

Low temperature perception by FER-TORC1 triggers root hair growth

1 **Cell surface receptor kinase FERONIA linked to nutrient sensor TORC1 signaling**
2 **controls root hair growth at low temperature in *Arabidopsis thaliana***

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35 **Abstract**

36
37 Root hairs (RH) are excellent model systems for studying cell size and polarity since they elongate
38 several hundred-fold their original size. Their tip growth is determined both by intrinsic and
39 environmental signals. Although nutrient availability and temperature are key factors for a
40 sustained plant growth, the molecular mechanisms underlying their sensing and downstream
41 signaling pathways remain unclear. Here, we identified that low temperature (10°C) triggers a
42 strong RH elongation response involving the cell surface receptor kinase FERONIA (FER) and the
43 nutrient sensing TOR Complex 1 (TORC1). In this study, we found that FER is required to perceive
44 limited nutrient availability caused by low temperature. FER interacts with and activates TORC1-
45 downstream components to trigger RH growth. In addition, the small GTPase Rho-related protein
46 from plants 2 (ROP2) is also involved in this RH growth response linking FER and TORC1. We also
47 found that limited nitrogen nutrient availability can mimic the RH growth response at 10°C in a
48 NRT1.1-dependent manner. These results uncover a molecular mechanism by which a central
49 hub composed by FER-ROP2-TORC1 is involved in the control of RH elongation under low
50 temperature and nitrogen deficiency.

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61 kinase.

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62 Introduction

63
64 Root hairs (RH) are cell outgrowths that develop as cylindrical protrusions from the root
65 epidermis in a developmentally regulated manner¹. RHs are able to expand in a polar manner
66 several hundred times their original size in a couple of hours to reach water-soluble nutrients in
67 the soil, to promote interactions with the local microbiome, and to support the anchoring of the
68 plant². RH growth is controlled by the coordination of a plethora of environmental and
69 endogenous factors^{3,4}. Recently, an autocrine mechanism of RH growth was described where
70 RALF1-FER complex recruits and phosphorylates the early translation initiation factor 4E1
71 (eIF4E1) to enhance protein synthesis of specific mRNAs, including the RH growth master
72 regulator *ROOT HAIR DEFECTIVE SIX-LIKE4 (RSL4)*⁵. The observation that RH cells can respond to
73 their local environment in a cell-autonomous manner within minutes⁶, points towards
74 mechanisms that directly modulate the growth machinery at the growing apex.

75
76 It is known that macronutrient availability is a key factor that promotes rapid RH growth^{3,7,8}
77 Recently, we showed that under low temperature conditions (10°C), nutrient availability in the
78 media is reduced and triggers an enhanced RH growth that is suppressed if nutrients are
79 increased⁹. We specifically showed that RH growth of WT Col-0 plants is highly responsive to
80 increasing nutrient concentrations (from 0.5X Murashige and Skoog (MS) to 2.0X MS) with regular
81 concentration gelling agents (0.8%). High nutrient concentration impairs RH growth even if they
82 are exposed to low temperatures. In a similar manner, an increase in agar concentration (from
83 0.8% to 2.5%) in the MS medium, which likely restrains nutrient mobility and nutrient uptake¹⁰⁻¹³
84 blocked low temperature-induced RH elongation⁹. These observations suggested that low
85 temperature restrict nutrient mobility and availability in the culture medium, leading to
86 promotion of polar RH growth. It is still unclear which specific nutrients are affected and the
87 signaling pathways involved in the temperature effect that triggers this RH growth response.
88 Diminished nutrient availability is known to activate RH expansion through a transcriptional
89 reprogramming governed by the transcription factors (TF) RHD6-RSL4⁹. Specifically, a novel
90 ribonucleoprotein complex composed of lncRNA AUXIN-REGULATED PROMOTER LOOP (APOLO)
91 and the TF WRKY42 forms a regulatory hub to activate RHD6 by shaping its epigenetic
92 environment and integrate low temperature signals governing RH growth^{9,14}. In addition, we
93 recently identified new molecular components involved in low temperature RH growth. We
94 found that cell wall-apoplastic related peroxidases, PRX62 and PRX69, are important for low
95 temperature triggered RH growth by inducing changes in ROS homeostasis and cell wall EXTENSIN
96 insolubilization¹⁵. Although relevant advances have been achieved in our understanding on how
97 RH growth occurs at low temperature, it is still unclear how nutrient availability caused by low
98 temperature is perceived in the RH cells and the extent of molecular responses that control RH
99 growth.

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100
101 TARGET OF RAPAMYCIN (TOR) is an evolutionarily conserved Ser/Thr protein kinase in all
102 eukaryotic organisms that acts as a central growth regulator controlling metabolism and protein
103 synthesis^{16–18}. TOR is found in at least two distinct multiprotein complexes called TOR Complex 1
104 and 2 (TORC1 and TORC2) in animal cells, although only TORC1 has been experimentally validated
105 in plants^{19,20}. The *Arabidopsis* TORC1 complex is encoded by one *TOR* gene (*AtTOR*)²¹, two
106 *REGULATORY-ASSOCIATED PROTEIN OF TOR (RAPTOR 1A and 1B)* genes^{22–27}, and two *LETHAL*
107 *WITH SEC THIRTEEN 8 (LST8)* genes²⁸. In contrast to the embryo-lethal *tor*-null mutant lines,
108 *raptor1b* and *lst8-1* mutants are viable but show significant growth defects and developmental
109 phenotypes²⁹. Some canonical downstream targets of TOR are conserved in plants, such as the
110 40S ribosomal S6 kinase (S6K) which stimulates protein synthesis^{24,30–33}. In plants, TORC1 complex
111 plays a key role during many stages of the plant life cycle by controlling both anabolic and
112 catabolic downstream processes. In addition, TORC1 is activated by nutrient availability and
113 inactivated by stresses that alter cellular homeostasis^{18,34–38}. The TORC1 complex senses and
114 integrates signals from the environment to coordinate developmental and metabolic processes
115 including hormones (e.g., auxin), several nutrients (e.g., nitrogen and sulfur), amino acids and
116 glucose^{16,27,39–43}. However, the underlying molecular mechanism by which TORC1 operates at a
117 single plant cell level has yet to be elucidated.

118
119 To date, few upstream regulators of TOR have been described in plants. Among them, there is a
120 subfamily of small GTPase Rho-related protein from plants (ROPs) involved in the spatial control
121 of cellular processes by signaling to the cytoskeleton and vesicular trafficking. Particularly, ROP2
122 activates TOR in response to auxin and nitrogen signals^{43–45}. ROP2 is described as a monomeric
123 GTP-binding protein that participates in many cellular signaling processes, including the polar
124 growth of pollen tubes and root hairs^{46–49}. ROP activation is regulated by ROP guanine nucleotide
125 exchange factors (ROP-GEFs) which interact with several receptor-like kinases (RLKs) including
126 the *CrRLK* family member *FERONIA (FER)*⁴⁷. During the elongation of RH, ROP2 is activated by the
127 ROPGEF1-FER interaction. This process is also regulated by ROP-GEF4 and ROP-GEF10^{47,50}. FER
128 was also linked to carbon/nitrogen balance during plant growth⁵¹. Even more, it was shown that
129 the cytoplasmic kinase domain of FER and its partner RIPK (RPM1-induced protein kinase)
130 interact with TOR and RAPTOR-1B forming a complex to positively modulate the TOR pathway
131 under low nitrogen nutrient conditions⁵². Currently, it is clear that plant growth can be regulated
132 via the FER-TOR pathway as it links amino acid and/or nutrients signaling with true leaves
133 development⁵¹. This evidence suggests that nutrient-mediated sensing at the cell surface level
134 triggers a response using the TOR signaling pathway. However, it is still unknown which are the
135 environmental signals activated in RH by the FER-TORC1 pathway in RH. Considering that low
136 temperature stress can result in enhanced RH growth^{9,14}, this stress condition was used as a
137 proxy to investigate environmental signals that activate RH cell elongation. Our research reveals a

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138 novel mechanism in which FER, ROP2 and TORC1 are necessary to regulate RH growth in
139 *Arabidopsis* in the context of the nitrogen nutrient availability as caused by low temperature
140 stress.

141

142 Results

143

144 **FER is required to trigger a strong RH growth response at low temperature.** We asked which
145 might be the RH surface protein involved in perceiving or transducing the low temperature
146 stimulus. Since CrRLK1L FER was shown as an important hub between the cell surface signaling
147 and downstream processes during RH growth^{5,47}, we tested whether *fer-4* null mutant or *fer-5* (a
148 truncated version with shortened kinase domain (KD)) can respond to low temperature stimulus
149 (**Figure S1**). Both *fer* mutants failed to trigger RH growth at low temperature (**Figure 1A**). In
150 contrast, the mutant *eru*, impaired in the related CrRLK1L ERULUS (ERU), previously linked to RH
151 growth and cell wall integrity processes with very short RH phenotype at 22°C^{53,54}, was able to
152 react to low temperature which triggered RH growth response although to a lower extent than
153 wild-type Col-0 plants (**Figure 1A**). This indicates that FER, but not ERU, might be involved
154 specifically in this growth response despite being phylogenetically related⁵⁴. Then, a fluorescent
155 translational reporter line of FER (*pFER:FER-GFP*), was used to study *FER* expression in roots and
156 RH. FER protein levels are increased clearly activated in root cells under low temperature
157 stimulus (**Figure 1B**), and during RH growth (**Figure 1C**). Distribution of FER-GFP in the clear zone
158 and punctate compartments in the cytoplasm was not affected by the temperature change from
159 22°C to 10°C. However, accumulation of FER-GFP at the plasma membrane was increased at low
160 temperature (at 10°C) (**Figure 1C**). Finally, phosphorylated, and non-phosphorylated levels of FER
161 (FER-p/FER) were quantified at both temperatures. Under low temperature, almost half of the
162 FER protein levels were present as a non-phosphorylated form and after 3 days of growth, almost
163 all FER protein was present as a phosphorylated version (**Figure 1D**). This indicates that low
164 temperature not only increases FER protein levels at the plasma membrane of growing RH, but
165 also promotes a complete phosphorylation of FER protein. This is possibly enhancing the
166 interaction with putative partners and triggering the activation of downstream signaling
167 components of RH polar growth.

