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# **Enhanced cell-cell contact stability and decreased N-cadherin-mediated migration upon Fibroblast Growth Factor Receptor-N-cadherin cross-talk**

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***Running Head:* Strengthening N-cadherin cell-cell contacts by FGFR.**

***Conflict of interest:*** The authors declare that they have no conflict of interest.

***Abbreviations:***

FGF: Fibroblast Growth Factor

FGFR: Fibroblast Growth Factor Receptor

FP: Fluorescent Protein

FN: Fibronectin

FRAP: Fluorescence Recovery After Photobleaching

GFP: Green Fluorescent Protein

Ncad: N-cadherin

PBS: Phosphate Buffer Saline

RTK: Receptor Tyrosine Kinase

SEM: Standard Error of the Mean

## Abstract

N-cadherin adhesion has been reported to enhance cancer and neuronal cell migration either by mediating actomyosin-based force transduction or initiating Fibroblast Growth Factor Receptor (FGFR)-dependent biochemical signalling. Here we show that FGFR1 reduces N-cadherin-mediated cell migration. Both proteins are co-stabilised at cell-cell contacts through direct interaction. As a consequence, cell adhesion is strengthened, limiting the migration of cells on N-cadherin. Both the inhibition of migration and the stabilisation of cell adhesions require the FGFR activity stimulated by N-cadherin engagement. FGFR1 stabilises N-cadherin at the cell membrane through a pathway involving Src and p120. Moreover, FGFR1 stimulates the anchoring of N-cadherin to actin. We found that the migratory behaviour of cells depends on an optimum balance between FGFR-regulated N-cadherin adhesion and actin dynamics. Based on these findings we propose a positive feedback loop between N-cadherin and FGFR at adhesion sites limiting N-cadherin-based single cell migration.



## Introduction

Cell adhesion and migration are central processes in morphogenesis, wound healing and cancerogenesis. Cells adhere and migrate on extracellular matrices thanks to their integrin receptors<sup>23</sup>. Many cells such as border cells in the *Drosophila* egg chamber<sup>58</sup>, neuronal precursors<sup>25, 39</sup> or cancer cells also adhere and migrate on the plasma membrane of the adjacent cells. In this case, cell migration is mediated by cadherins which physically hold cells together<sup>43</sup>. Changes in expression or function of cadherins have major impacts on cell migration during neural development<sup>59, 62, 70</sup> and tumour cell invasion<sup>28, 74, 76</sup>.

Cadherins are the homophilic ligands of *Adherens Junctions* (AJ) involved in the cohesion of solid tissues<sup>47</sup>. Cadherins provide anchorage between neighbouring cells thanks to their interaction with the contractile actomyosin network via catenins<sup>46</sup>. E-cadherin is required for epithelial cell cohesion<sup>21</sup> and is recognised as a tumor suppressor<sup>28, 73</sup>. N-cadherin, the neuronal cadherin, although required for the cohesive interaction of neuroepithelial cells<sup>26</sup>, mediates weaker cell-cell adhesion and is also associated with physiological and pathological cell migration in a large range of tissues<sup>16, 51, 65</sup>. N-cadherin ensures weak adhesion between post-mitotic neurons and radial glial cells allowing radial neuronal migration<sup>17, 25</sup>. Its active endocytosis and turnover maintain proper steady-state level of N-cadherin at the cell surface allowing the effective locomotion of neurons<sup>25</sup>. It is also required for long distance migration of tangentially migrating interneuron precursors<sup>39</sup>. Moreover, N-cadherin stimulates neurite outgrowth<sup>2, 3, 37, 42, 77</sup> *in vitro*. Two pathways have been involved: (i) the mechanical coupling of cadherins to actomyosin cytoskeleton, which generates the traction forces necessary to propel the growth cones<sup>2, 19</sup> and (ii) the activation of FGFR-dependent biochemical signalling cascades<sup>4, 77</sup>.

FGFRs belong to the family of single pass transmembrane Receptor Tyrosine Kinases. Binding of their ligands, FGFs, triggers intracellular signalling cascades playing key roles during development and pathogenesis<sup>36, 45</sup>. Loss of expression of FGFR1 in mice disrupts the migration of epidermal cells from the primitive streak. This phenotype can be rescued by down-regulating E-cadherin-mediated intercellular adhesion<sup>10, 11, 15, 79</sup>. In *Drosophila*, the migration of tracheal cells requires FGFR signalling, which regulates cytoskeletal reorganisation<sup>8, 35, 54</sup>.

Dysfunctions of N-cadherin and FGFRs both induce pathological migrations that are most visible in cancers. N-cadherin upregulation correlates with increased motility and invasiveness of dysplastic cells in melanoma<sup>38</sup>, bladder<sup>61</sup>, prostate<sup>31</sup>, lung<sup>49</sup> or breast

cancers<sup>48</sup>. Mutations in FGFRs are associated to pancreatic, endometrial, bladder, prostate, lung and breast cancers<sup>57, 75</sup>. Literature reports on a synergistic action between N-cadherin and FGFRs in the regulation of epiblast stem cells pluripotency<sup>67</sup>, ovarian cells survival<sup>71</sup> and osteogenic cells differentiation<sup>13</sup>. Overexpression of N-cadherin in mEpiSC cells prevents the downregulation of FGFR at the plasma membrane after FGF2 addition<sup>67</sup>. FGF and N-cadherin maintain granulosa and ovarian cells viability *in vitro* by stimulating FGFR phosphorylation<sup>71</sup>. The expression of a constitutively active form of FGFR increases the expression of N-cadherin reinforcing cell-cell adhesion in human osteogenic cells<sup>13</sup>. A functional relationship between FGFR and N-cadherin has been reported during neurite outgrowth<sup>4, 77</sup>. FGFR and N-cadherin co-cluster and interact at the surface of neuronal cells<sup>4, 72</sup>. The expression of a dominant negative FGFR inhibits neurite growth stimulated by N-cadherin<sup>5</sup>. In breast cancer cells, N-cadherin overexpression increases cell migration<sup>22</sup>. N-cadherin prevents FGFR from undergoing ligand-induced internalisation, resulting in FGFR stabilisation at the plasma membrane and sustained FGFR signalling<sup>64</sup>. In human pancreatic cancer xenografts, inhibition of FGFR leads to a decrease in N-cadherin expression and cell invasion<sup>66</sup>. Altogether, these data suggest that N-cadherin and FGFR synergise to generate signals that regulate the migratory behaviours of normal and cancer cells.

Little is known however about the combined effects of N-cadherin and FGFR activities on cell adhesion and migration. To dissect the reciprocal interplay between FGFR1 and N-cadherin, we expressed both receptors in HEK cells and analysed the consequences on N-cadherin-dependent cell adhesion and cell migration using a single cell migration model on N-cadherin coated lines. Both proteins are co-recruited and co-stabilised at cadherin-mediated cell contacts through direct interaction of their extracellular domains. As a consequence, N-cadherin-mediated cell contacts are strengthened, limiting the migration of cells on N-cadherin coated surfaces. Both the inhibition of N-cadherin-mediated migration and the stabilisation of N-cadherin at cell contacts require FGFR activity, which is itself stimulated by N-cadherin engagement. We further show that FGFR1 stabilises N-cadherin at the cell membrane by decreasing its internalisation. FGFR1 expression triggers an increase of activated Src but does not affect significantly the phosphorylated p120 catenin on tyrosine 228. However, both p120 and Src are involved in the stabilisation of N-cadherin at cell-cell contacts and in the negative regulation of N-cadherin-mediated migration induced by FGFR1. Moreover, FGFR1 stimulates the anchoring of N-cadherin to actin. Finally, we found that the migratory behaviour of cells depends on an optimum balance between FGFR-regulated N-cadherin adhesion and actin dynamics.

## Results

### FGFR1 expression inhibits N-cadherin-mediated cell migration

To study single N-cadherin-mediated cell migration and the impact on this migration on Fibroblast Growth Factor Receptor 1 (FGFR1), we developed a model in which isolated N-cadherin transfected (Ncad cells) and N-cadherin/FGFR1 double transfected HEK cells (Ncad/FGFR cells) were allowed to adhere and migrate on 10  $\mu\text{m}$  width Ncad-Fc-coated stripes, in the absence of exogenously added FGF (**Fig. 1A, Video 1**). Both Ncad and Ncad/FGFR cells adhered to the surface while untransfected HEK cells did not (data not shown). Ncad cells migrated efficiently covering a total displacement up to 400  $\mu\text{m}$  over 20 hours with very few inversions of direction of migration, while Ncad/FGFR cells were almost stationary (**Fig. 1B-E**). Treatment of Ncad/FGFR cells with PD173074, a FGFR kinase inhibitor, restored the migratory behaviour of Ncad/FGFR cells close to that of Ncad cells (**Fig. 1A-E**), indicating that the inhibition of N-cadherin-mediated migration by FGFR1 requires the receptor kinase activity. This effect is specific to N-cadherin mediated adhesion, as we observed that the expression of FGFR1 did not impact on the migration of Ncad expressing cells on fibronectin coated surfaces (data not shown). Moreover, we verified that the blockade of FGFR activity strongly increased the migration of cells endogenously expressing N-cad and FGFR, i.e., C2C12 mouse myoblastic, U2OS human osteosarcoma and 1205Lu human metastatic melanoma (**Fig. S1**), that otherwise only moved at a similar speed as Ncad/FGFR HEK cells. Altogether these data indicate that FGFR1 strongly impairs the migration of cells on N-cadherin in a process depending on its kinase activity, but in the absence of exogenous FGF. Accordingly, further experiments were all performed in the absence of FGF.

