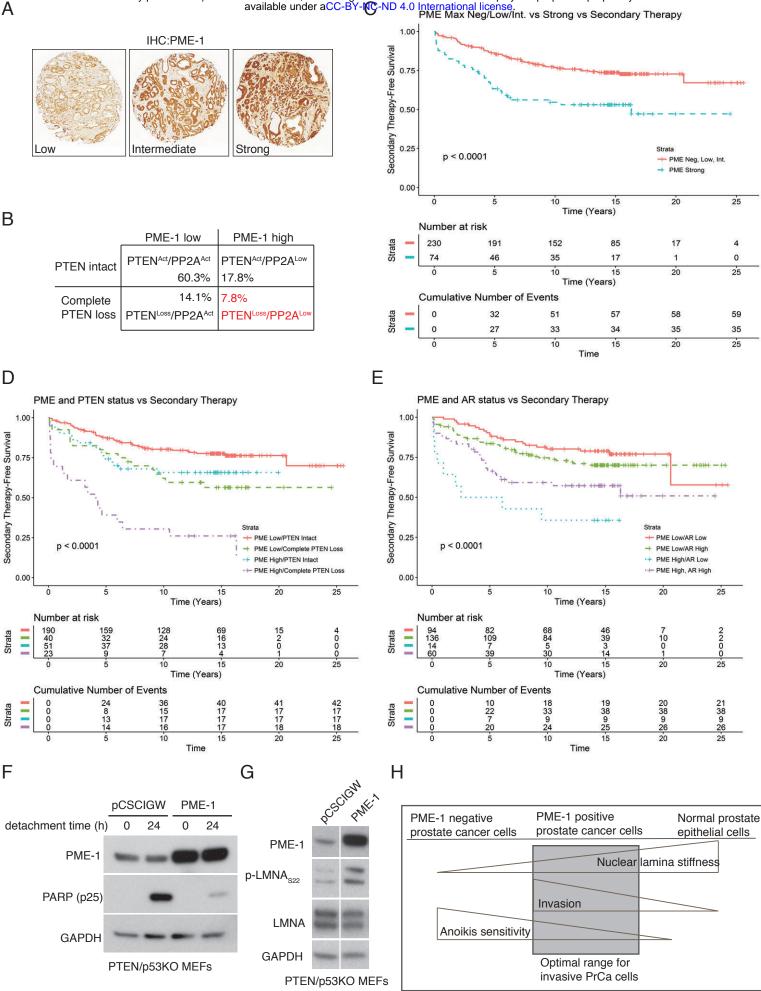


Table 1. Demographics of the radical prostatectomy patient cohor	t.
	Total Cohort (n=358)
Age at RP, years (median, IQR) (n = 358)	64.0 (59.2 – 67.9)
Preoperative PSA, ng/ml (n, %) (<i>n</i> = 283)	
≤10.0	143 (50.5)
10.1-20.0	89 (31.45)
>20.0	51 (18.0)
Grade group at RP (n, %) (n = 358)	
1	93 (26.0)
2	93 (26.0)
3	114 (31.8)
4	45 (12.6)
5	13 (3.6)
pT (n, %) (<i>n</i> = 334)	
2	202 (60.5)
≥3	132 (39.5)
Lymph node status (n, %) (n = 358)	
Negative	342 (97.2)
Positive	10 (2.8)
Follow-up time after RP, years (median, range) (n=358)	15.7 (0.7-28.6)
Death from any cause $(n, \%)$ $(n = 358)$	172 (48.0)
Death from prostate cancer (n, %) ($n = 358$)	33 (9.2)
Patients receiving secondary therapy after RP (n, $\%$) (n = 358)	124 (34.6)

Table 2. Univariate Analysis of PME IHC and clinical v			
	PME IHC Low (n = 238)	PME IHC High (n = 82)	P-Value
Age at RP, years (mean, SD) (n = 320)			
< 60	81 (34.0)	24 (29.3)	0.490ª
60 - 70	132 (55.5)	46 (56.1)	
> 70	25 (10.5)	12 (14.6)	
Preoperative PSA, ng/ml (n, %) (n = 254)			
≤10.0	100 (52.1)	32 (51.6)	1.000 ª
10.1-20.0	59 (30.7)	19 (30.7)	
>20.0	33 (17.2)	11 (17.7)	
Grade group at RP (n, %) (<i>n</i> = 320)			
1	74 (31.1)	5 (6.1)	< 0.001 ª
2	65 (27.3)	17 (20.7)	
3	73 (30.7)	36 (43.9)	
4	22 (9.2)	17 (20.7)	
5	4 (1.7)	7 (8.6)	
pT (n, %) (<i>n</i> = 299)			
2	150 (67.3)	28 (36.8)	< 0.001ª
3-4	73 (32.7)	48 (63.2)	
Lymph node status (n, %) (n = 314)			
Negative	228 (97.4)	78 (97.5)	0.974ª
Positive	6 (2.6)	2 (2.5)	
Secondary Therapy (n, %) (n = 320)			
No	171 (71.8)	39 (47.6)	< 0.001ª
Yes	67 (28.2)	43 (52.4)	
Death from any cause (n, %) (n = 320)			
Yes	104 (43.7)	46 (56.1)	0.055ª
No	134 (56.3)	36 (43.9)	
Death from prostate cancer (n, %) (n = 320)			
Death due to PC	20 (8.4)	11 (13.4)	0.197ª
Alive, or dead from other causes	218 (91.6)	71 (86.6)	
^a Fisher's exact test			