168

169 **TORC1 pathway is involved in low temperature induced RH growth.** Recently, it was shown that
170 FER-RIPK interacts with TOR-RAPTOR1B and phosphorylates it leading to TOR pathway activation
171 in the context of nutrient perception and regulation of global metabolism⁵². We then asked
172 whether TOR might be also involved in the RH growth process under low temperature. We used a
173 β -estradiol (es)-induced TOR knockdown (*tor-es*, es-induced RNAi silencing of TOR) line⁵⁵ which
174 showed short RH as reported previously³². Notably, induction of TOR silencing with estradiol
175 completely blocked the RH response to low temperature (**Figure 2A**). Similarly, inhibition of TOR

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176 kinase activity with AZD-8055⁵⁶, an ATP-competitive inhibitor of mTOR kinase activity, abolished
177 the RH growth at both 22°C and low temperature (**Figure 2B**). Overexpression of TOR enhanced
178 RH growth at both temperatures assayed (**Figure 2C**). These results suggest that the TOR pathway
179 is involved in the RH growth and specifically in RH low temperature growth response and might
180 operate downstream of FER. At the transcriptional level, TOR root expression level is upregulated
181 up to three-folds at 10°C vs 22°C (**Figure S2**). We tested whether the growth response at low
182 temperature in RH is also affected in mutants of the TORC1 pathway and some downstream
183 components. All the plant mutants tested (*raptor1b*, *rps6b*, and *lst8-1*) were unable to respond to
184 the low temperature treatment. They exhibited a broad spectrum of RH phenotypes as compared
185 to Wt Col-0 plants (**Figure 2C**). For instance, *raptor1b* behaved similarly to *tor-es* mutant, while
186 *rps6b* showed an intermediate phenotype and *lst8* was comparable to Wt Col-0 plants (at 22°C)
187 (**Figure 2C**). In addition, overexpression of TOR and the S6 KINASE 1 (S6K1), a direct downstream
188 target of TOR, showed enhanced RH growth at 22°C. Since previous reports showed
189 phosphorylation of the S6K1 can be used to monitor TOR protein kinase activity in plants^{32,55}, we
190 measured ratios of S6K-p/S6K. Strong induction of the S6K-p/S6K ratio was observed in Wt Col-0
191 under low temperature (**Figure 2D**). These results suggest TORC1 and some downstream
192 components (e.g. S6K and RPS6b) are required to promote RH growth under low temperature.

193
194 **FER facilitates TOR polar localization in RH and activation of TORC1 pathway at low**
195 **temperature.** Since TORC1 activation under a plethora of stimuli usually triggers phosphorylation
196 of the downstream factor S6K (S6K-p), we asked whether low temperature responses regulated
197 by FER might control this output. We investigated the molecular mechanisms by which TORC1-
198 S6K are induced at 10°C and we tested whether this might be mediated by FER. Since *fer-4* and
199 *tor-es* showed similar RH phenotypes at low temperature (**Figure 2A-B**), possibly indicating that
200 they may act in the same pathway, we first tested whether FER and TOR interaction is enhanced
201 under 10°C vs 22°C. By performing a co-immunoprecipitation (Co-IP) analysis, we found that FER-
202 TOR interaction is enhanced under low temperature (**Figure 3A**). This result suggests an
203 interaction between FER and TOR kinase is implicated in RH growth under low temperature. This
204 corroborates previous findings that showed direct interaction of the FER kinase domain and the
205 N-terminal domain of TOR (aa 1-1449 of TOR, NTOR)⁵². Next, we evaluated TOR levels in growing
206 RH by immunolocalization. We found that FER is required for the apical accumulation of TOR
207 since TOR lost the polar pattern in *fer-4* root hairs. However, the level of RH tip-localized TOR
208 detected was similar at both, low temperature, and control conditions (22°C) (**Figure 3B-C**). This
209 result indicates that localization of TOR at RH tip is dependent on FER. Then, we tested whether
210 FER is involved in S6K activation by TOR. We measured S6K-p/S6K protein ratio under both
211 growth conditions in Wt Col-0, *fer-4*, *fer-5*, and FER^{K565R} (which has been shown to abolish FER
212 autophosphorylation and transphosphorylation in an *in vitro* kinase activity assay^{57,58}, although
213 some activity might remain⁵⁹). Interestingly, low temperature enhanced S6K-p levels after 3 days

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214 of growth in Wt Col-0 this was partially suppressed in all three *fer* mutants tested (**Figure 3D**).
215 This suggests that FER controls the level of TORC1 activation under low temperature in RH.
216 Collectively, these results indicate that TOR localization in the RH tip and TORC1 activation is
217 dependent on FER and enhanced at low temperature.

218
219 **Nitrate perception and transport mediated by NRT1.1 controls RH growth at low temperature.**
220 Nitrate is a key nitrogen nutrient required for plant growth and development^{60–62}. The
221 CHLORINA1/NITRATE TRANSPORTER1.1 (CHL1/NRT1.1) is the only known nitrate receptor
222 (transporter and receptor)^{63–65} as well as for auxin transport⁶⁶. NRT1.1 belong to the low affinity
223 nitrate transporter family. However, when NRT1.1 is phosphorylated at threonine 101, it behaves
224 as a high-affinity NO₃⁻ (nitrate) transporter, giving this protein a dual-affinity capability^{67–70}.
225 Previous studies have shown that nitrogen and specifically nitrate is important for TOR
226 signaling^{43,71}. Since low temperature growth conditions reduce nutrient availability in the media
227 and trigger a strong response in RH growth^{9,14}, we hypothesized that the nitrate signaling
228 pathway was involved. We found that high levels of nitrate (18.8 and 37.6 mM) supplied in the
229 M407 media without nitrogen (see methods) were able to partially repress low temperature-
230 mediated RH growth while low levels of nitrate (0.5 mM) did not affect growth under this
231 temperature (**Figure 4A**). As a reference, previous experiments were carried out with 0.5X MS
232 media containing 9.3 mM of nitrate and several other compounds (see **Table S1**). This result
233 indicated nitrate can impact RH cell elongation at low temperature, particularly under low
234 nutrient mobility environment. Next, we tested whether NRT1.1 was involved in this low
235 temperature RH growth response. The NRT1.1 null mutants *chl1-5*, *chl1.9* (NRT1.1 harbors a
236 substitution P492L that suppresses its root nitrate uptake activity⁸), as well as the *CHL1*^{T101D}
237 (which mimics phosphorylated version of NRT1.1) showed strong RH growth response regardless
238 of temperature conditions (**Figure 4B**), similarly to *S6K1 OE* and *TOR OE* lines (**Figure 2C**), but to a
239 lower extent in terms of RH elongation. Only the *CHL1*^{T101A} (dephosphorylated version of NRT1.1)
240 was able to slightly respond to the change in temperature. As expected, similar levels of S6K-
241 p/S6K were detected in the *chl1-9* line at both temperatures (**Figure 4C**). When these NRT1.1
242 mutants were grown under high (18.8 mM) or low (0.5mM) nitrate concentrations (high N/low N)
243 at both temperatures (at 22°C and at 10°C), a similar RH phenotype was detected between both
244 nitrate conditions either at 22°C or 10°C (**Figure 4D**). In contrast, Wt Col-0 RH was sensitive to
245 both low temperature and nitrate levels. Taken together, these results suggest low temperature
246 may restrict nitrate accessibility to the RH linked to a lower mobility in the media affecting
247 NRT1.1-mediated signaling upstream of TORC1-S6K activation.

248
249 **ROP2 is required for low temperature RH growth.** Previous studies have shown a key role of
250 ROP2 in the regulation of RH polarity and elongation^{48,72,73}, as an important molecular link
251 between FER and downstream components involved in RH growth⁴⁷. In addition, ROP2 promotes

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252 the activation of TOR and its relocation to the cell periphery and induces the downstream signal
253 transduction pathway⁴⁴. More importantly, ROP2 was shown to integrate diverse nitrogen and
254 hormone signals for TOR activation⁴³. We asked whether low temperature (10°C) is able to
255 change ROP2 targeting (as ROP2p:ROP2-mCitrine) to the plasma membrane in roots and RH. Low
256 temperature (10°C) led to noticeable increase of ROP2 fluorescence intensity in root cells (**Figure**
257 **5A**), in the apical and subapical cytoplasm, and at the apical plasma membrane of RH (**Figure 5B**).
258 Then, we observed that *rop2* and *rop2 rop4* abolishes the differential growth responses at 10°C
259 and produces a very short RH phenotype (**Figure 5C**). On the contrary, ROP2 OE triggers a
260 constitutive RH growth at both temperatures (**Figure S3**) similarly to *S6K1 OE* and *TOR OE* lines
261 (**Figure 2C**). In addition, when ROP2 OE is expressed in the *fer* mutant background *fer-8* (ROP2
262 OE/*fer-8*; **Figure S3**), this strong growth effect disappears highlighting the role of FER on the ROP2
263 function. On the other hand, while high levels of TOR enhance RH growth regardless of the
264 temperature treatments, the constitutively active ROP2 version (CA-ROP2) is able to abolish TOR
265 activation of RH growth (**Figure 5C**). The levels of S6K-p/S6K in TOR OE, with constitutive long RH
266 at both temperatures, was reduced by the presence of CA-ROP2 (in TOR OE/CA-ROP2), which
267 correlates with a diminished RH growth (**Figure 5D**). Taken together, we suggest a direct function
268 of ROP2 (and possibly ROP4) at the RH tip. ROP2 function is dependent on the presence of FER
269 and it modulates TOR activity in the RH growth responses to low temperature.