Ncad/FGFR cells were more spread than Ncad cells, a trend that was reverted in the presence of the FGFR inhibitor (**Fig. 1C, Video 1**). The mean spreading area of Ncad/FGFR cells of  $764.7 \pm 24.9 \mu\text{m}^2$  was reduced to  $562.6 \pm 10.6 \mu\text{m}^2$  in the presence of the inhibitor, which is close to the values measured for Ncad cells ( $302.8 \pm 13.4 \mu\text{m}^2$ ). Plotting the mean cell speed as a function of cell area confirmed an inverse correlation between these two parameters: the more the cells spread on N-cadherin, the slower they migrate (**Fig. 1F**). Ncad/FGFR cells displayed an extensive spreading and a reduced migration speed. Thus, the reduced migration of FGFR1 expressing cells could result from a strengthening of cadherin-mediated adhesion on the Ncad-Fc coated lines.

## **N-cadherin and FGFR1 are co-stabilised at cell-cell contacts**

We hypothesised that FGFR1 increases N-cadherin-mediated cell-cell adhesion by affecting the dynamics of junctional N-cadherin. To test this hypothesis, we performed dual wavelength FRAP (Fluorescence Recovery After Photobleaching) experiments at cell-cell contacts of cells expressing DsRed-Ncad or GFP-FGFR1 (alone) or both, in the absence of added FGF (**Fig. 2A, B**). Expressed alone, N-cadherin displayed a mobile fraction at the cell-cell contacts of  $61.3 \pm 2.7$  %. Coexpression of FGFR1 decreased this value to  $38.2 \pm 3.4$  % while treatment with the FGFR inhibitor restored N-cadherin mobile fraction level ( $62.3 \pm 2.3$  %) close to that of Ncad cells. Expression of N-cadherin also significantly decreased the mobile fraction of junctional FGFR1 (**Fig. 2C**). Similar experiments were performed using mCherry-tagged E-cadherin (mCherry-Ecad) instead of DsRed-Ncad, revealing that the E-cadherin expression did not affect the dynamics of FGFR1 at cell-cell contacts (**Fig. 2C**). FGFR1 expression did not modify the mobile fraction of E-cadherin at cell-cell contacts (**Fig. 2D**). Thus, FGFR1 and N-cadherin specifically co-stabilise each other at N-cadherin-mediated contacts. This co-stabilisation may lead to the strengthening of N-cadherin adhesion, which may explain the increased spreading of cells on Ncad-coated lines upon FGFR1 expression.

## **FGFR1 stimulates junctional N-cadherin accumulation and strengthens cell-cell contacts**

To confirm the strengthening effect of FGFR1 on N-cadherin-mediated adhesion, we quantified the accumulation of N-cadherin at cell-cell contacts in cell monolayers. Cell-cell contacts accumulated more N-cadherin and were straighter for Ncad/FGFR than for Ncad cells (**Fig. S2A**). FGFR inhibitor treatment inhibited the effect of FGFR1 on both the straightness and junctional accumulation of N-cadherin, indicating that the kinase activity of the receptor is required for the strengthening of N-cadherin-mediated cell contacts. Accordingly, analysing N-cad distribution at cell-cell contacts in cell doublets grown on fibronectin-coated lines, as well as in suspended cell doublets, revealed an increased accumulation of junctional N-cadherin in the presence of FGFR1 (**Fig. S2B, S3**).

To determine the impact of this junctional N-cadherin stabilisation on cell-cell contact stability, we followed by live imaging the disassembly of cell-cell contacts upon chelation of  $\text{Ca}^{2+}$  ions in cell monolayers (**Fig. 3A**). Quantitative analysis of cell-cell contact life-time

following  $\text{Ca}^{2+}$  depletion indicated that Ncad cells were dissociated in 2 minutes while it took almost 6 minutes to dissociate Ncad/FGFR cells. This cell-cell contact stabilisation was prevented by FGFR activity inhibition. Notice that the inhibitor had no effect on the dissociation of intercellular contacts of Ncad cells. To test more directly the effect of FGFR1 on the strength of N-cadherin-mediated adhesion, we probed the response to force developed at contacts between Ncad-Fc-coated magnetic beads and Ncad or Ncad/FGFR cells. Beads were left to interact with the cell surface for 30 minutes, before being probed for displacement under force by approaching a magnetic rod. After calibration, one can estimate the actual forces at which the N-cadherin-mediated adhesions between the bead and the plasma membrane were disrupted (**Fig. 3B, Video 2**). Significantly fewer beads were displaced or detached from the cell surface of Ncad/FGFR cells indicating that the binding strength was higher compared to Ncad cells. The inhibition of the FGFR activity restored bead detachment/displacement level close to the one observed for Ncad cells (**Fig. 3C**). For the population of beads that were detached under force, the mean breaking distance was of  $28.5 \pm 0.9 \mu\text{m}$  for Ncad cells and  $14.3 \pm 0.6 \mu\text{m}$  for Ncad/FGFR cells, respectively. Inhibition of FGFR increased the breaking distance to  $21.4 \pm 0.9 \mu\text{m}$  in Ncad/FGFR cells, corresponding to rupture forces of  $5.9 \pm 0.1 \text{ nN}$ ,  $7.3 \pm 0.1 \text{ nN}$  and  $6.5 \pm 0.1 \text{ nN}$ , respectively. Thus, FGFR1 and its kinase activity increase the mechanical resistance of N-cadherin adhesion.

### **N-cadherin and FGFR1 interaction promotes FGFR1 activation**

We described so far the effects of FGFR1 on N-cadherin-mediated adhesion and migration, both requiring the kinase activity of the receptor although no exogenous FGF ligand was added. Furthermore, FGFR1 and N-cadherin co-localised and co-stabilised at the cadherin-mediated cell contacts. Therefore, we hypothesised that the increased residence of FGFR at cell-cell contacts induced by N-cadherin-mediated adhesion could induce an activation of the receptor relying on direct interaction of these two proteins, as previously reported in neuronal cells<sup>4</sup>. To confirm this hypothesis, the level of binding of Ncad-Fc to immobilised FGFR1 extracellular domain was measured using an optical biosensor. Results showed a direct interaction between N-cadherin and FGFR1 ectodomain with an affinity, calculated from the kinetic parameters of the interaction, of  $106 \pm 25 \text{ nM}$  (**Fig. 4A and Suppl. Table I**). This interaction was confirmed by co-immunoprecipitation of GFP-FGFR1

from protein extracts of HEK cells co-expressing the two proteins (**Fig. 4B**). The co-precipitation was strongly reduced when FGFR kinase activity was inhibited.

Then, we tested whether N-cadherin could induce FGFR activation in the absence of exogenous FGF. Our results revealed that FGFR1 phosphorylation was significantly increased in Ncad/FGFR cells compared to cells only expressing FGFR1 (**Fig. 4C**). To further provide evidence that FGFR1 was activated by N-cadherin-mediated adhesion, we probed the activation of Erk1/2, a well-known downstream effector of FGFR signalling, following  $\text{Ca}^{2+}$  switch in C2C12 cells that express endogenous N-cadherin<sup>18</sup> and FGFRs<sup>32</sup> (**Fig. S4**). Addition of  $\text{Ca}^{2+}$  for 10 minutes to  $\text{Ca}^{2+}$ -depleted cells significantly increased Erk1/2 phosphorylation in the absence, but not in the presence of the FGFR inhibitor, strongly suggesting that N-cadherin engagement triggers the activation of the FGFR1. Altogether our results suggest a two-way communication between FGFR1 and N-cadherin resulting from their direct interaction. The stabilisation of FGFR1 by N-cadherin at cell-cell contacts allows its activation. The activation of FGFR1 could in turn increase junctional N-cadherin stabilisation, responsible for the observed strengthening of N-cadherin-mediated cell adhesion and reduction of N-cadherin-dependent cell migration.