440 **FIGURE LEGENDS**

441 Figure 1. PME-1 inhibits anoikis in PTEN-deficient prostate cancer cells

A. The effect of PME-1 depletion, utilizing two independent siRNAs, was characterized
by colony formation assays (10 days of growth) in two PTEN-deficient PCa cells lines
PC-3 and DU-145. The graphs show the quantified colony area, as mean fold change

445 +/- SD compared to siScr. N=3.

B. PME-1 knock-down effect on anchorage-independent growth in soft agar assays
(14 days of growth) in both PC-3 and DU-145 cells. The graphs show the quantified
colony area, as mean +/- S.D. fold change compared to siScr. N =2, p-value: 2-sided
t-test between colony areas in 15-20 image fields.

450 **C**. PME-1 inhibition effect on cell viability of PC-3 on low attachment plates, as 451 measured by Cell Titer-GLO luminescence assay. The relative cell viability was 452 calculated as compared to siScr (100%). N=2, Shown in mean + S.D, 2-sided t-test.

D. siScr and siPME-1 transfected PC-3 cells were plated 72h post-transfection on
normal or low attachment plates for 24h, before collection and lysis, and subsequently
analyzed by Western blot for cPARP induction.

E. Apoptosis induction, as measured by PARP cleavage, was studied by Western blot
analysis in CRISPR/Cas9 generated PC-3 cells, with and without non-targeting gRNA
(lanes 1-3), a pool of PME-1 gRNA transfected cells (lane 4), and a single cell subclone
of PME-1 targeted cells (lane 5), after 24h detachment.

F. Mechanosensitive PARP cleavage in PME-1-depleted PC-3 cells. The cells were
cultured on soft (0.5 kPa) or stiff (50 kPa) hydrogel for 24h before being scraped into
PBS, spun down, lyzed and analyzed for protein expression. Results are
representative of two independent experiments.

464

Figure 2. PME-1 supports *in vivo* anoikis resistance and survival of prostate cancer cells in circulation.

A. The effect of PME-1 effect on anchorage-independent growth of PC-3 cells *in vivo*was tested by Chicken embryo chorioallantoic membrane (CAM) assay. PC-3 cells
were transiently transfected with either siScr or siPME-1, and 24h post-transfection
placed on the CAM. Growth of tumors was followed for 3-5 days.

B-C. Immunohistological stainings of dissected tumors were conducted using
antibodies for PME-1, VIM, TUNEL and Ki67, and subsequently the percentage of
TUNEL- and Ki67-positive were quantified. Shown is mean + S.E.M, p-value; Mann
Whitney test.

D-E. The survival of siScr and siPME-1 transfected PC-3 was studied by microinjecting
them into the common cardinal vein of zebrafish embryos 72h post-transfection. After
overnight incubation the embryos were imaged by fluorescence stereomicroscopy.
The number of surviving fluorescent tumor cells per was counted manually from the
images using FIJI and statistical analysis was performed using non-parametric
Kruskal-Wallis test.

481

482 Figure 3. PME-1 associates with nuclear lamina and regulates its deformability

483 A. Proximity ligation assay (PLA) was used to confirm the physical association of PME-

484 1 with Lamin-A/C in PC-3 cells. Cleraly decreased PLA signal in PME-1-1 transfected

485 cells confirm the specificity of PLA reaction. DNA was stained using DAPI

B. PME-1 depletion by transient siRNA transfection induced dephosphorylation of the indicated nuclear lamina constituents in HeLa cells. Shown is Log2 fold change in abundance of the indicated phosphopeptide 72 hours after transfection. Included are phosphopeptides detected with 1% FDR accuracy and for which FDR for

490 reproducibility between 3 parallel samples was < 10%. The data was retrieved from491 (28).

492 **C-D**. CRISPR/Cas9 generated control and PME-1 KO cells were analyzed by 493 micropipette aspiration, showing representative images (C) and quantification (D) of 494 nuclear protrusions (um at different time-points).

E. The effect of PME-1 knock-down in PC-3 cells on the phosphorylation of Lamin-A/C
S22 ans S392 was confirmed by Western blot analysis, after trypsin-induced
detachment for the indicated time (0-210 minutes) prior to collection and lysis of
samples. The signal was quantified as compared to total Lamin-A/C and GAPDH.

F. Quantification depicting the phosphorylation of Lamin-A/C S22 in PME-1-depleted
and control PC-3 cells, 48h post-transfection and 24h after seeding on soft hydrogel
(2 kPa PAG). After culture the cells were fixed, stained and analyzed by confocal
microscopy. Mean +/- SD, Mann-Whitney test, n = 84-117 cells from two independent
experiments.