270
271 **FER-ROP2-TORC1 hub is required for nitrate starvation-mediated RH growth response.** Recently
272 it was reported that RALF1-FERONIA complex is able to interact with TOR kinase under low N
273 conditions modulating the TORC1 downstream signaling pathway⁵². As described above, we
274 found that Col-0 grown at 22°C under high N condition have a shorter RH phenotype as compared
275 to low N condition. This growth response is consistent with previous studies in which the RH
276 length in *Arabidopsis* and other species decrease as the nitrate concentration increases although
277 the molecular mechanism remained unknown^{60,74}. To gain a better insight into the role of FER
278 and TOR in this process, we decided to test the *fer* mutants, the TORC1 pathway mutants and
279 selected overexpressing lines (TOR and S6K) in low (0.5 mM) and high nitrate (18.8 mM) media
280 conditions (**Figure 6A** and **Figure S4**). Under low nitrate conditions, the RH response was
281 suppressed in both *fer* mutants and in the *tor-es* inducible mutant compared to Col-0. This result
282 supports the idea that FER and TOR are necessary for the RH growth mediated by low nitrate
283 conditions. The *TOR OE* line showed a RH growth phenotype similar to Col-0 under low N and a
284 higher RH growth in excess of N. It is well known that nitrogen starvation causes enhanced root
285 and RH growth, whereas an excess of nitrate inhibits primary root growth⁷⁵, because of osmotic
286 stress⁷¹. Furthermore, plants overexpressing TOR in a nitrate excess medium have a longer
287 primary root phenotype⁷¹. According to our data that *TOR OE* line showed longer RH than Col-0
288 under normal conditions (**Figure 6A**). The constitutive overexpression of TOR can lead not only to
289 a longer primary root but also to longer RH when plants are grown on a high nitrate media to

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290 alleviate the nutrient stress. The constitutive active mutant of ROP2 suppressed the TOR OE RH
291 phenotype while *rop2* mutant was unable to respond to contrasting levels of nitrates. A model
292 was proposed where ROP GTPases and the cytoplasmic and apoplasmic pH fluctuations can
293 regulate RH tip growth in a nitrogen supply dependent manner thus maintaining the
294 unidirectional growth during the RH elongation⁷⁷. Additionally, nitrate and ammonium levels
295 restore TOR activation under nitrogen-starvation condition via ROP2 activation⁴³. We then tested
296 if other components on the TORC1 complex are also involved in the low nitrate perception linked
297 to RH growth (**Figure 6B**). Similarly to their low temperature phenotype, *raptor1b* and *lst8-1* do
298 not respond with differential RH growth under low nitrate. Mutation of RAPTOR1B resulted in a
299 strong reduction of TOR kinase activity, leading to massive changes in carbon and nitrogen
300 metabolism⁷⁸. The RH phenotype derived from the overexpression of S6K resembled that of TOR
301 OE line (**Figure 6A, 6B**). Altogether, our results show that low-nitrate (low N) RH responses are
302 similar to those determined for low temperature. This is consistent with the idea that nitrate is
303 one of the main signals that trigger RH expansion upon perception by NRT1.1, and subsequently
304 transduction by FER, ROP2 and TORC1 pathway. We then tested how changed are the levels of
305 S6K-p/S6K as a readout of TORC1 activation in low N and high N conditions in FER, TOR silenced
306 line (*tor-es*) and NRT1.1 mutants compared to Wt Col-0 (**Figure 6C**). First, Wt Col-0 showed an
307 increased in S6K-p/S6K ratio in low nitrogen similarly to the low temperature effect. On the
308 contrary, and as expected, *fer-4*, *fer-5* and *tor-es* showed much reduced levels of S6K-p/S6K in
309 both nitrate conditions but much lower in low nitrogen. Importantly, the effect of low N on
310 increasing S6K-p levels is abolished in NRT1.1 mutants (*chl1-9*, *chl1-5* and *CHLT101D*), which is
311 similar to the results observed under low temperature conditions. These results indicate that
312 nitrate effect on S6K-p/S6K ratio depends on NRT1.1 and the FER-TOR complex.

313
314 Since an integrative gene regulatory network analysis of TF–target interactions positioned TGA1
315 and its homolog TG4 as the most influential TFs in the nitrate response^{79,80}, we decide to test if
316 they play a role in the response to low temperature in RHs and low/high levels of Nitrogen
317 (**Figure S5**). As expected *tga1 tga4* double mutant failed to respond to low temperature and low
318 nitrogen while TGA1 OE showed a constitutive growth response regardless the temperature and
319 nitrogen levels. This confirms that nitrate, NRT1.1 and its downstream signaling pathway
320 including the transcriptional responses controlled by TGA1-TG4 act an important pathway in RH
321 growth at low temperature.

322

323 Discussion

324

325 Growth and development of plants and animals are based on nutrient and hormonal signaling
326 that constitute as the main regulatory networks in eukaryotes. Unraveling the functions of the
327 regulatory hubs and their detailed molecular mechanisms are critical for deeper understanding of

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328 these signaling pathways. In all cells, there is a central hub composed by the evolutionarily
329 conserved TOR protein kinase that integrates nutrient and energy levels coupled to stress
330 signaling networks to further modulate cell growth⁸¹⁻⁸⁴. In addition, the cell surface receptor FER
331 acts as a versatile sensor of signals coming from the environment^{51,85,86}. Specifically, our study
332 uncovers a new molecular mechanism by which plants use FER-ROP2-TORC1 signaling pathway to
333 control RH elongation under low nutrient conditions, specifically nitrate, induced by low
334 temperature stress (**Figure S6**).

335
336 In this work, we discovered that the cell surface FER receptor regulates TOR apical localization
337 and further downstream activation, both controlling S6K phosphorylation linked to RH growth.
338 Based on the results obtained and previous evidences, here we propose a model in which the
339 FER-ROP2-TORC1 axis might be able to regulate the RH growth under low temperature with
340 remarked attention to variable nitrate conditions (**Figure S6**). In agreement with this, it was
341 previously shown that ROP2 in response to auxin is an upstream effector of TOR activation⁴⁴ and
342 by inorganic and organic nitrogen inputs⁴³. In addition, FER was shown to directly activate the
343 TOR/RAPTOR1B signaling pathway under low inorganic and organic nitrogen conditions⁵² and is
344 an upstream regulator of the ROPGEF4-ROP2 signaling pathway that controls ROS-mediated RH
345 development^{47,87}. In addition, ROP2 promotes TOR accumulation close to the cell periphery⁴⁴,
346 suggesting that plasma membrane-localized FER together with ROP2 function in regulating the
347 relocation of TOR, close to the RH tip. Interestingly, such a local, non-genomic regulatory circuit
348 may also explain the rapid, cell-autonomous growth regulation in RH cells observed under
349 asymmetric conditions in the dual-flow-RootChip⁶. All these findings are in line with our results
350 and support the idea of the existence of the FER-ROP2-TORC1 pathway being highly relevant for
351 RH growth in a changing nutritional environment. Interestingly, we recently showed that FER
352 regulates localized protein synthesis during polar RH cell growth^{5,88}, suggesting that FER-TOR may
353 also regulates polar protein synthesis in RHs. All the data together clearly highlight a key role of
354 the FER-ROP2-TORC1 pathway as a core complex in sensing nutrients (e.g. nitrates) to direct RH
355 growth.