### **FGFR1 stabilises N-cadherin at the plasma membrane through downregulation of endocytosis.**

To determine whether FGFR1 expression also increases N-cadherin prevalence at the plasma membrane, we performed cell surface biotinylation on Ncad and Ncad/FGFR HEK cells. The fraction of cell surface exposed biotin-labelled N-cadherin was significantly higher in Ncad/FGFR cells than in Ncad cells. It was strongly decreased in Ncad/FGFR cells following treatment with the FGFR inhibitor (**Fig. 5A**). Thus, FGFR1 favours the accumulation of N-cadherin at the plasma membrane in a process depending on its kinase activity. A first hint on the way FGFR1 may regulate N-cadherin availability at the cell surface was given by imaging DsRed-Ncad and analysing its distribution in Ncad or Ncad/FGFR cells thanks to flow cytometry imaging (**Fig. 5B**). Accordingly, when imaging DsRed-Ncad in cells migrating on fibronectin-coated lines (**Video 3**), we observed N-cadherin vesicles trafficking from the leading edge to the rear of the cells. These vesicles were more prominent in Ncad than in Ncad/FGFR cells, suggesting that the trafficking of N-cadherin was reduced in the latter (**Fig. 5C**). We thus questioned the role of endocytosis in the

regulation of cell surface N-cadherin by FGFR1 by quantifying N-cadherin endocytosis following the internalisation of biotinylated cell surface proteins (**Fig. 5D, Fig. S5**). The N-cadherin endocytic pool was significantly reduced in Ncad/FGFR cells compared to Ncad cells. This effect was significantly reduced in the presence of FGFR inhibitor, although the inhibition was far from complete (**Fig. S5**). Moreover, treatment of Ncad and Ncad/FGFR cells with hydroxyl-dynasore, an endocytosis inhibitor decreased the fraction of endocytosed N-cadherin in Ncad cells to levels measured for Ncad/FGFR cells (**Fig. 5D**). These data support the notion that FGFR1 upregulates N-cadherin prevalence at the plasma membrane by inhibiting its endocytosis, a process that could contribute to the reinforcement of N-cadherin-mediated cell contacts.

### **The effects of FGFR1 on N-cadherin-mediated adhesion and migration involves p120**

p120 has been reported to stabilise cadherins at cell-cell contacts by regulating their trafficking<sup>24</sup> either to the plasma membrane<sup>7</sup> or from the plasma membrane to endocytic compartments<sup>12, 78</sup>. In particular, it would do so by binding and masking an endocytic signal conserved in classical cadherins<sup>50</sup>. We thus asked whether the interaction of N-cadherin with p120 was involved in the regulation of N-cadherin endocytosis by FGFR1. First, we measured the intensity of GFP-p120 fluorescence along Ncad or Ncad/FGFR cells in contact (doublets). The results showed an increased recruitment of GFP-p120 at cell-cell contacts in Ncad/FGFR cells compared to Ncad cells (**Fig. 6A**). Then, to test the implication of p120, we coexpressed FP tagged FGFR1 and the NcadAAA mutant. The AAA mutation at position 764 in E-cadherin<sup>68</sup> and N-cadherin<sup>69</sup> was described to impair their binding to p120. FRAP experiments on Ncad/FGFR and NcadAAA/FGFR cells revealed that the mobile fraction of the mutated N-cadherin was significantly higher than that of the wild type molecule (**Fig. 6B**), suggesting that the binding of p120 to N-cadherin was involved in the stabilisation of N-cadherin induced by FGFR1. To test whether the ability of N-cadherin to bind p120 also affects N-cadherin-mediated cell migration, we compared the migration of Ncad/FGFR and NcadAAA/FGFR cells (**Fig. 6C, D and Video 4**). NcadAAA/FGFR cells displayed reduced spreading areas and increased migration speeds compared to Ncad/FGFR cells (**Fig. 6C**). They migrated at a speed very similar to the one of Ncad cells (see **Fig 1D**). Interestingly, the mutation drastically reduced the propensity of N-cadherin to interact with FGFR1 (**Fig. S6**).

Thus, preventing the binding of N-cadherin to p120 strongly inhibits the junctional N-cadherin stabilisation and the single cell migration inhibition induced by FGFR1.

Next, we analysed the expression levels of p120 in Ncad, FGFR and Ncad/FGFR cells (**Fig. S7A**). However, the cellular levels of p120 were not significantly affected by the expression of FGFR1. The N-terminal phosphorylation domain of p120, containing tyrosine residues phosphorylated by Src family kinases, has been reported to regulate negatively N-cadherin stability at the plasma membrane<sup>24, 34, 40</sup>. We thus analysed the phosphorylation on Y228 of p120 (**Fig. S7B**). However, no significant effect of FGFR1 on the phosphorylation of p120 on this site was observed. Thus, although p120 has been reported as a substrate of Src<sup>27, 40</sup>, itself a well-known downstream target of FGFR1<sup>60, 80</sup>, the involvement of p120 in regulating N-cadherin trafficking upon FGFR1 may not rely on a Src-dependent tyrosine phosphorylation pathway. Accordingly, Src inhibition by PP2 had no effect on the level of p120 phosphorylation on Y228 in Ncad/FGFR cells (**Fig. S7B**).

### **The effects of FGFR1 on N-cadherin-mediated adhesion and migration involve Src family kinases**

We then investigated the levels of Src activation in N-cadherin immunocomplexes in Ncad and Ncad/FGFR cells (**Fig. 7A**). FGFR1 expression led to an increase of the phosphorylation in the Src catalytic domain, while FGFR inhibition prevented this increase, suggesting that FGFR1 kinase activity is responsible for the increase in activated Src associated to N-cadherin. In order to determine the involvement of Src in the stabilisation of N-cadherin-mediated adhesion induced by FGFR1 expression, we analysed the effect of Src inhibition on the mobility of junctional N-cadherin. FRAP experiments revealed that the inhibition of Src by PP2 restored high levels of mobile junctional N-cadherin in Ncad/FGFR cells, comparable to those found in cells that do not express the receptor (**Fig. 7B**). When Ncad/FGFR cells were submitted to the single cell migration assay on N-cadherin in the presence of PP2 they displayed a strong stimulation of their migration properties with a mean speed of migration comparable to the one of Ncad cells (**Fig. 7C and Video 5**). Thus, Src inhibition counteracts both the stabilisation of N-cadherin cell-cell contacts and the inhibition of N-cadherin-mediated migration induced by FGFR1 expression. These data suggest that the activation of Src by FGFR1 in N-cadherin complexes may regulate the stability of junctional cadherin and the migratory response of N-cadherin expressing cells.



## FGFR1 stiffens the anchoring of N-cadherin to actin network

The anchoring of cadherins to actomyosin has been reported as a major mechanosignalling leading to cell-cell contact reinforcement and neuronal cell migration<sup>19, 46</sup>. To evidence the effect of FGFR activity on the mechanical link between N-cadherin and actin, we analysed the retrograde flow of actin in the lamellipodia of LifeAct-GFP expressing C2C12 myogenic cells spread on N-cadherin as a proxy of the coupling of cadherin to the treadmilling actin<sup>56, 63</sup>. The speed of F-actin rearward flow was increased by 40% in cells treated with the FGFR inhibitor compared with cells treated by the vehicle alone (**Fig. 8A; Video 6**), indicating that FGFR activity stimulates the coupling of N-cadherin to the actin cytoskeleton.

To test whether this mechanocoupling modulation was instrumental in regulating N-cadherin-mediated cell migration, Ncad and Ncad/FGFR HEK cells were treated with the Arp2/3 inhibitor CK 666 and analysed for their migration on Ncad-coated lines (**Fig. 8B; Video 7**). While the inhibition of branched actin polymerisation almost fully abrogated the migration of Ncad cells, it significantly increased the migration of Ncad/FGFR cells on N-cadherin (compare **to Fig. 1**). These observations indicate a bimodal implication of actin polymerisation in N-cadherin-mediated adhesion that is necessary for the migration of cells displaying mild adhesion (Ncad cells), but prevents the migration of tightly adhering Ncad-FGFR cells, likely through the destabilisation of adhesions. Cell migration on N-cadherin thus requires an optimal adhesion that depends on the strength of the N-cadherin-F-actin mechanocoupling. To further support this hypothesis, we analysed the implication of myosin II, also contributing to the stabilisation of cadherin adhesion. Treatment with the myosin II inhibitor similarly blocked the migration of Ncad cells and stimulated the one of Ncad/FGFR cells (**Fig. S8**). Altogether, these data indicate that FGFR1 activity increases the coupling of N-cadherin complexes to the underlying cytoskeleton. The resulting strengthening of N-cadherin-mediated contacts contributes to the inhibition of cell migration on the N-cadherin substrate.