G-H. PC-3 cells, in which the endogenous Lamin-A/C expression was suppressed by shRNA was analyzed by Western blot, and upon confirmation of Lamin-A/C suppression (E) transiently transfected with either wild-type Lamin-A/C (WT) or with a Lamin-A/C mutant with both serine 22 and 392 mutated to alanine (S22A/S392A). The effect of Lamin-A/C phosphorylation on cell viability was analysed, as measured by CellTiter-GLO assay. Shown in mean + S.D, 2-sided t-test, n=2.

510

511 Figure 4. PME-1 overexpression associates with PTEN loss and therapy relapse

512 of PTEN negative PCa patients

A. PME-1 protein expression in PCa tissue microarray material from 358 patients
 treated primarily with radical prostatectomy in the Helsinki University Hospital between

515 1983 and 1998 was assessed by immunohistochemical (IHC) stainings. PME-1 protein
516 content was scored using 4-tier scale; negative (not shown), low, intermediate and
517 strong expression.

518 **B**. Assessment of the activity status of PTEN and PP2A in the patient populations by

519 using genetic deletion of PTEN and PME-1 overexpression as surrogate markers

520 **C-E**. Kaplan-Meyer analysis of time to secondary therapies after primary treatment

based on PME-1 status alone (C) and in combinations with PTEN deletion (D) and AR
expression (E).

F) Effect of stable PME-1 overexpression in anoikis in mouse embryo fibroblasts (MEFs) from a PTEN/p53 KO mouse model after 24h incubation in low attachment conditions. Apoptosis induction was assessed by Western blot utilizing antibodies for cleaved PARP.

527 G. Effect of stable PME-1 overexpression in S22 phosphorylation of Lamin-A/C in
 528 PTEN/p53 KO MEFs.

H. Schematic representation of prostate cell properties, such as nuclear lamina stiffness, invasion and anoikis sensitivity, in relation to PME-1 protein levels and cell malignancy. Based on the results and existing literature, we propose that during cellular transformation nuclear lamina becomes softened to allow cell migration, whereas PME-1 overexpression prevents excess softening of the lamina to the levels that would threaten the viability of these cells.

535

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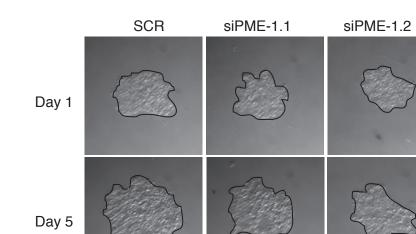
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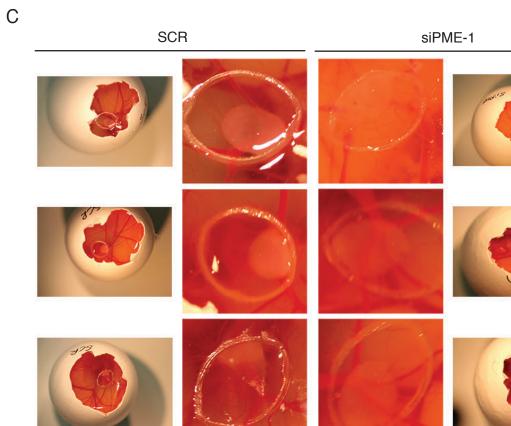
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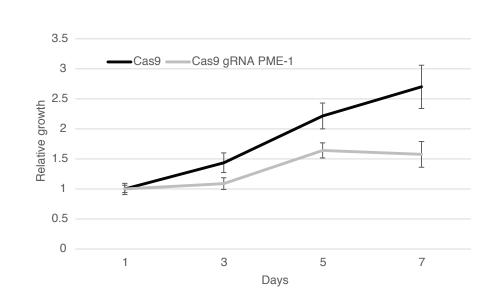
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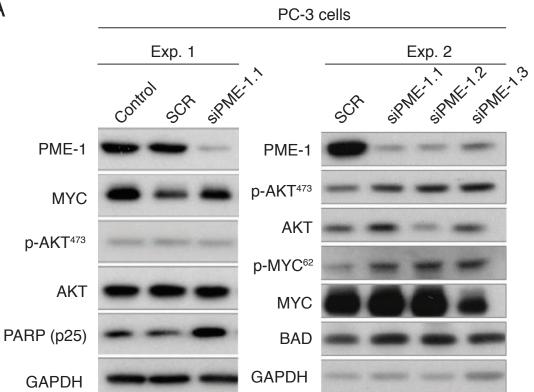
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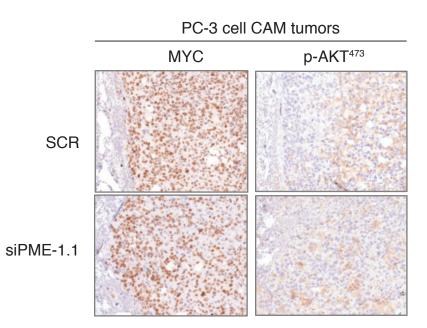
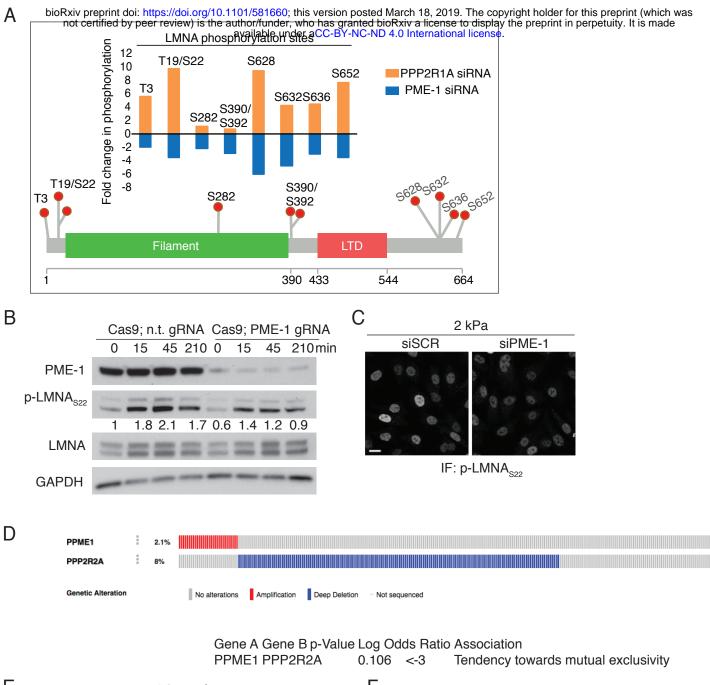
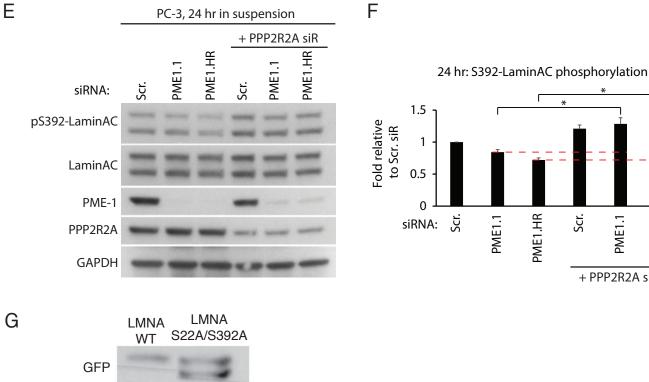