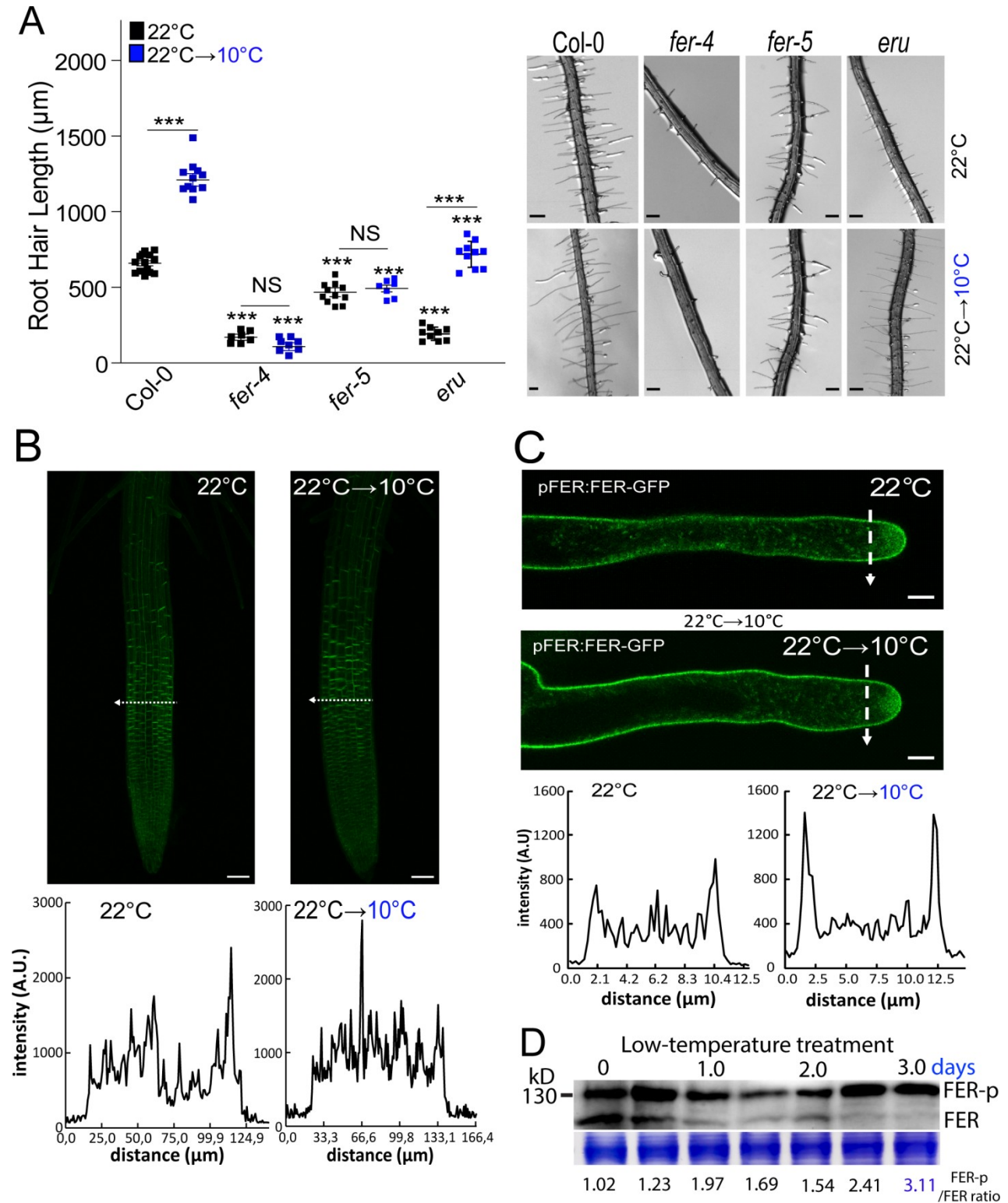
356
357 Nitrogen is an essential nutrient for plants that when scarce limits plant growth. Although
358 inorganic nitrogen (e.g. nitrate) is a major source of nitrogen for plants in soil, organic forms of
359 nitrogen, like amino acids, can also be assimilated by plants from soil^{89,90}. Changes in nitrate
360 concentrations in the media mimicked this response at low temperature and NRT1.1 mutants are
361 insensitive at RH growth level regardless the temperature or nitrate levels. We hypothesize that
362 low nitrate in the media might rapidly induce the levels of mature-active RALF1 (and possibly
363 other RALF peptides), which might then activate the FER-ROP2-TORC1 pathway and the
364 downstream fast cell elongation effect to search for further nutrients sources. This autocrine
365 mechanism dependent on RALF1-FER growth activation has been demonstrated for RH⁸⁸, but not

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366 in a context that involves low temperature/low nitrate and TORC1 pathway. It is expected that
367 low nitrates levels will impact on amino acid homeostasis. In line with our results, branched-chain
368 amino acids have been shown to serve as upstream activators of TOR in plants⁴⁰. TOR is also
369 involved in mediating amino acid-derived metabolic regulatory signals that influence respiratory
370 activity and plant metabolic rate⁹¹. Specifically, a total of 15 proteinogenic amino acids can
371 reactivate TOR under inorganic-N starvation conditions⁴³. In addition, the FER-TOR pathway
372 responds to Gln, Asp, and Gly, reinforcing that amino acids serve as conserved upstream
373 regulators of the TOR signaling pathway in animals and plants⁵². It is tempting to speculate that
374 under low nitrates condition (and low temperature), the RH cell compensates with an altered
375 amino acid homeostasis to trigger TOR activation linked fast cell growth.

376
377 Finally, several questions in our proposed model remain to be answered in future studies. How
378 does low nitrate concentration (low temperature) triggers FER-TORC1 activation?. Does RALF1
379 and other RALFs respond to low nitrate to activate FER?. Does FER activate NRT1.1 directly?.
380 Recently, it was shown that FER-regulated ROP2 triggers a new mechanism of negative regulation
381 in the case of rhizosphere microorganisms such as *Pseudomonas*⁷⁶. Although not tested here,
382 FER-ROP2-TORC1 signaling coupled to enhanced RH growth under low temperature (low-
383 nutrients) might be linked to the root growth in specific soil conditions to select favorable
384 microbiota in the soil. This pathway may integrate complex signals from soil nutrients and
385 microbiota in the rhizosphere to plant root cell growth mechanism.

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386

387 **Figure 1. High levels of FER in its phosphorylated form are required to trigger the low**
 388 **temperature RH growth.**

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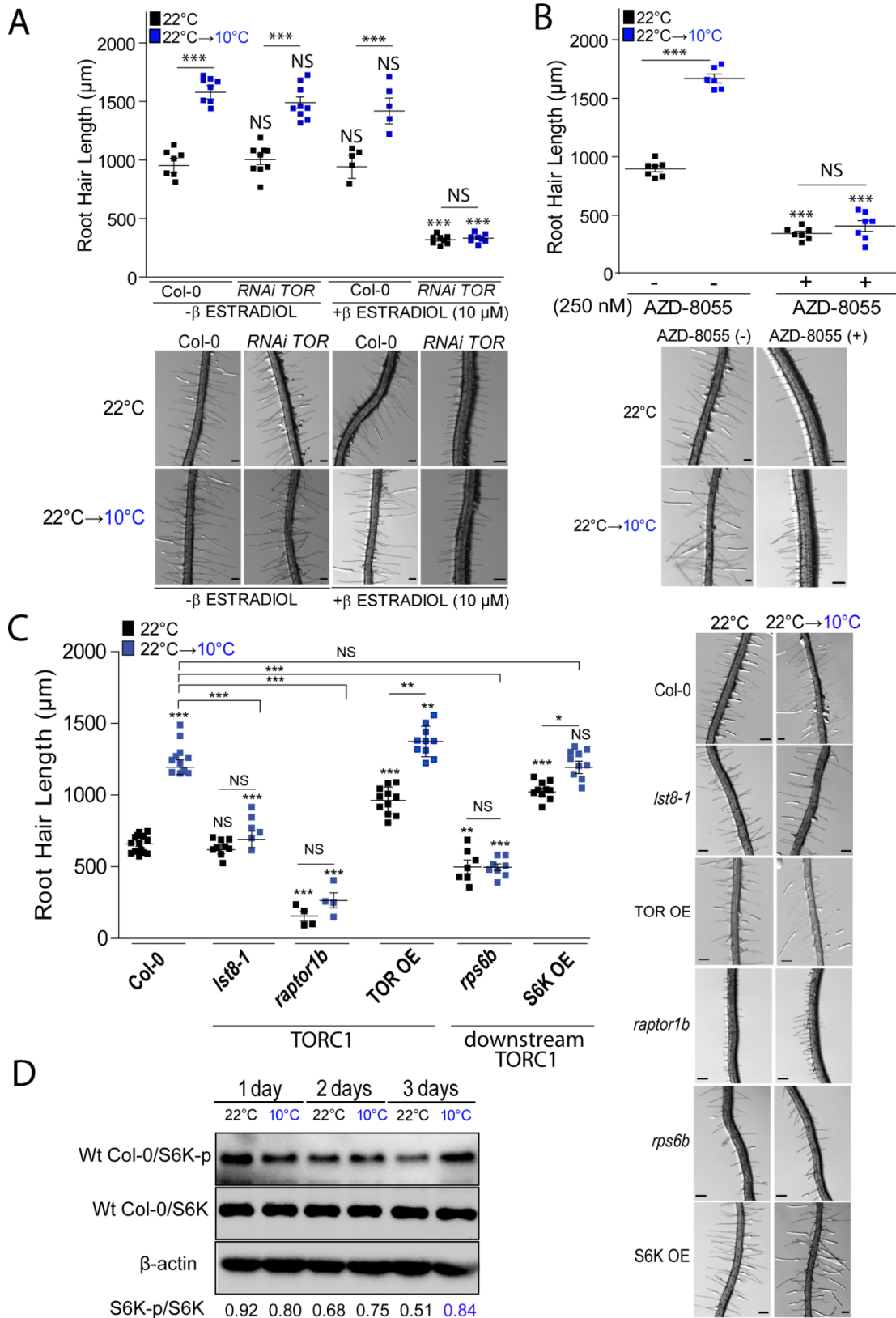
389 **(A)** Scatterplot of RH length of Col-0, of *fer-4*, *fer-5*, and *eru* mutants grown at 22°C or at 10°C. RH
390 growth is enhanced at low temperature in Wt Col-0 and *eru* mutant but not in the *fer-4* and *fer-5*
391 mutants. Each point is the mean of the length of the 10 longest RHs identified in the maturation
392 zone of a single root. Data are the mean \pm SD (N=10 roots), two-way ANOVA followed by a
393 Tukey–Kramer test; (***) $p < 0.001$, NS= non-significant. Results are representative of three
394 independent experiments. Asterisks indicate significant differences between Col-0 and the
395 corresponding genotype at the same temperature or between the same genotype at different
396 temperatures. Representative images of each genotype are shown on the right. Scale bars= 300
397 μm .