## Discussion

Although cadherin and FGFR dysfunctions are observed in cancer, their relation to cell migration and invasion remain unclear. N-cadherin facilitates either cell adhesion or cell migration, whereas FGFRs are either enhancers or repressors of cell migration. In light of reported cell type specific cadherin/tyrosine kinase growth factor receptor, E-cadherin and EGFR<sup>44</sup> and VE-cadherin and VEGFR<sup>6, 20</sup>, functional interactions, a crosstalk between N-cadherin and FGFR has been proposed<sup>4, 51, 64, 77</sup>, although the mechanisms by which it may affect cell adhesion and migration remained unclear.

To mimic N-cadherin-dependent neural or cancer cells migration over neighbouring cells, we set up a model system of isolated N-cadherin or N-cadherin/FGFR1 expressing cells migrating on recombinant N-cadherin-coated lines. We describe here a complex interplay between N-cadherin engagement and FGFR1 activation positively regulating the strength of cell-cell adhesion and decreasing cell migration on N-cadherin, which occurs in the absence of added FGF. FGFR1 expression dramatically blocked N-cadherin-dependent single cell migration. This inhibition was associated to an increased cell spreading due to a strengthening of N-cadherin-mediated adhesion. FGFR1 led to the reinforcement of N-cadherin adhesion as demonstrated by the increased recruitment and stabilisation of junctional N-cadherin. We do not know whether FGFR1 regulates directly the “cis”- or “trans”- clustering of N-cadherin that may affect its stability at the plasma membrane<sup>46</sup>. However, this stabilisation was associated to an increased resistance of cell-cell contacts to calcium depletion, to an increase in the coupling of cadherin complexes to the actin treadmilling and to a rise in the mechanical strength of cell contacts. The rupture force of N-cadherin-mediated bead-cell contacts measured here was in the same range than those reported for doublets of N-cadherin expressing S180 cells ( $7.7 \pm 1.4$  nN)<sup>9</sup>. This rupture force was significantly increased by FGFR1 expression.

Altogether, our data strongly support the hypothesis that FGFR1 blocks cell migration on N-cadherin by strengthening N-cadherin adhesion. This behaviour is reminiscent of the reported biphasic relationship between cell migration of cells on fibronectin and the strength of integrin mediated cell-substratum adhesion<sup>53, 55</sup>. Cell migration is enhanced with increasing adhesion up to a threshold, above which further increases in adhesion acts to the detriment of migration. Accordingly, cells expressing only N-cadherin were poorly spread and supported cycles of both adhesion and deadhesion, allowing them to migrate and invert their polarity

whereas cells expressing also FGFR1 remained tightly spread on the cadherin-coated substrate preventing migration. Pharmacological treatments altering actin polymerisation and actomyosin contraction, which destabilise cadherin adhesions<sup>46</sup>, blocked the migration of N-cadherin expressing cells, but stimulated the migration of cells expressing FGFR1. By analogy, the positive effect of FGFR signalling on the migration of neuronal cell growth cones<sup>4,77</sup> may be related to the intrinsic weak adhesion of neuronal cells.

The cellular responses reported in this study, including the regulation of N-cadherin stability at the plasma membrane and of the mechanocoupling between N-cadherin and actin require the receptor activation. Although we cannot exclude that FGFR1 could be activated by FGFs endogenously produced by HEK cells, these experiments have all been performed in the absence of exogenous FGF. Thus, we provide evidence that N-cadherin engagement by itself stimulates the activation of FGFR1, in agreement with previous observations made in neuronal cells<sup>4,72</sup>. The presence of N-cadherin strongly decreases the mobility of junctional FGFR1 suggesting that the receptor was trapped in adhesion complexes. This process is N-cadherin specific, as junctional FGFR1 stabilisation was not observed with E-cadherin. We confirmed by co-immunoprecipitation, but also using purified proteins, that FGFR1 and N-cadherin interact through their extracellular domain. This may be essential for FGFR1 activation; decreasing the mobility of the receptor and/or increasing its local density at cell-cell contacts may stimulate its dimerisation and cross phosphorylation.

The sustained activation of FGFR1 significantly increased N-cadherin levels at the plasma membrane. The level of expression of the p120 catenin has been reported to stabilise junctional cadherin by preventing their internalisation<sup>12, 24, 50</sup>. Accordingly, we found that a mutant of N-cadherin impaired for its binding to p120 was not stabilised at cell-cell junctions in FGFR1 expressing cells. Cells expressing this mutated cadherin together with FGFR1 spread poorly on N-cadherin and migrated at high speed. Thus, we propose that p120 is involved in the regulation of N-cadherin stabilisation at cell adhesion sites by FGFR1. It has been reported that the phosphorylation of this catenin may induce its dissociation from cadherin allowing endocytosis of the latter<sup>24</sup>. Thus, the negative effect of FGFR1 on p120 phosphorylation might be a relay to stabilise junctional N-cadherin. However, no changes were observed in the cellular levels of p120 tyrosine phosphorylation upon FGFR1 expression, in agreement with a previous study reporting that mutation of Y228 and other prominent Src-associated p120 phosphorylation sites did not noticeably reduce the ability of E-cadherin to assemble AJs<sup>41</sup>. Alternatively, p120-dependent endocytosis of N-cadherin upon

FGFR expression may rely on phosphorylation on serine residues, which have been reported to regulate p120 functions<sup>14, 33</sup>. Alternatively, the increased stability of N-cadherin at cell-cell contact may depend by the reported regulatory action of p120 on Rho GTPases activity<sup>1, 52</sup>.

The kinase activity of the receptor was only partially involved in the inhibition of N-cadherin endocytosis. Moreover, the N-cadherin mutant impaired for p120 binding displayed a reduce association to FGFR1, indicating that the regulation of N-cadherin availability at the plasma membrane by FGFR1 involves additional pathways. Accordingly, we unravelled an involvement of Src in the cellular response to FGFR1 expression, which is unrelated to p120. FGFR1 increased the amount of activated Src associated to N-cadherin immunocomplexes and blocking pharmacologically Src activity reverted the effect of FGFR1 on cell spreading and migration. The inhibition of Src had a blocking effect on the stabilisation of junctional N-cadherin induced by FGFR1, indicating that Src is also involved in the mechanocoupling between N-cadherin complexes and actomyosin. Altogether, N-cadherin stability at the plasma membrane inversely correlates with the migratory properties of the cells on N-cadherin substrates. It is important here to recall that, in the case of the radial migration of cortical neurons *in vivo*, efficient migration on radial glia requires an active recycling of N-cadherin in neurons<sup>17, 25, 29</sup>. In this system, both the blockade of N-cadherin recycling and N-cadherin overexpression induced abnormal stabilisation of cell-cell contacts and impaired cell migration.

Taken together, these data reveal the existence of a pathway controlled by FGFR1 and N-cadherin and regulating N-cadherin-dependent cell-cell adhesion and cell migration. FGFR1 and N-cadherin are co-recruited and co-stabilised at the cell-cell adhesions. This leads to sustained activation of FGFR1, which in turn promotes N-cadherin accumulation at the plasma membrane, strengthens N-cadherin mediated cell-cell contacts and N-cadherin mechanocoupling to actin. Adhesion between migrating cells and N-cadherin-expressing cellular substrates is increased therefore decreasing cell migration. This mechanism could be used by cancer cells to engraft to the vessel wall or the host tissue. In less adherent cells, such as neurons or for cancer cells in other locations or considering different type of cancer cells, depending on the level of expression of N-cadherin and the dynamics of the actomyosin cytoskeleton, the same pathway may promote cell migration or cell anchoring.

## **Materials and Methods**

### ***Plasmid constructions***

The construct encoding GFP-FGFR1 was constructed as described in supplementary methods using the mouse fgfr1-IIIc full sequence (gift from D. Ornitz, University of Washington).

### ***Cell culture and transfection***

HEK 293 (HEK), C2C12, 1205Lu and U2OS cells were grown, transfected and selected, as described in supplementary methods.

### ***Drug treatments***

The FGFR kinase activity inhibitor, PD173074 (Sigma, 20 nM final concentration), and the Src family proteins inhibitor, PP2 (Abcam, 100 nM final concentration), were added in the medium 30 minutes prior the beginning and maintain throughout the experiments. Hydroxy-dynasore (Sigma, 100 nM final concentration) was incubated for 1 hour.

### ***FGFR activation and $Ca^{2+}$ switch assay***

C2C12 cells cultures were starved in serum-free medium 24 hours and then treated for 5 minutes with 1ng/ml of FGF2<sup>30</sup>. Alternatively, starved cultures were first treated with 4 mM EGTA for 20 minutes and then washed and incubated in the presence of 5 mM of  $Ca^{2+}$ .

### ***Ncad-Fc line guided cell migration***

Guided cell migration on 10  $\mu$ m wide N-cadherin coated lines was performed as described in supplemental methods, in the absence of exogenously added FGF.

### ***Fluorescence Recovery after Photobleaching***

Dual wavelength fluorescence recoveries after photobleaching (FRAP) was performed at 37°C (see supplemental methods) and analysed as reported previously<sup>63</sup>.