Figure S2





GAPDH

Figure S3

PME1.1

+ PPP2R2A siRNA

PME1.HR

Scr.

PME1.HR

А

PME vs PTEN St	atus.				PME vs A	R Status.				PME vs ERG Status.				
	PME Low	PME High	Total	Р		PME Low	PME High	Total	Р		PME Low	PME High	Total	Р
PTEN Intact	193 (81.1)	57 (69.5)	250	0.043 ^ª	AR Low	97 (40.8)	14 (17.1)	111	< 0.001 ^ª	ERG Negative	128 (53.8)	27 (32.9)	155	0.001 ^ª
Complete PTEN	45 (18.9)	25 (30.5)	70		AR High	141 (59.2)	68 (82.9)	209		ERG Positive	110 (46.2)	55 (67.1)	165	
Total	238	82	320		Total	238	82	320		Total	238	82	320	

^a Fisher's exact test

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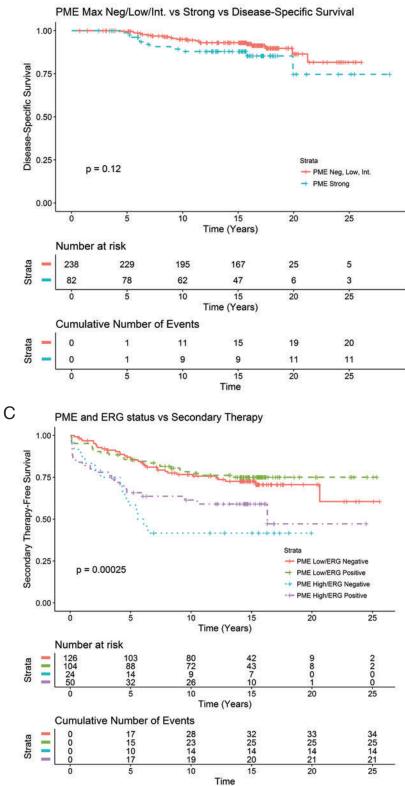


Figure S4

694 SUPPLEMENTARY FIGURE LEGENDS

695

696 Figure S1. PME-1 supports anoikis resistance of PC-3 cells

- 697 A. Colony growth assays of PC-3 cells were conducted on low attachment plates, after
- 698 PME-1 knock-down by two independent siRNAs, and the growth was compared to
- 699 siScr transfected cells.
- 700 **B**. The effect of PME-1 KO, by CRISPR/Cas9, on cell growth on low attachment plates
- 701 was studied for seven days.
- 702 C. Chicken embryo chorioallantoic membrane (CAM) assay was used to test the effect
- 703 of PME-1 on the anchorage-independent growth of PC-3 cells in vivo. After transient
- transfections the cells were placed on the CAM, and the growth was followed for 3-5
- 705 days. The figure shows three representative pictures of siScr and siPME-1 tumors.
- 706

Figure S2. PME-1 silencing has no effect AKT or MYC signaling in PC-3 cells *in vitro* or *in vivo*

- 709 **A-B**. The effect of PME-1 silencing on AKT and MYC phosphorylationwas assayed in
- vitro by Western blot (A) and *in vivo* by immunohistochemistry on CAM tumors (B).
- 711

712 Figure S3. PME-1 inhibits PP2A-mediated Lamin-A/C dephosphorylation

- 713 A. Eight Lamin-A/C phosphopeptides/10 phospho-sites, negatively regulated by PME-
- 1 depletion and positively regulated by PP2A inhibition (siRNA targeting the scaffold
- subunit PPP2R1A) as identified by a LC-MS/MS phosphoproteome screen (32).