398 **(B-C)** Increased level of FER in root cells after its activation by low temperature. **(B)** Confocal
399 images of root apex of a fluorescent translational reporter line of FER (*pFER:FER-GFP*) at 22°C
400 and after transfer from ambient to low temperature (22°C→10°C). On the Bottom: semi-
401 quantitative evaluation of the GFP fluorescence intensity across the root tip at 22°C and after
402 transfer from ambient to low temperature at the transition zone, indicated by a interrupted
403 white line. Fluorescence intensity is expressed in arbitrary units (AU), N=4-6 roots. Results are
404 representative of three independent experiments. Scale bars: 50 μm .

405 **(C)** Confocal images of short root hairs of a fluorescent translational reporter line of FER
406 (*pFER:FER-GFP*) at 22°C and after transfer from ambient to low temperature (22°C→10°C) On the
407 bottom: semi-quantitative evaluation of the GFP fluorescence intensity across the subapical root
408 hair zone at 22°C and after transfer from ambient to low temperature, indicated by a interrupted
409 white line. Fluorescence intensity is expressed in arbitrary units (A.U.), N=6-8 roots, 8-11 root
410 hairs. Results are representative of three independent experiments. Scale bars=5 μm .

411 **(D)** Phosphorylation levels on FER (FER-p in *pFER:FER-GFP*) increases after 3 days at low
412 temperature in roots. Protein loading control (Coomasie Blue) is indicated below. FER-p/FER
413 ratios were analyzed by ImageJ.

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Figure 2. TORC1 signaling pathway is required for low temperature triggered RH growth.

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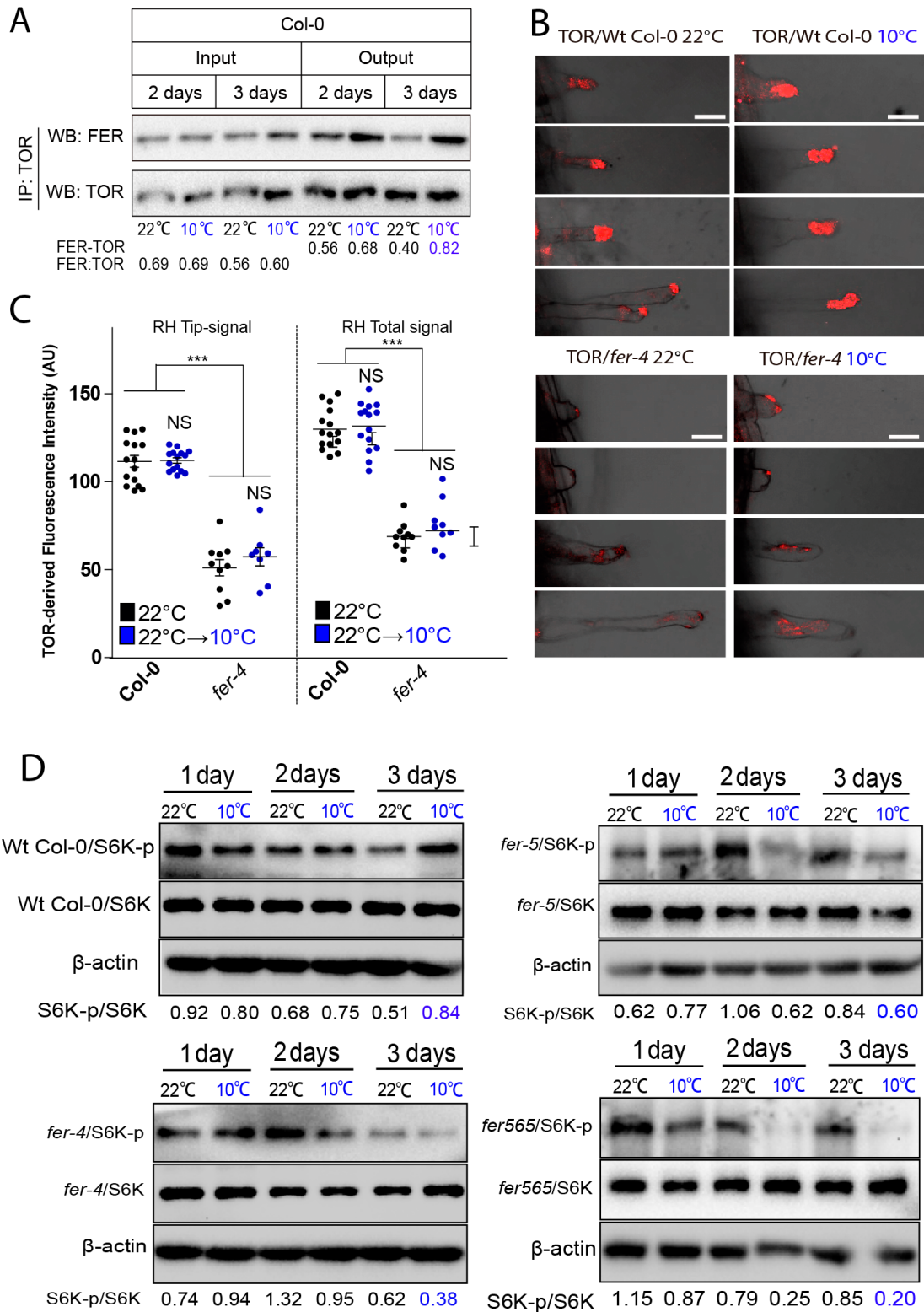
417 **(A)** Scatterplot of RH length of Col-0 and *tor-es* line grown at 22°C or at 10°C. Differential growth
418 of RH at low temperature is suppressed in the estradiol inducible *RNAi TOR* line. Each point is the
419 mean of the length of the 10 longest RHs identified in the maturation zone of a single root. Data
420 are the mean \pm SD (N=7-10 roots), two-way ANOVA followed by a Tukey–Kramer test; (***)
421 $p < 0.001$, NS=non-significant. Results are representative of three independent experiments.
422 Asterisks indicate significant differences between Col-0 and the corresponding genotype at the
423 same temperature or between the same genotype at different temperatures. Representative
424 images of each line are shown below. Scale bars=300 μ m.

425 **(B)** Differential growth of RH at low temperature is abolished in the Col-0 treated with 250 nM of
426 TOR inhibitor, AZD-8055. Each point is the mean of the length of the 10 longest RHs identified in
427 the maturation zone of a single root. Data are the mean \pm SD (N=7 roots), two-way ANOVA
428 followed by a Tukey–Kramer test; (***) $p < 0.001$, NS=non-significant. Results are representative
429 of three independent experiments. Asterisks indicate significant differences between Col-0 and
430 the corresponding genotype at the same temperature or between the same genotype at different
431 temperatures. Representative images of each line are shown below. Scale bars=300 μ m.

432 **(C)** RH elongation of TORC1 and downstream TORC1 signaling pathway mutants under low
433 temperature. Each point is the mean of the length of the 10 longest RHs identified in the
434 maturation zone of a single root. Data are the mean \pm SD (N=7-12 roots), two-way ANOVA
435 followed by a Tukey–Kramer test; (*) $p < 0.05$, (**) $p < 0.01$, (***) $p < 0.001$, NS=non-significant.
436 Results are representative of three independent experiments. Asterisks indicate significant
437 differences between Col-0 and the corresponding genotype at the same temperature or between
438 the same genotype at different temperatures. Representative images of each line are shown on
439 the right. Scale bars=300 μ m.

440 **(D)** Analysis of the phosphorylation state of S6K (S6K-p/S6K ratio) in Col-0. A representative
441 immunoblot is shown of a three biological replicates. S6K-p/S6K ratio was analyzed by ImageJ.