### ***Magnetic tweezers***

A homemade magnetic tweezer was the source of the magnetic field gradient used to pull Ncad-Fc coated paramagnetic microbeads attached to the cells (see supplemental methods). For the measurement of the rupture force of N-cadherin-mediated bead-cell contacts, Ncad or Ncad/FGFR cells seeded on 10  $\mu$ m-width fibronectin coated lines were

incubated with 4.5  $\mu\text{m}$  magnetic Ncad-Fc coated beads for 30 minutes, then unbound beads were washed out. The magnetic microneedle was approached while cells and the moving tip were imaged in phase contrast (every 10 milliseconds during 2 minutes).

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## **Conflict of interest**

The authors declare that they have no conflict of interest.

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## Figure Legends

**Figure 1: FGFR decreases the migration of N-cadherin expressing cells on N-cadherin coated lines.** DsRed-Ncad (Ncad) and DsRed-Ncad/GFP-FGFR (Ncad/FGFR) expressing HEK cells were seeded at low density on 10  $\mu\text{m}$ -width Ncad-Fc coated lines in the absence or in the presence of FGFR inhibitor (Ncad/FGFR+inh) and imaged in phase contrast every 6 minutes during 20 hours (see Video 1). **(A)** Representative kymographs of the displacement over 10 hours of two cells for each condition. **(B)** The trajectories of single cells Ncad ( $n = 26$ ), Ncad/FGFR ( $n = 22$ ) and Ncad/FGFR+inh ( $n = 25$ ) were manual tracked and the cell displacement plotted over time. **(C)** Representative kymographs of 1-hour long cell displacements imaged at higher magnification (see Video 1). **(D, E)** Histograms representing the mean cell body speed and the frequency of inversion in migration direction, respectively, for Ncad, Ncad/FGFR and Ncad/FGFR+inh cells (\*\*  $p \leq 0.01$ , \*\*\*  $p \leq 0.0001$ , ANOVA multi-comparison test, Newman-Keuls post-test). **(F)** Plots of the mean cell body speed as a function of cell surface area for the three cell populations.

**Figure 2: FGFR and N-cadherin co-stabilise each other at cell-cell contacts.** **(A)** Representative images of FRAP experiments performed at the cell-cell contacts of DsRed-Ncad HEK cells: Fluorescent signal before (Pre-bleach), immediately after bleaching (Bleach) and 110 sec after the bleach (Post-bleach). Red rectangles represent the bleached region at cell-cell contacts. Scale bar: 40  $\mu\text{m}$ . **(B)** Left: normalised DsRed-Ncad fluorescence recovery curves for Ncad, Ncad/FGFR and Ncad/FGFR+inh cells, respectively.  $n = 28$  from 3 independent experiments. Right: mean Ncad mobile fraction  $\pm$  SEM, \*\*\*  $p \leq 0.001$ ; ns: non-significant, ANOVA multiple comparison test,  $n = 28$  from 3 independent experiments. **(C)** Left: normalised fluorescence recovery for GFP-FGFR in FGFR, FGFR/Ecad and FGFR/Ncad cells, respectively.  $n = 15$ . Right: mean FGFR mobile fraction  $\pm$  SEM, \*\*\*  $p \leq 0.0001$ ; ns: non-significant, ANOVA multi-comparison test,  $n = 15$  from 3 independent experiments. **(D)** Left: normalised mCherry-Ecad fluorescence recovery in mCherry-Ecad (grey) and mCherry-Ecad/FGFR (black) cells, respectively.  $n = 18$  from 3 independent experiments, Right: mean mCherry-Ecad mobile fraction  $\pm$  SEM, ns: non-significant, student t-test,  $n = 18$ .

**Figure 3: FGFR strengthens N-cadherin-mediated cell-cell contacts and reinforces N-**

**cadherin anchoring to the cell cortex.** (A) Ncad, Ncad/FGFR, Ncad/FGFR + inh cells cultured at confluence over 1 mm<sup>2</sup> square fibronectin-coated-patterns were treated with EGTA then imaged for DsRed-Ncad every 30 seconds during 15 min. **(top left)** Low magnification images taken after 5 min of EGTA treatment. Scale bar = 40 µm. **(bottom left)** Examples of kymographs of the DsRed-Ncad signal along a line perpendicular to the cell-cell contact starting from EGTA addition (t<sub>0</sub>) for Ncad, Ncad/FGFR and Ncad/FGFR+inh cells. **(right)** Contact dissociation time upon EGTA addition as determined from the kymographs for the three conditions, plus Ncad+inh cells. \*\* p≤0.01; \*\*\* p≤0.001, ANOVA multi comparison test, Newman-Keuls post-test, n = 60 contacts. (B) Left: magnetic tweezers experimental set up used to evaluate the anchorage and rupture force of N-cadherin mediated bead-cell contacts. Right: Calibration curve of the magnetic tweezers determined as described in material and methods. The force is exponentially inversely correlated with the bead-magnetic needle distance. (C, left) Representative images of Ncad-coated beads before and after tweezer-induced detachment from the cell membrane. **(middle)** Distribution of the responses of Ncad beads to the magnetic field in three classes (release, displacement, and immobility) for Ncad (n = 60), Ncad/FGFR (n = 65) and Ncad/FGFR+inh (n = 50) cells. **(right)** Bead-cell contact disruption forces calculated from the Stoke equation for Ncad, Ncad/FGFR and Ncad/FGFR + inh HEK cells (\*\*\* p ≤ 0.001, ANOVA multi comparison test, Newman-Keuls post-test).

**Figure 4: N-cadherin and FGFR associate leading to increased activation of FGFR.** (A) Binding of Ncad-Fc to FGFR1 extracellular domain. Kinetics of Ncad-Fc to immobilised FGFR1 extracellular domain was measured as described under “Materials and Methods.” Left panel: Ncad-Fc at different concentrations was added to a FGFR1-derivatised cuvette and the association reaction was followed for 200 s. Data were collected three times a second. The concentration of Ncad-Fc is indicated. Data shown are the result of one representative experiment out of three. Right panel: relationship between the extent of binding (response in arc s) of the association reactions shown in left and Ncad-Fc concentration. All results are summarised in supplementary table 1. (B) GFP-FGFR was immunoprecipitated with anti-GFP bead from protein extracts of Ncad, FGFR and Ncad/FGFR cells. Immunoprecipitates, together with total protein extracts, were then analysed by Western blot using anti-Ncad and anti-GFP (FGFR) antibodies. The histogram shows the ratio of N-cadherin bound to GFP-FGFR on N-cadherin in total extract, determined from the quantification of 3 independent



immunoblots then converted to percentage. \*\*  $p \leq 0.01$ , Student's  $t$  test,  $n = 3$ . (C) To detect FGFR phosphorylation GFP-FGFR immuno-precipitates were immunoblotted with anti-P-Tyr and anti-GFP (FGFR) antibodies. The histogram shows the ratio of P-Tyr on GFP-FGFR signals as a quantification of the degree of phosphorylation of FGFR in the different extracts. \*\*  $p \leq 0.01$ , Student  $t$  test,  $n = 3$ .

**Figure 5: FGFR increases N-cadherin cell surface accumulation by reducing its endocytosis.** (A) Analysis of cell surface expression of N-cadherin. After surface biotinylation at cold and removal of unfixed biotin, Ncad, Ncad/FGFR and Ncad/FGFR+inh cells were immediately lysed and protein extracts subjected to precipitation by streptavidin beads. GFP transfected HEK cells were used as control. Total extracts and streptavidin bound fractions (plasma membrane exposed fractions) were then immunoblotted with anti-N-cadherin antibodies. The histogram shows the quantification of N-cadherin exposed at the plasma membrane over total N-cadherin content for the three conditions. \*\*\*  $p \leq 0.001$ , ns: non-significant, ANOVA multi comparison test, Newman-Keuls post-test,  $n = 4$ . (B) Analysis of Ncad internal pool by flow cytometry imaging. Ncad and Ncad/FGFR cells were non-enzymatically detached, then processed for flow cytometry imaging in bright field, and for dsRed-Ncad and GFP-FGFR fluorescence imaging. Masks were defined on bright field images to separate cell membrane and internal cell areas on each cell. Applied to the fluorescence images they allowed to extract an internalisation score as described in Materials & Methods. FGFR reduces the internalisation score of N-cadherin molecule by 17% (1.09 U.I versus 1.32 U.I). Experiences were repeated 4 times, over populations of 150.000 cells for each condition in each experiment. (C) Ncad and Ncad/FGFR cells were seeded on Ncad-coated stripes of 10  $\mu\text{m}$ , then after 4 hours, preparations were imaged at 63X for Ds-Red Ncad. The panels show the maximum projection of 1  $\mu\text{m}$  thick confocal sections encompassing the whole cell thickness. Arrow-heads show N-cadherin puncta trafficking from the leading edge to the rear of Ncad expressing cells. The histogram shows the quantification of the percentage of cells with such puncta. \*\*  $p \leq 0.01$ ; non parametrical  $t$  test;  $n = 15$ ,  $n = 20$  cells for Ncad and Ncad/FGFR cells, respectively. (D) Analysis of N-cad endocytic fraction following cell surface biotinylation. Freshly biotinylated Ncad and Ncad/FGFR cells were switched to 37°C for 40 minutes to allow endocytosis to resume in the presence or in the absence of dynasore, then subject to a reducing wash in order to remove remaining medium exposed biotin. Left: cells were lysed and protein extracts subjected to precipitation by streptavidin beads, then

anti-N-cadherin bound (total) and streptavidin bound fractions (endocytosed) were immunoblotted with anti-N-cadherin antibodies. Right: The histogram shows the ratio of endocytosed over total Ncad in each extract. \*\*\*  $p \leq 0.001$ ; ns: non-significant, ANOVA multiple comparison test,  $n = 3$  experiments.