B. The effect of PME-1 KO by Crispr/Cas in PC-3 cells on the phosphorylation of
Lamin-A/C S22 after trypsin-induced detachment. The Lamin-A/C S22 signal intensity
was guantified and compared to total Lamin-A/C levels.

C. Immuofluoresence analysis of phosphorylation of Lamin-A/C S22 in PME-1depleted and control PC-3 cells, 48h post-transfection and 24h after seeding on soft
hydrogel (2 kPa). After culture the cells were fixed, stained and analyzed by confocal
microscopy. Quantification of the data is shown in figure 3F.

D. Analysis of human PCa samples in cBioPortal shows loss of PPP2R2A in 8% of the
samples.

725 E,F. To show the PP2A-dependence of Lamin-A/C phosphorylation regulation by

PME-1, PC-3 cells were transiently depleted of PME-1 and co-depleted of the PP2A

727 B-subunit PPP2R2A. After transfection, cells were cultured on low attachment plates

for 24h prior to sample collection, lysis and subsequent analysis of p-Lamin-A/C, PME-

1 and PPP2R2A as compared to GAPDH by Western blot. (F) Relative change as
compared to siRNA was quantified.

G. Western blot analysis of GFP-tagged Lamin-A/C WT and S22A/S392 mutant
 overexpression in PC-3 cells silenced for endogenous WT-Lamin-A/C.

733

734 Figure S4. PME-1 overexpression associates with total PTEN loss in PCa

735 **A**. PME-1 status was correlated to previously assessed PTEN, ERG and AR status.

736 PME-1 expression significantly associated with complete PTEN loss, but also with

737 ERG positivity and high AR expression status, as analysed by Fisher's exact test.

B. Association of PME-1 status and disease-specific survival in the PCa tissue
microarray material from 358 patients treated primarily with radical prostatectomy in
the Helsinki University Hospital between 1983 and 1998.

- 741 **C.** Kaplan-Meyer analysis of time to secondary therapies after primary treatment
- 742 based on PME-1 status in combination with ERG.

743

745 Material and Methods

746

- 747 Cell lines
- PC-3 and DU-145 prostate cancer cell line were obtained from ATCC and cultured in
- 749 RPMI. Pten^{Δ/Δ}; Trp53^{Δ/Δ}; Isl-tdTomato mouse embryonic fibroblasts were provided by
- Lloyd Trotmann and cultured in DMEM. All growth media were supplemented with 10%
- 751 heat-inactivated FBS (Biowest), 2 mmol/L l-glutamine, and penicillin (50
- 752 units/mL)/streptomycin (50 µg/mL; pen/strep). Cells were confirmed to be
- 753 mycoplasma-free and grown at 37° in a humidified atmosphere of 5% CO².
- 754 siRNAs
- 755 Scrambled (SCR.2): CGU ACG CGG AAU ACU UCG A
- 756 PME-1.1 (PME-1.3): GGA AGU GAG UCU AUA AGC A
- 757 PME-1.2 (PME-1 hr2): GAA UGA AAC UGG CAA GGA U
- 758 PP2R2A: CUG CAG AUG AUU UGC GGA U

759 Validated PP2A B-subunit siRNA was purchased from Qiagen, PME-1 siRNAs from

- 760 Eurofins Genomics (MWG).
- 761 LMNA: ON-TARGETplus human siRNA SMARTpool (L-004978-00-0005)
- 762 Antibodies

The following antibodies were used at the indicated dilutions: PME-1: Santa Cruz Biotechnology sc-20086 (H-226) Western blotting 1:1000; PME-1: Santa Cruz Biotechnology sc-25278 (B-12) Immunohistochemistry 1:1000, Immunofluorescence 1:100; PPP2R2A: Cell Signalling Technology #5689 Western blotting 1:1000; cleaved PARP: Abcam ab32064 [E51] Western blotting 1:1000; GAPDH: Hytest 5G4-6C5 Western blotting 1:500.000 c-MYC: Abcam ab32072 [Y69] Western blotting 1:1000; p-Myc S62:Abcam ab78318 Western blotting 1:1000; Akt1/2/3: Cell signalling #9272