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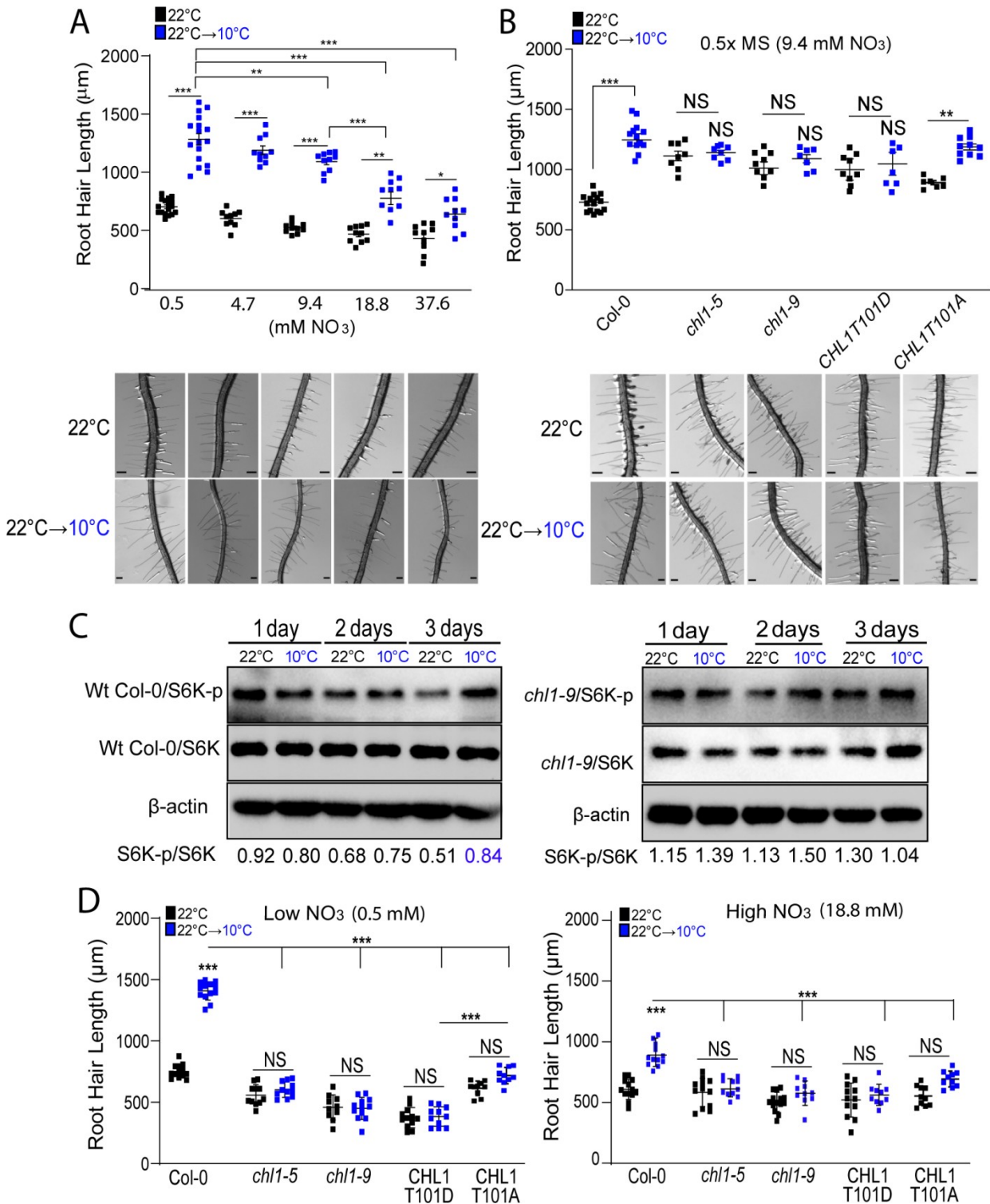
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Figure 3. FER controls polar TOR localization and subsequent activation thus increasing phosphorylation of S6K under low temperature.

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- 446 **(A)** Enhanced FER-TOR interaction at low temperature in roots by Immunoprecipitation (IP). A
447 representative experiment of three replicates is shown.
- 448 **(B)** Representative images showing TOR immunolocalization in RHs is FER-dependent. Scale
449 bar=10 μ m.
- 450 **(C)** Apical and total TOR signal quantification in RHs showed in (B). A ROI at the RH tip of the
451 fluorescent signal or at the entire RH TOR-derived fluorescent signal was measured. Fluorescence
452 AU data are the mean \pm SD (N=10-15 root hairs), two-way ANOVA followed by a Tukey–Kramer
453 test; (***) $p < 0.001$, NS= non-significant. Results are representative of two independent
454 experiments. Asterisks on the graph indicate significant differences between genotypes.
- 455 **(D)** Analysis of the phosphorylation state of S6K in Wt Col-0, *fer-4*, *fer-5*, and *FER*^{K565R} mutants
456 Arabidopsis roots. Phosphorylation of S6K (S6K-p) is enhanced at low temperature and requires
457 FER active kinase. A representative immunoblot is shown of a three biological replicates. S6K-
458 p/S6K ratio was analyzed by Image J. Wt Col-0 immunoblot is the same of the Figure 2D.

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461 **Figure 4. Nitrate acts as a RH growth signal perceived by NRT1.1 and triggered by TORC1**
462 **pathway at low temperature growth.**

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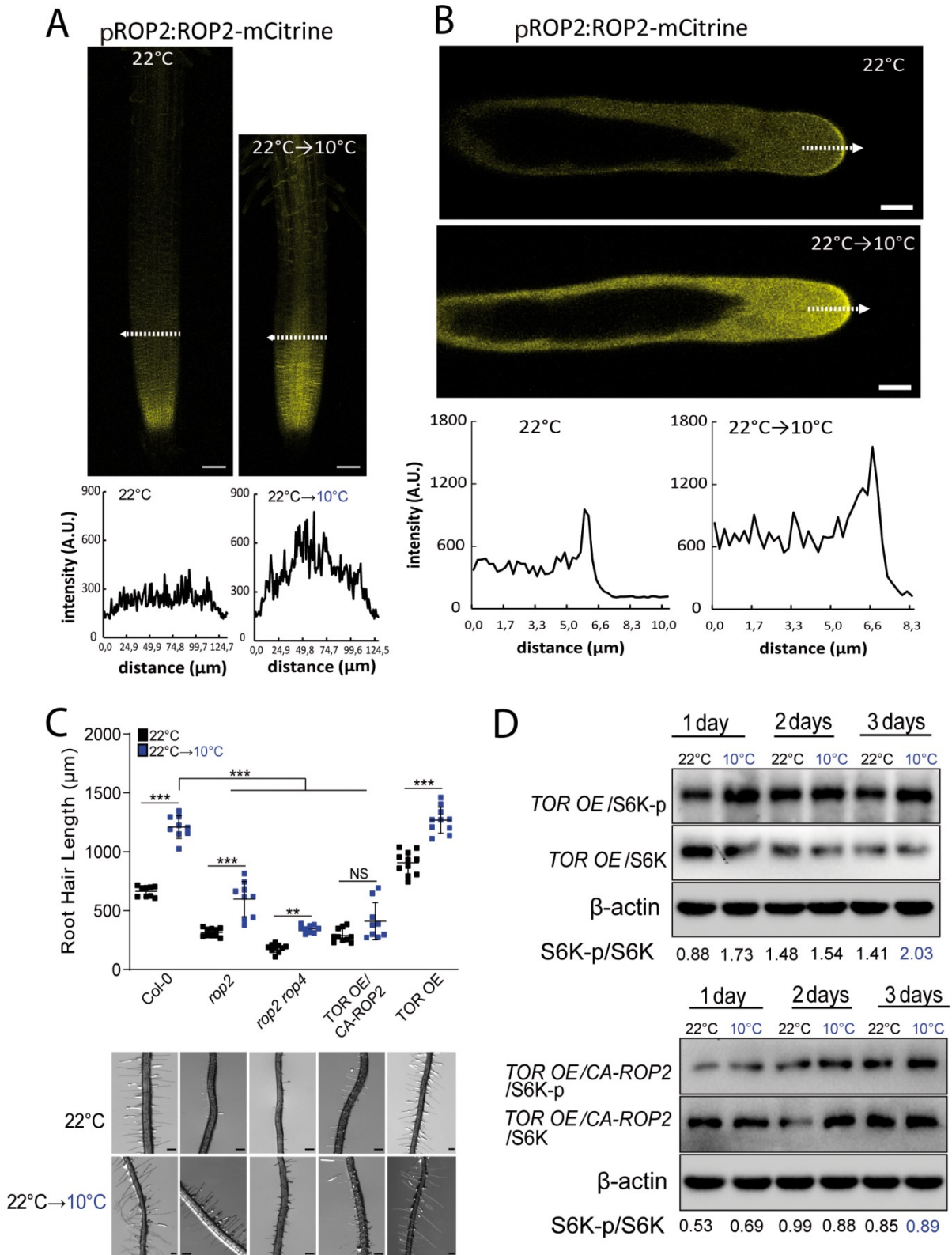
463 **(A)** High nitrate supplemented in M407 media (18.8 and 37.6 mM) partially suppresses low
464 temperature enhanced RH growth in Wt Col-0 plants. Each point is the mean of the length of the
465 10 longest RHs identified in the maturation zone of a single root. Data are the mean \pm SD (N=10
466 roots), two-way ANOVA followed by a Tukey–Kramer test; (***) $p < 0.001$. Results are
467 representative of three independent experiments. Asterisks indicate significant differences
468 between 0.5mM N concentration and every other N concentration at the same temperature or
469 between the same N concentrations at different temperatures. Representative images of Col-0
470 under the different nitrate concentrations are shown below. Scale bars=300 μ m.

471 **(B)** Low temperature RH growth is regulated by the nitrate sensor NRT1.1 (CHL1). Scatter-plot of
472 RH length of Col-0 and NRT1.1 mutants in 0.5X MS (contains 9.4 mM Nitrates) grown at 22°C or
473 10°C. Each point is the mean of the length of the 10 longest RHs identified in the maturation zone
474 of a single root. Data are the mean \pm SD (N=10-15 roots), two-way ANOVA followed by a Tukey–
475 Kramer test; (***) $p < 0.001$, NS=non-significant. Results are representative of three independent
476 experiments. Asterisks indicate significant differences between Col-0 and the corresponding
477 genotype at the same temperature or between the same genotype at different temperatures.
478 Representative images of each line are shown below. Scale bars=300 μ m.

479 **(C)** Analysis of the phosphorylation state of S6K in Wt Col-0 and *chl1-9* mutant roots. A
480 representative immunoblot is shown of a three biological replicates. S6K-p/S6K ratio was
481 analyzed by ImageJ. Wt Col-0 immunoblot is the same of the Figure 2D and Figure3D.

482 **(D)** RH growth responses under high and low nitrate (18.8 mM and 0.5 mM, respectively) in Wt
483 Col-0, *chl1-5*, *chl1-9* mutants, and *CHL1*^{T101D} grown at 22°C or 10°C. Each point is the mean of the
484 length of the 10 longest RHs identified in the maturation zone of a single root. Data are the mean
485 \pm SD (N=10-15 roots), two-way ANOVA followed by a Tukey–Kramer test; (***) $p < 0.001$, NS= non-
486 significant. Results are representative of three independent experiments. Asterisks indicate
487 significant differences between Col-0 and the corresponding genotype at the same temperature
488 or between the same genotype at different temperatures.