**Figure 6: p120 is involved in the stabilisation of N-cadherin at cell-cell contacts and the decreased migration induced by FGFR expression.** (A) Ncad and Ncad/Flag-FGFR cells were transfected with GFP-p120 and seeded on fibronectin coated lines. Mean p120 intensities along the cell length with 0 as junctional end and 1 as free end of the cell was calculated on 25 cell doublets. p120 junctional accumulation was higher in Ncad/FGFR cell doublets than in Ncad doublets. (B) FRAP experiments were performed on cells expressing GFP-FGFR1 and either DsRed-Ncad or mCherry-NcadAAA. Curves show Ncad and NcadAAA normalised fluorescence recoveries over time for Ncad/FGFR (black) and NcadAAA-FGFR (red) cells (mobile fraction  $0.29 \pm 0.1$  and  $0.50 \pm 0.1$ , respectively ( $n = 25$ ), (\*\*\*,  $p \leq 0.0001$ , Student t test). (C) Ncad/FGFR and NcadAAA/FGFR cells were seeded on Ncad-Fc coated stripes and imaged every 6 minutes during 20 hours. Left: examples of Ncad/FGFR and NcadAAA/FGFR individual cell displacements over 1 hour. Right: histograms representing the mean cell speeds as a function of cell areas. (D) Plots show the displacement in function of time for Ncad/FGFR (left), NcadAAA/FGFR (right) cells with respectively  $n = 30$ ,  $n = 40$  cells. Histograms show the mean speed of Ncad/FGFR (black), NcadAAA/FGFR (red) cells (\*\*\*\*,  $p \leq 0.0001$ , ANOVA multi-comparison test).

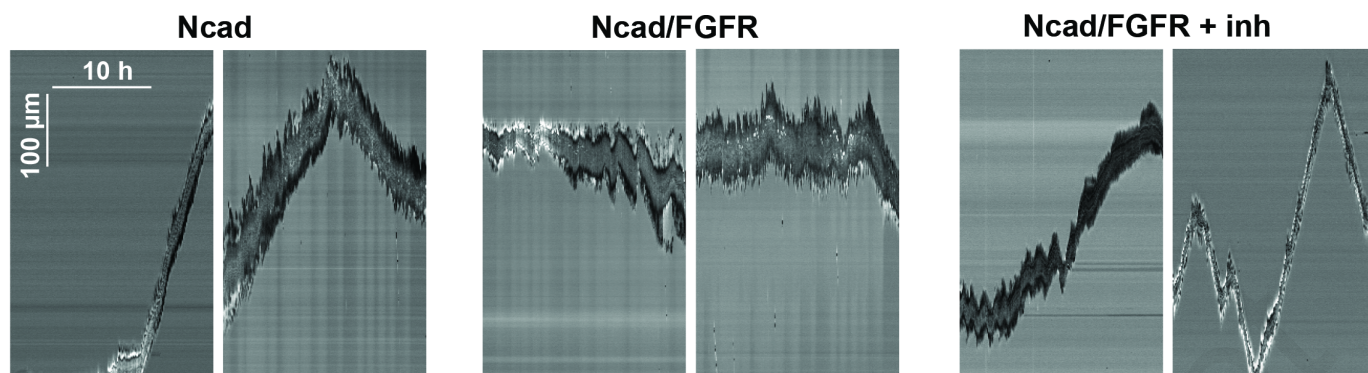
**Figure 7: Src activity is involved in the stabilisation of N-cadherin at cell-cell contacts and the decreased migration induced by FGFR expression.** (A) Western blot detection of N-cad, Src and phosphorylated Src (P-Src) in the Ncad immunoprecipitates. Histogram shows the ratio of phosphorylated Src calculated as the ratio of P-Src band intensity on Src band intensity. (B) FRAP experiments were performed on DsRd-N-cadherin at cell-cell contacts of Ncad and Ncad/FGFR in the absence or in the presence of Src inhibitor. Curves and histograms show Ncad and NcadAAA normalised fluorescence recoveries over time and extracted mobile fractions  $\pm$  SEM, \*\*\*  $p \leq 0.001$ ; ns: non-significant, ANOVA multiple comparison test,  $n = 18$ ). (C) Migration of Ncad/FGFR and Ncad/FGFR cells treated with the Src inhibitor on Ncad-Fc coated lines. Graph shows the cumulative cell displacements in

function of time and histogram the mean cell migration speeds. (\*\*  $p \leq 0,01$ , \*\*\*  $p \leq 0.0001$ , ANOVA multi-comparison test, Newman-Keuls post-test).

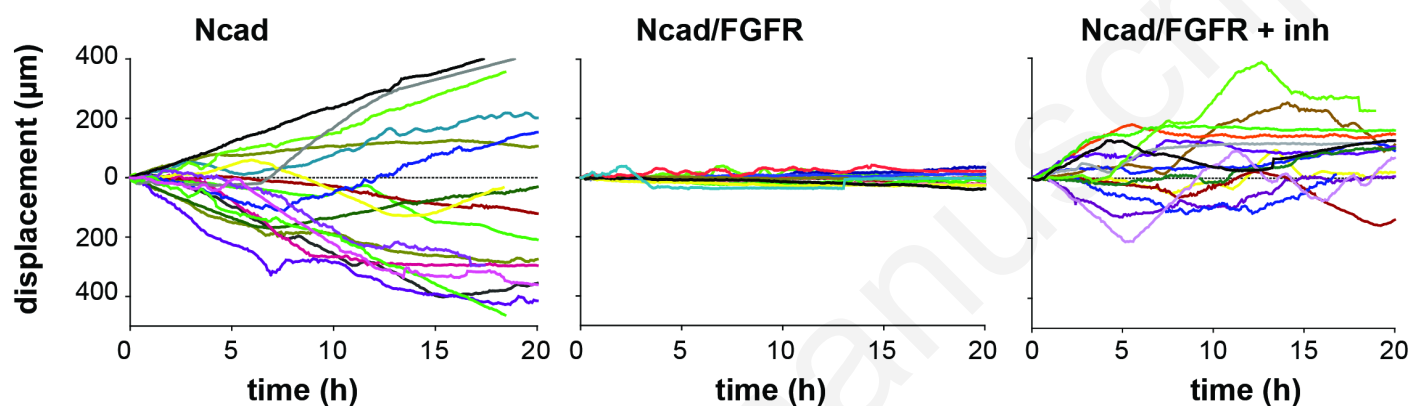
**Figure 8: FGFR promotes N-cadherin-F-actin functional mechanocoupling.** (A) LifeAct-GFP expressing C2C12 cells were seeded on Ncad-Fc coated surfaces for 2 hours, then treated with or without FGFR inhibitor for 1 hour, and then imaged for 5 minutes at a frequency of two images per second. Left: still images of the LifeAct-GFP signal, scale bar = 20  $\mu\text{m}$ . Inserts on the right represent examples of kymograph constructed along the two pixel-wide yellow lines (1–3), Right: the actin retrograde flow was quantified by kymograph analysis. Right: the histogram shows the mean actin retrograde flow speed for C2C12 ( $n = 140$  kymographs from 24 cells) and C2C12 + inh ( $n = 156$  kymographs from 25 cells) cells, (\*\*\*\*  $p \leq 0.0002$ , Student  $t$  test). (B) Migration on Ncad-Fc coated lines of Ncad and Ncad/FGFR cells treated or not with the Arp2/3 inhibitor. Left: kymographs of the displacement over 10 hours of three cells for each condition. Middle: cumulative displacements of cells in function of time (20 hours) for Ncad ( $n = 25$ ), Ncad/FGFR ( $n = 25$ ), Ncad + CK666 ( $n = 26$ ), Ncad/ FGFR + CK666 ( $n = 22$ ) conditions. Right: histograms representing the mean cell speed for each condition (\*\*  $p \leq 0,01$ , \*\*\*  $p \leq 0.0001$ , ANOVA multi-comparison test, Newman-Keuls post-test).