770 Western blotting 1:2000; p-Akt S473: Cell Signalling Technology #4060 Western 771 blotting 1:1000; p-Akt T308: Cell Signalling Technology #13038 Western blotting 772 1:1000; Lamin A/C: Santa Cruz Biotechnology sc-20681 (H-110) Western blotting 773 1.10,000; Lamin A/C: Santa Cruz Biotechnology sc-6215 (N-18) Western blotting 1:1000 Immunofluorescence 1:100; Lamin A/C: Santa Cruz Biotechnology sc-7292 774 775 (636) Immunofluorescence 1:250; p-Lamin S22: Cell Signalling Technology #2016 776 Western blotting: 1:500 Immunofluorescence 1:50 or 1:100; p-Lamin S392: Abcam 777 ab58528 Western blotting 1:5000, Immunofluorescence 1:200

778 siRNA/plasmid transfection

For siRNA transfections Lipofectamine RNAi MAX (Thermo Fisher Scientific) was
used following the manufacturer's instructions. In most cases 2.5x10⁵ cells were
seeded on a 6-well plate one day before transfection to reach 60-70% confluency.
Cells were then transfected with 25pmol siRNA, 7.5µl siRNA MAX per 6-well and
assayed 48-72h after transfection unless otherwise stated.

For plasmid transfections Lipofectamine 3000 (Thermo Fisher Scientific) was used
following the manufacturer's instructions. In most cases 3-4 x10⁵ cells were seeded
on a 6-well plate one day before transfection to reach 80-90% confluency. Cells were
then transfected with 2.5µg DNA, 5µl P3000 reagent, 3.75µl Lipofectamine 3000 per
6-well and assayed 24-48h after transfection unless otherwise stated.

789 Colony formation assay

Cells were reseeded at low confluency (1-4x10³) on a 12-well dish 24h after siRNA transfection and grown for around 10 days to allow formation of cell colonies. Colonies were then fixed with ice-cold methanol and stained with 0.2% crystal violet solution (made in 10% ethanol) for 10 minutes at room temperature each. Excess stain was removed by repeated washing with PBS. Plates were dried and scanned with Epson

795 perfection V700 scanner. Quantifications were performed with ColonyArea ImageJ

plugin (48), and graphs were plotted using the area % values.

797 Anchorage-independent colony formation assay

798 For the anchorage-independent colony formation assay, which correlates typically with 799 in vivo tumorigenicity, 2x10⁴ cells were resuspended in 1.5ml growth medium 800 containing 0.4% agarose (4% Agarose Gel, Termo Fisher Scientific Gibco; top layer) 801 and plated on 1ml bottom layer containing growth medium and 1.2% agarose in a 12-802 well plate. After 14 days of growth, colonies were stained over night with 1mg/ml Nitro 803 blue tetrazolium chloride (NBT; Molecular Probes) in PBS. Colonies were imaged 804 using a Zeiss SteREO Lumar V12 stereomicroscope. Analysis was done using the 805 ImageJ software. First, the background was subtracted using the rolling ball function 806 with a radius of 50µm, then auto-thresholding was applied to separate the colonies. 807 Area percentage was calculated using the ImageJ built-in function 'Analyze Particles' with exclusion of particles smaller than 500µm² that are not considered colonies. 808

809 Cell viability assay on low attachment plates

Cell viability on 96 Well Clear Black Round Bottom Ultra Low Attachment Spheroid Microplates (Corning) was determined by using the CellTiter-Glo[®] Luminescent Cell Viability Assay (Promega). For this purpose, a small number of cells (250-1000/well) were seeded on the plates and grown for the indicated number of days. Cells were then lysed in the CellTiter-Glo[®] reagent by vigorous pipetting and luminescence was read with a Synergy H1 reader.

816 Western blotting

Western blot protein lysates were prepared in 1x RIPA buffer (150 mM sodium
chloride, 1.0% NP-40, 0.5% sodium deoxycholate, 0.1% SDS and 50 mM Tris, pH 7.5)
containing PhosSTOP[™] phosphatase and cOmplete[™], EDTA-free protease inhibitors

820 (Roche). The DNA in the protein samples was sheared by sonication and the protein amount was estimated using the Pierce[™] BCA Protein Assay Kit (Thermo Fisher 821 822 Scientific). Lysates were usually separated on 4–20% Mini-PROTEAN® TGX[™] Gels 823 (Biorad) and transferred by wet blotting to PVDF membranes (Millipore). Unspecific 824 antibody binding was blocked with 5% non-fat dry milk in TBS-T. Incubation for most of the primary antibodies was performed overnight at 4°C in either 5% non-fat dry milk 825 826 or 5% BSA for most of the phospho-specific antibodies. For detection, HRP- labelled 827 secondary antibodies (DAKO) followed by incubation with Pierce™ECL Western 828 Blotting Substrate (Thermo Fisher Scientific) was used.

829 Generation of Pten^{Δ/Δ}; Trp53^{Δ/Δ} mouse embryonic fibroblasts

Pten^{Δ/Δ}; Trp53^{Δ/Δ}; IsI-tdTomato MEFs were generated as previously described (49). To
stably overexpress PME-1 in those cells, PME-1 cDNA was cloned into pCSCIW-2
lentiviral construct which was then packaged into lentivirus. Following transduction,
GFP positive cells were sorted and overexpression of PME-1 was confirmed by
western blotting.