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Figure 5. ROP2 is required to triggers RH growth at low temperature.

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492 **(A-B)** Increased level of ROP2 after its activation by low temperature.

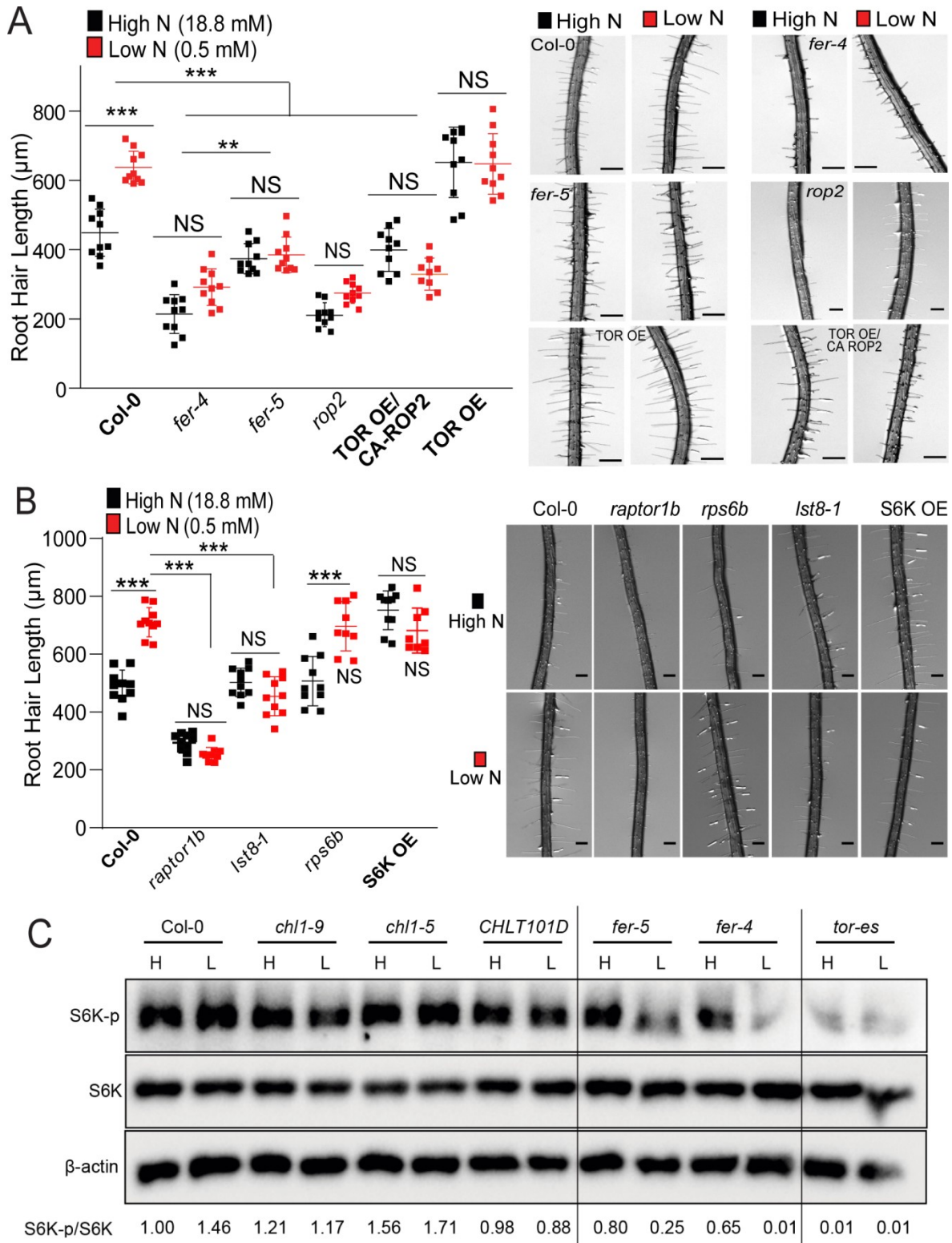
493 **(A)** Confocal images of root apex of a fluorescent translational reporter line of ROP2
494 (*pROP2:ROP2-mCitrine*) at 22°C (A) and after transfer from ambient to low temperature (22°C→
495 10°C). On the bottom: semi-quantitative evaluation of the mCitrine fluorescence intensity across
496 the root tip at 22°C and after transfer from ambient to low temperature at the transition zone,
497 indicated by a interrupted white line. Fluorescence intensity is expressed in arbitrary units (A.U.),
498 N=4-5 roots. Results are representative of three independent experiments. Scale bars=50 µm.

499 **(B)** Increased level of ROP2 in the plasma membrane of RHs after its activation by low
500 temperature. Confocal images of short RHs of a fluorescent translational reporter line of ROP2
501 (*ROP2p:ROP2-mCitrine*) at 22°C and after transfer from ambient to low temperature (22°C→10°
502 C). On the bottom: semi-quantitative evaluation of the mCitrine fluorescence intensity across the
503 RH apex at 22°C and after transfer from ambient to low temperature, indicated by a interrupted
504 white line. Fluorescence intensity is expressed in arbitrary units (A.U.), N=4-5 roots, 6-8 root
505 hairs. Results are representative of three independent experiments. Scale bars=5 µm.

506 **(C)** ROP2 is required for RH at low temperature meanwhile *rop2*, *rop2 rop4* and *CA-ROP2/TOR OE*
507 blocks RH growth. Each point is the mean of the length of the 10 longest RHs identified in a single
508 root. Data are the mean ± SD (N=10-15 roots), two-way ANOVA followed by a Tukey–Kramer test;
509 (**) $p < 0.01$, (***) $p < 0.001$. Results are representative of three independent experiments.
510 Asterisks indicate significant differences. Representative images of each line are shown below.
511 Scale bars=300 µm.

512 **(D)** Analysis of the phosphorylation state of S6K in *TOR OE* and *TOR OE /CA-ROP2* Arabidopsis
513 roots. Phosphorylation of S6K (S6K-p) is enhanced in *TOR OE* and suppressed in *TOR OE/ROP2-CA*.
514 A representative immunoblot is shown of a three biological replicates. S6K-p/S6K ratio was
515 analyzed by Image J.

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518 **Figure 6. Low nitrate perception relays in FER, ROP2 and TORC1 to trigger RH growth.**

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519 **(A)** Scatterplot of RH length of Col-0, *fer-4*, *fer-5*, *rop2*, TOR OE/CA-ROP2 and TOR OE grown in
520 low (0.5 mM) and high nitrate (18.8 mM) conditions in M407 media, both at 22°C. Each point is
521 the mean of the length of the 10 longest RHs identified in the maturation zone of a single root.
522 Data are the mean \pm SD (N=10 roots), two-way ANOVA followed by a Tukey–Kramer test; (**)
523 $p < 0.01$, (***) $p < 0.001$, NS=non-significant. Results are representative of three independent
524 experiments. Asterisks indicate significant differences between the same genotype at different N
525 concentration or between different genotypes at the same N concentration. Representative
526 images of each line are shown on the right. Scale bars=300 μ m.

527 **(B)** Scatterplot of RH length of Col-0, *raptor1b*, *lst8-1*, *rps6b* and S6K OE mutants grown in low
528 and high N conditions. Each point is the mean of the length of the 10 longest RHs identified in the
529 maturation zone of a single root. Data are the mean \pm SD (N=10 roots), two-way ANOVA followed
530 by a Tukey–Kramer test; (***) $p < 0.001$, NS=non-significant. Results are representative of three
531 independent experiments. Asterisks indicate significant differences between the same genotype
532 at different N concentration or between different genotypes at the same N concentration.

533 **(C)** Analysis of the phosphorylation state of S6K in Wt Col-0, in NRT1.1 mutants (*chl1-9*, *chl1-5*,
534 CHLT101D), and in FER mutants (*fer-4*, *fer-5*, and *FER^{K565R}*). A representative immunoblot is shown
535 of a three biological replicates. S6K-p/S6K ratio was analyzed by Image J.

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536 **Experimental Procedures**

537
538 **Plant Material and Growth Conditions.** *A. thaliana* Col-0 ecotype was used as a wild-type plant.
539 To test the low and high nitrate response a 0.5X M407 medium without N, P nor K (M407,
540 PhytoTechnology Laboratories, <https://www.phytotechlab.com/>) supplemented with 0.8% plant
541 agar (Duchefa, Netherlands); 1.17mM MES, 0.625 mM KH₂PO₄ monobasic and 0.5mM KNO₃ (low
542 nitrate medium) 9.4mM, 18.8 mM and 37.6 mM KNO₃ (high nitrate medium) was used. Media
543 composition is detailed in **Table S1**. Plants were grown in the above media in continuous light
544 (120 μmol.sec⁻¹.m⁻²) either: 8 days at 22°C in the case of *tor-es* line assay, 8 days at 22°C for
545 nitrate response assay or 5 days at 22°C + 3 days at 10°C for the low temperature and nitrate
546 response combination assay. For the rest of the low temperature experiments plants were grown
547 on regular 0.5X MS agar plates. For the imaging of fluorescence intensity distribution in root tips
548 and RH, seedlings were grown for 3 days at 22°C + 3-4 days at 10°C. All mutants and transgenic
549 lines are listed in **Table S2**. *chl1-5*, *chl1-9* and T101D mutants were kindly donated by Dr Yi-Fang
550 Tsay. *tor-es* was kindly donated by Dr Ezequiel Petrillo, *lst8-1*, *rps6b*, *raptor1b* and overexpressing
551 line S6K1 (S6K OE) kindly donated by Dr Elina Welchen and TOR OE and TOR OE/CA-ROP2 kindly
552 donated by Dr. Lyubov Ryabova.