# Figure 1

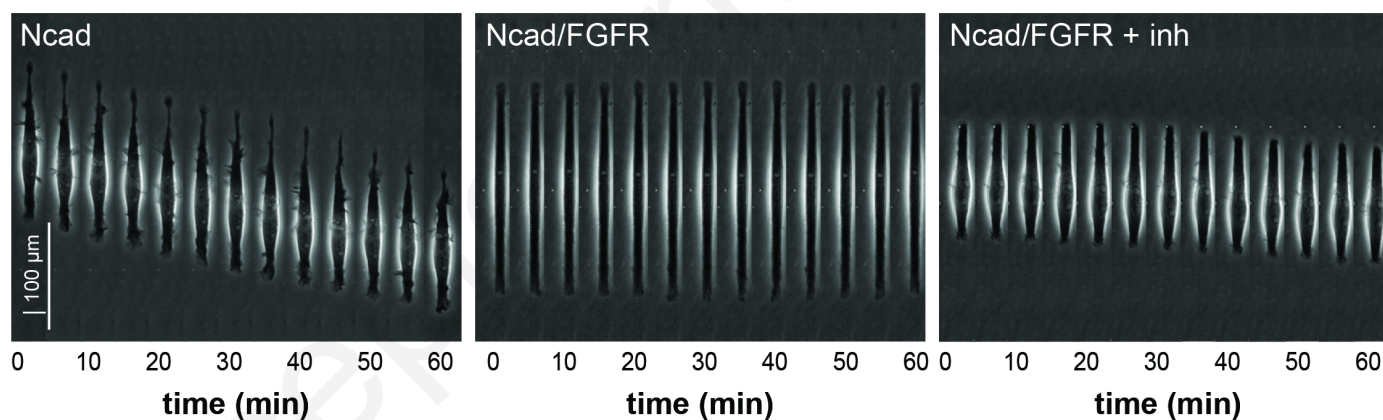
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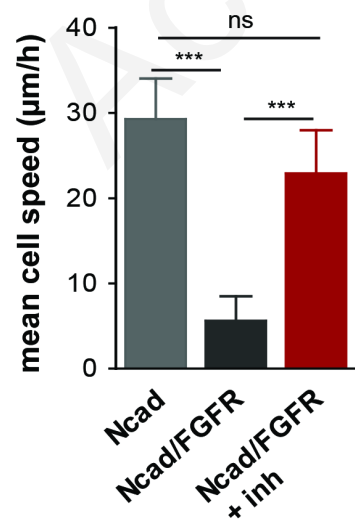
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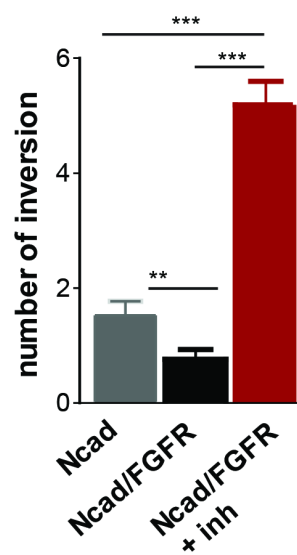
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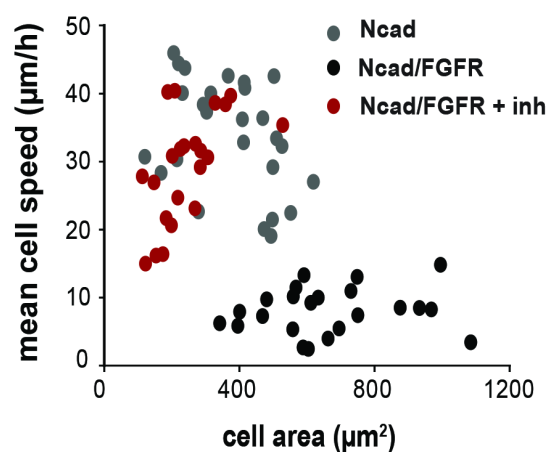
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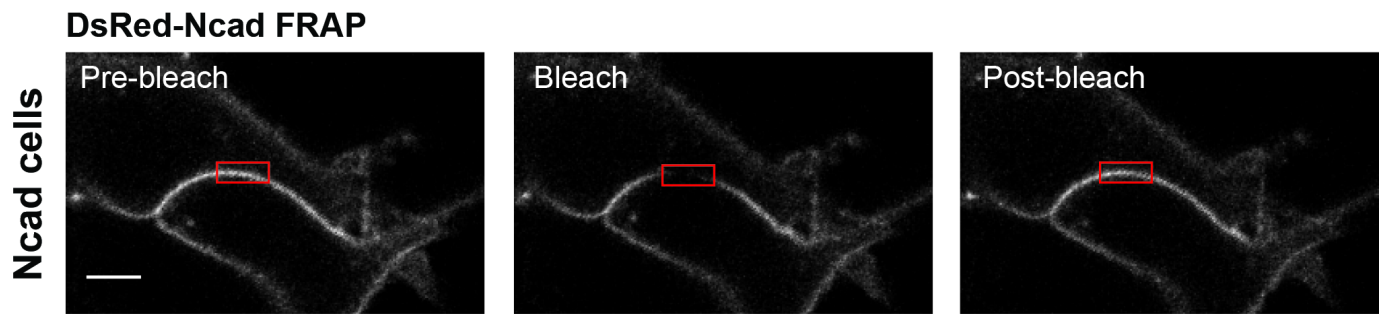


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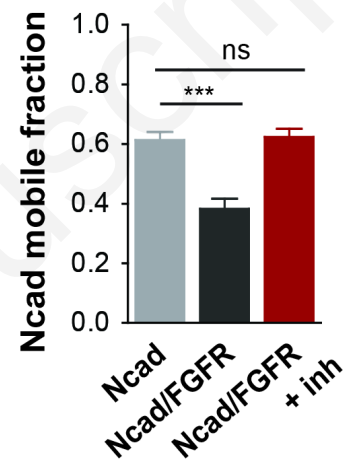
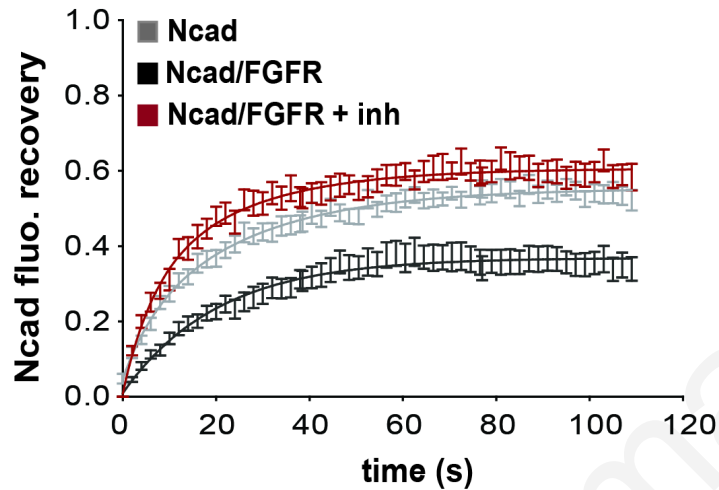