835 CRISPR/Cas9 mediated PME-1 knock-out

To generate PME-1 knock-out cells, PC-3 cells were transduced with lentivirus 836 containing lentiCas9-Blast construct (Addgene, Feng Zhang lab) and selected with 837 838 growth medium containing 4µg/ml blasticidin for about one week. In a second round, cells were infected with lentivirus sgRNA against PME-1 exon 3 (sequence: 839 840 ACTTTTCGAGTCTACAAGAGTGG) cloned into pKLVflipedU6gRNA PB BbsI PGKpuro2ABFP construct. After 841 puromycin selection 842 (2µg/ml in growth medium), PME-1 KO on the protein level was confirmed by western 843 blotting. Single cell clones were then obtained by single-cell sorting and the knock-out confirmed sequencing (Primers. Forward 844 was by

845 CACCGCTTTTCGAGTCTACAAGAGGT;

reverse

TAAAACGAAGATCTGTCTGCAGAAAC). Cas9, non-targeting sgRNA expressing
cells served as a negative control in the functional assays.

848 Chick chorioallantoic membrane (CAM) assay

849 To start chick embryonic development for the chorioallantoic membrane assay, 850 fertilized eggs were kept rotating in an incubator at 37°C and 50-60% humidity for four 851 days. After that initial incubation time, a small hole was introduced at the sharp edge 852 of the eggs and sealed with parafilm. Following four more days of incubation on 853 stationary racks, the hole was enlarged and a plastic ring was placed on top of blood 854 vessels in the chorioallantoic membrane. Next, 1x10⁶ PC-3 cells in 20µl volume of a 1:1 mixture of ice-cold PBS and matrigel were pipetted inside the ring on the 855 856 membrane. The hole was then covered with parafilm and the eggs were incubated for 857 three more days. At day twelve of embryonic development, the animals were sacrificed 858 by freezing the eggs for 15min and the tumor cell mass was dissected from the 859 membrane and processed for further analysis.

860 Immunohistochemistry

861 Hematoxylin/eosin staining and immunohistochemistry were performed on 3-um-thick sections of 4% paraformaldehyde-fixed and paraffin-embedded tissues. Following 862 rehydration, endogenous peroxidase was blocked by incubation in 50% MetOH, 1% 863 864 H_2O_2 . Subsequent antigen retrieval was performed with the 2100 Retriever (Aptum) in 865 R- Universal Buffer. Unspecific antibody binding was blocked with 10% goat serum in 2% BSA/PBS prior to overnight incubation at 4°C with the primary antibody. For 866 867 detection, the DAKO EnVision peroxidase system, followed by incubation with 0.01% 868 diaminobenzidine (Sigma-Aldrich) was used.

869 Zebrafish *in vivo* dissemination assay

870 The xenotransplantation of zebrafish embryos was performed as described in detail in (35) with some modifications. PC-3 cells were washed with PBS, and stained with 871 872 CellTracker Green CMFDA dye (5uM, Thermo Fisher Scientific) and detacher and 873 detached with trypsin-EDTA in a single incubation step at +37°C. Subsequently, cells 874 were pelleted by centrifugation and washed with PBS twice. This was followed by 875 filtration through 40µM mesh into FACS tube (BD Falcon, 352235) and pelleting cells 876 by centrifugation. Finally, cells were resuspended into 30µl of injection buffer (2%PVP 877 in PBS) and kept on ice until transplanted.

878 Zebrafish (*Danio rerio*) of pigmentless casper strain ($roy^{-/-}$; mitfa^{-/-})(50) was 879 used in the experiments under licence no. MMM/465/712-93 (issued by Finnish 880 Ministry of Forestry and Agriculture) and following legislation: the European 881 Convention for the Protection of Vertebrate Animals used for Experimental and other 882 Scientific Purposes and the Statutes 1076/85 and 62/2006 of The Animal Protection 883 Law in Finland and EU Directive 86/609. The embryos were obtained through natural 884 spawning, and embryos were cultured in E3-medium (5 mM NaCl, 0.17 mM KCl, 0.33 885 mM CaCl₂, 0.33 mM MgSO₄) at +33°C. At 2 or 3 days post-fertilization, the embryos 886 were anesthesized with 200mg/ml Tricaine and embedded in 0.7% low-melting point 887 agarose. Subsequently, the cell suspension was microinjected into common cardinal 888 vein (duct of Cuvier) of the embryo using glass capillaries (Transfertip), CellTramVario 889 microinjector and InjectMan micromanipulator (all from Eppendorf). Embryos were 890 liberated from the agarose gel using forceps and successfully transplanted embryos 891 were selected to the experiment. After overnight incubation at +33°C, the embryos 892 were anesthesized again with Tricaine and imaged using Zeiss StereoLumar V12 893 fluorescence stereomicroscope. The number of surviving cells was counted manually 894 from the images using FIJI and statistical analysis was performed using non-

895 parametric Kruskal-Wallis test (GraphPad Prism 6.05 software, GraphPad Software,