553
554 **Pharmacological Treatments.** For all experiments, plants were grown first on solid 0.5X MS
555 medium at 22°C for 5 days in continuous light. According to the specific treatment plants were
556 transferred to plates containing regular solid 0.5X MS supplemented with 100nM IAA (auxin
557 treatment), 250 nM AZD-8055 (TOR inhibition) or 10 μM β-estradiol (*tor-es* line low temperature
558 treatment) and grown at 22°C for 5 days + 3 days at 10°C in continuous light. RH phenotype was
559 quantified after each 3 days span. For nitrate response treatment plants were transferred to
560 plates containing low or high nitrate solid media supplemented with 10 μM β-estradiol (*tor-es*
561 line), grown at 22°C for 3 days in continuous light and RH phenotype quantified.

562
563 **Root hair phenotype.** Seeds were surface sterilized and stratified in darkness for 3 days at 4°C.
564 Then grown *in vitro* on a specific condition and medium in a plant growth chamber in continuous
565 light (120 μmol.sec⁻¹.m⁻²) at 22°C and/or 10°C. The quantitative analyses of RH phenotypes of Col-
566 0 and transgenic lines were made the last day of the growth conditions described in the two
567 previous sections. In the case of low temperature treatment, the measurements were done after
568 5 days at 22°C and after 3 days at 10°C. For that purpose, 10 fully elongated RH from the
569 maturation zone were measured per root under the same conditions from each treatment and
570 control. Images were captured using an Olympus SZX7 Zoom Stereo Microscope (Olympus, Japan)
571 equipped with a Q-Colors digital camera and Q Capture Pro 7 software (Olympus, Japan). Results
572 were expressed as the mean ± SD using the GraphPad Prism 8.0.1 (USA) statistical analysis

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573 software. Results are representative of three independent experiments, each involving 7–20
574 roots.

575
576 **Confocal Microscopy.** For measurements of fluorescence intensity distributions after cold stress
577 (22°C→10°C) in root tips and root hairs of *pFER::FER-GFP* and *pROP2::ROP2-mCitrine* lines
578 confocal laser scanning microscopy Zeiss LSM 710 (Carl Zeiss, Germany) was used. For image
579 acquisition, 10x/0.3 NA EC Plan-Neofluar objective for root tips, and 40x/1.4 Oil DIC Plan-
580 Apochromat objective for root hairs were used. GFP signal was excited with 488 nm argon laser
581 at 4% laser power intensity and emission band of 493-549 nm. mCitrine signal was excited with
582 514 nm argon laser at 4% laser power intensity and emission band 519-583 nm. Fluorescence
583 intensity measurements (in A.U.) were generated using Zen Black 2011 software (Carl Zeiss,
584 Germany) and graphically edited in Microsoft Excel. Results are representative of three
585 independent experiments, each involving 1–2 roots and 1 to 4 hairs per root.

586
587 **Co-IP assay.** For Co-IP assays using A/G agarose and an anti-TOR antibody³², 30 µL of A/G beads
588 (Thermo Fisher Scientific Inc., 20421) was resuspended and washed three times using NEB buffer
589 (20 mM HEPES [pH 7.5], 40 mM KCl, 5 mM MgCl₂) before adding 8 µL of anti-TOR antibody or
590 preimmune serum as a negative control³² in a total volume of 500 µL of NEB buffer, followed by
591 incubation for 4 h at 4 °C. Col-0 seedlings were first grown at 22 °C for 5 days then transferred to
592 22 °C or 10 °C for 2 days or 3 days. For protein extraction from plants, the collected materials
593 were ground to a fine powder in liquid nitrogen and solubilized with NEB-T buffer (20 mM HEPES
594 [pH 7.5], 40 mM KCl, 5 mM MgCl₂, 0.5% Triton X-100) containing 1 × protease inhibitor cocktail
595 (Thermo Fisher Scientific Inc., 78430) and 1 × phosphatase inhibitor (Thermo Fisher Scientific Inc.,
596 78420) and incubated for 1 h on the ice. The extracts were centrifuged at 16,000 g at 4 °C for 15
597 min, and the resultant supernatant was incubated with prepared antibody-beads from the above
598 step. After overnight incubation at 4°C with rotation, the agarose beads were washed five times
599 with the NEB buffer and eluted with elution buffer (0.2 M glycine, 0.5% Triton X-100, pH 7.5).
600 Anti-FER and anti-TOR antibodies³⁰ were used for immunoblotting to detect the
601 immunoprecipitates.

602
603 **TOR immunolocalization.** Col-0 and *fer-4* seedlings were first grown at 22 °C for 5 days before
604 transferred to 22 °C or 10 °C for 3 days. Then seedlings were collected and incubated for 10 min
605 under vacuum (0.05 MPa) in phosphate-buffered saline (PBS) containing 4% paraformaldehyde
606 and 0.1% Triton X-100. Seedlings were washed gently three times (10 min for each wash) in PBS
607 and then the cell wall was digested in 2% Driselase (Sigma, D8037) in PBS for 18 min at 37 °C and
608 washed five times with PBS. The permeability of the seedlings was increased by incubating them
609 in 3% IGEPAL CA-630 (Sigma, 18896) and 10% DMSO in PBS for 18 min, followed by washed three
610 times with PBS. Seedlings were incubated in 2% bovine serum albumin (BSA) (Ameresco, 0332) in

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611 PBS for 1.5 h and then incubated with primary TOR antibody³² (antibody diluted 1:600 in 2% BSA)
612 for overnight at 4°C. The seedlings were washed with PBS for five times. Fluorophore-labeled
613 secondary antibody (goat–mouse secondary antibody, diluted 1:600 in 2% BSA) was incubated
614 with the samples at 37°C for 5 h in the dark. Seedlings were washed five times with PBS before
615 observation. Fluorescent signal detection and documentation were performed using a Nikon
616 confocal laser scanning microscope with a 560-nm band-pass filter for IF555 detection.

617
618 **S6K-p/S6K immunoblotting detection.** For immunological detection of S6K-p and S6K1/2, total
619 soluble proteins were extracted from 50 mg of plant materials grown as indicated previously with
620 100 µL 2 × Laemmli buffer supplemented with 1% Phosphatase Inhibitor Cocktail 2 (Thermo
621 Fisher Scientific Inc., 78430). Proteins were denatured for 10 min at 95°C and separated on 10%
622 or 8% SDS-PAGE. Rabbit AtTOR polyclonal antibodies (Abiocode, R2854-2), rabbit polyclonal
623 S6K1/2 antibodies (Agrisera, AS121855), and S6K1-p (phospho T449) antibody (Abcam,
624 ab207399) and ACTIN antibody (Abmart, M20009) were used for immunoblotting.

625
626 **Quantitative PCR (qPCR).** Total root RNA was extracted from plantlets grown *in vitro* at 22 °C and
627 10 °C using the RNeasy®Plant Mini Kit (QIAGEN, Germany). One microgram of total RNA was
628 reverse transcribed using an oligo(dT)₂₀ primer and the Super Script™ IV RT (Invitrogen,USA)
629 according to the manufacturer's instructions. cDNA was diluted 20-fold before PCR. qPCR was
630 performed on a LightCycler®480 Instrument II (Roche, USA) using 2 µL of 5X HOT
631 FIREPol®EvaGreen® qPCR Mix Plus(no ROX) (Solis BioDyne, Estonia), 2 µL of cDNA, and 0.25 µM of
632 each primer in a total volume of 10 µL per reaction. ACT2 (AT3G18780) gene was used as
633 reference for normalization of gene expression levels (ACT2 primers, F:
634 GGTAACATTGTGCTCAGTGGTGG R: CTCGGCCTTGGAGATCCACATC; TOR primers, F:
635 GAAGATGAAGATCCCGCTGA R: GCATCTCCAAGCATATTTACAGC⁴⁴). The cycling conditions were: 95
636 °C for 12 min., 35 cycles of 95°C for 15 sec., 60°C for 1 min. and finally a melting curve from 60°C
637 to 95 °C (0.05°/sec). Data were analyzed using the $\Delta\Delta C_t$ method⁹³ and LightCycler®480 Software,
638 version 1.5 (Roche). Two independent experiments with three biological and three technical
639 replicates per experiment, were performed.

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649 project ID 390686111) to G.G., and by the Czech Science Foundation GAČR (project Nr. 19-
650 18675S) to J.Š.

651

652

653 **Author Contribution**

654 J.M.P performed most of the experiments, analysed the data and helped in the writing process of
655 the manuscript. L.S. performed the S6K-p determinations, IP of TOR-FER and immunolocalization
656 assay of TOR. L.K., M.O. and J.Š. provided live cell imaging data on FER and ROP2. V.B.G., J.M.P.,
657 T.U.I., M.A.I., S-Z., Y.S., R.A.G., M.S., L.A.R., J.M.A., G.G., J.S. helped in the data analysis and
658 writing process of the manuscript. F.Y. designed research and analysed part of the data and
659 J.M.E. designed research, analysed the data, supervised the project, and wrote the paper. All
660 authors commented on the results and the manuscript. This manuscript has not been published
661 and is not under consideration for publication elsewhere. All the authors have read the
662 manuscript and have approved this submission.

663

664

665 **Competing financial interest**

666 The authors declare no competing financial interests. Correspondence and requests for materials
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