896 Inc.).

897 **Proximity ligation assay (PLA)**

898 Cells were grown to about 80% confluency on sterilized coverslips, fixed for 10min in 899 4% PFA and permeabilized for 10min with 0.5% Triton X-100 in TBS for proximity 900 ligation assay (PLA). The following steps were completed following the manufacturer's 901 instructions (Sigma-Aldrich). Briefly, unspecific antibody binding was blocked with the 902 provided blocking solution for 30min. The slides were then incubated with the primary 903 antibodies overnight at 4°C, with the mouse and rabbit probes for one hour at 37°C, 904 with the ligation mix for 30min at 37°C and the amplification mix for 100min at 37°C. 905 The washing steps in-between the individual steps were carried out with Buffer A 906 (Sigma), the final washing step with Buffer B (Sigma). Slides were mounted in mowiol 907 mounting medium and imaged with a Zeiss LSM780 confocal microscope.

908 Immunofluorescence

909 For immunofluorescence cells were fixed for 10min with 4% PFA and permeabilized 910 for 10min with 0.5% Triton X-100 in TBS. Unspecific antibody binding was blocked by 911 incubation with 10% serum from the host of the secondary antibody in 2% BSA/PBS 912 for 30min. Primary antibodies were diluted in 2% BSA/PBS and cells were usually 913 incubated with the primary antibody over-night at 4°C. As secondary antibodies 914 appropriate Alexa-Fluor conjugates (Thermo Fisher Scientific) were used at a 1:2000 915 dilution in PBS for 30min. Nuclei were stained with DAPI (4',6-diamidino-2-916 phenylindole).

917 Polyacrylamide hydrogels

35 mm glass bottom dishes (MatTek Corporation, P35G-1.0-14-C) were treated with
100 µl of Bind-silane solution [7.14% Bind-silane (Sigma, M6514) and 7.14% acetic

920 acid in absolute ethanol] for 15 min, washed twice with absolute ethanol and left to dry completely. A pre-polymer mix comprising 5.6% acrylamide (Sigma) and 0.078% N,N'-921 922 methylenebisacrylamide (Sigma) in PBS was prepared to obtain hydrogels with an 923 elastic modulus of 1.8-2 kPa. Polymerization was initiated by adding 2.5 µl of 20% ammonium persulfate (Bio-Rad) and 1 µl of N,N,N',N'-tetramethylethylenediamine 924 925 (Sigma). The solution was vortexed, 13 µl was added on top of the glass bottom dish, 926 a 13 mm glass coverslip was placed on the drop and the gel was left to polymerize for 927 1 h at room temperature. After polymerization, the dish was filled with PBS and the 928 coverslip was carefully removed. Hydrogels were made permissive for protein binding 929 by incubating them in 500 µl of Sulfo-SANPAH solution [0.2 mg/ml Sulfo-SANPAH 930 (Sigma, 803332) and 2 mg/ml N-(3-dimethylaminopropyl)-N'-ethylcarbodiimide 931 hydrochloride (Sigma, 03450) in 50 mM HEPES] for 30 min on slow agitation, followed 932 by a 10 min UV exposure (~30 mW/cm², 253.7 nm). Activated gels were washed three 933 times with PBS to get rid of residual Sulfo-SANPAH.

Alternatively, pre-activated polyacrylamide hydrogels of variable stiffness were ordered from Matrigen Life Technologies. 35 mm glass bottom dishes (SV3510-EC-0.5, SV3510-EC-50) were used for imaging and 6-well plates (SW6-EC-0.5, SW6-EC-50) for growing cells for lysis. All hydrogels and corresponding plastic controls were functionalized with bovine plasma fibronectin (Merck-Millipore, 341631) and collagen type I (Sigma, C8919) by incubating the dishes in 5 μ g/ml of each protein for 1-2 h at +37 °C, or overnight at +4 °C, before use.

For Western blot analysis from cells in hydrogels, 0.5 and 50 kPa hydrogel-coated 6well plates were immersed in cell culture medium for 30 min and seeded with 300,000 PME-1 or control siRNA-treated PC-3 cells per well. The cells were incubated on the gels for 24 h. Once the cultures reached 80-90% confluency, the cells were washed

once and scraped into cold PBS, spun down and lysed in radioimmunoprecipitation
assay (RIPA) buffer.

For immunofluorescence experiments, PME-1 or control siRNA transfected cells were 947 948 seeded on 2 kPa hydrogel-coated glass bottom dishes and incubated for an additional 949 24 h before fixing. Cells were fixed with 4% PFA for 10 min at room temperature, and 950 simultaneously permeabilized and blocked with 0.3% Triton in 10% horse serum 951 (Gibco) for 15 min at room temperature. All samples were incubated in primary 952 antibodies overnight at +4 °C, and stained with secondary antibodies for 1-2 h at room 953 temperature on the following day. The antibodies were diluted in 10% horse serum 954 before use.

955 Microscopy and image analysis

Fluorescent specimens were imaged using a laser scanning confocal microscope
LSM780, controlled by Zen 2010 (Zeiss), and the objective used was a 40x/1.2 W CApochromat objective (Zeiss). Images were analyzed using ImageJ (National
Institutes of Health) and CellProfiler (Broad Institute) softwares.