# Figure 2

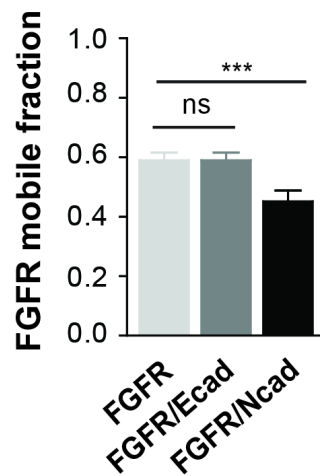
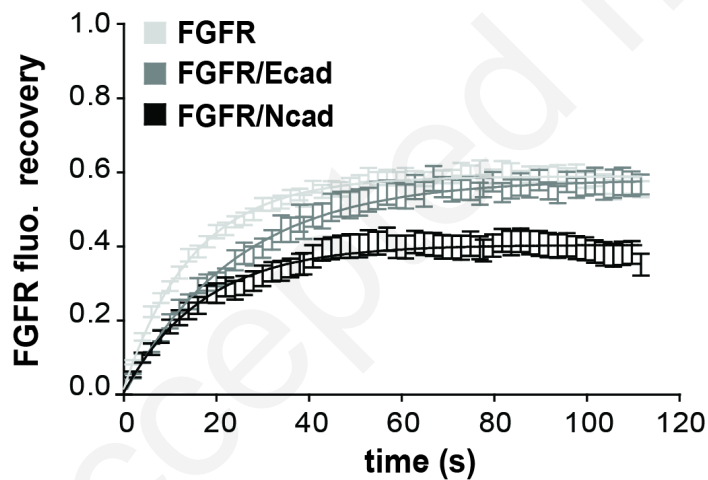
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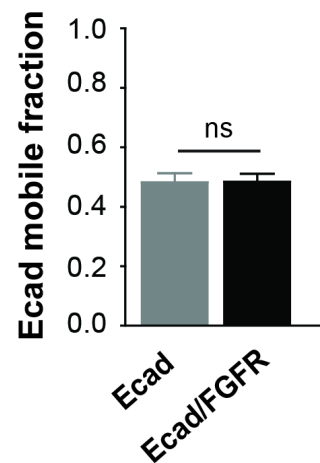
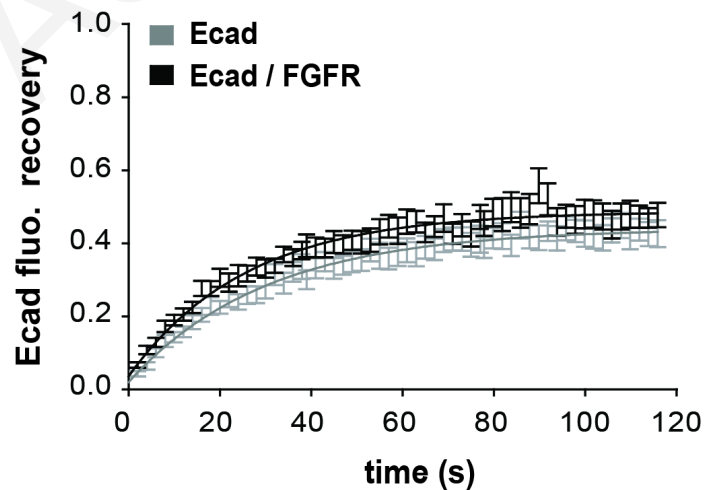
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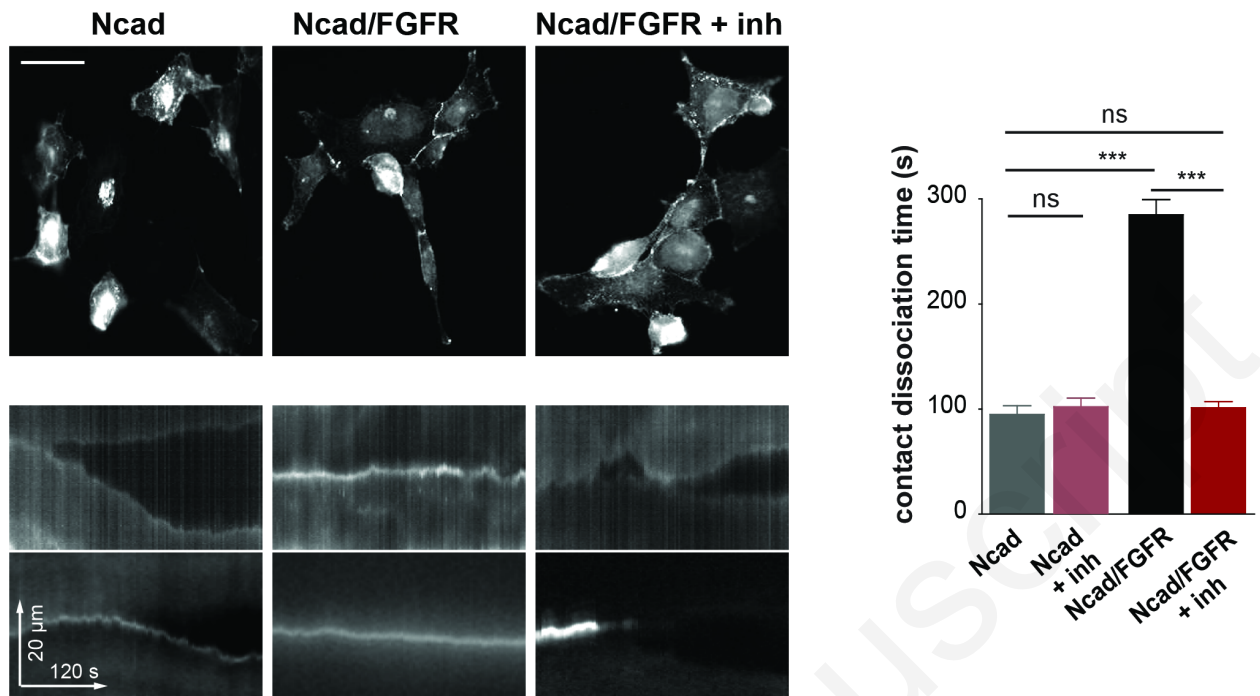


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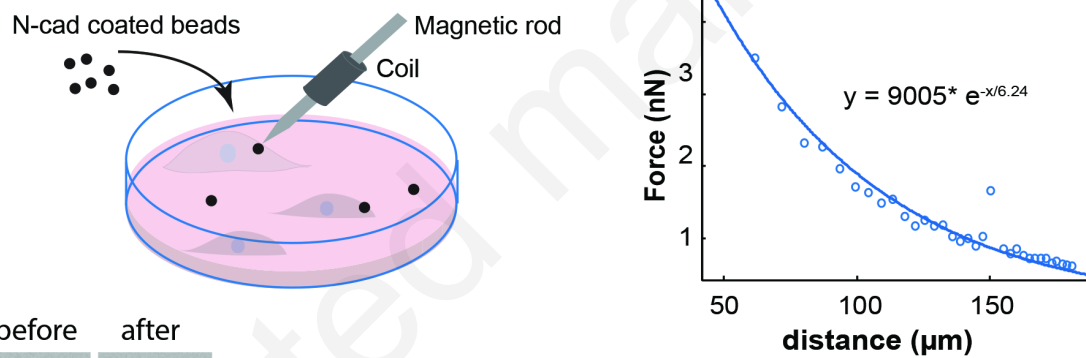


# Figure 3

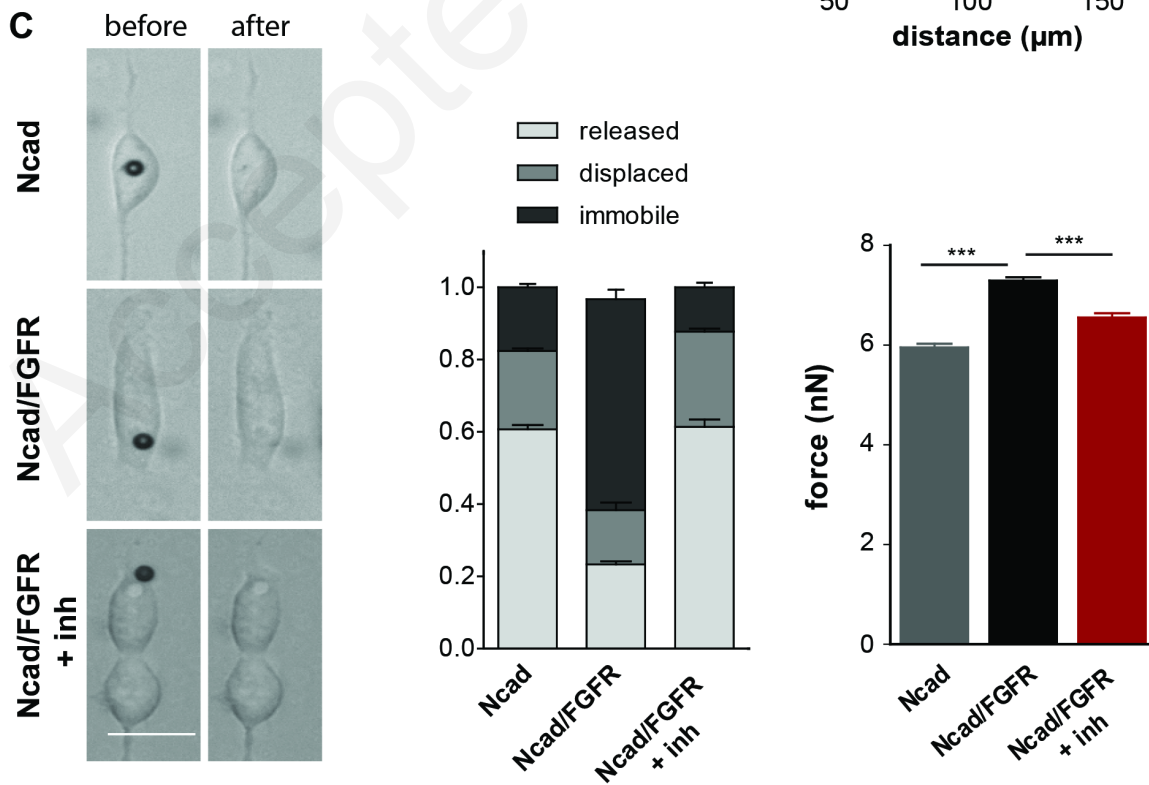
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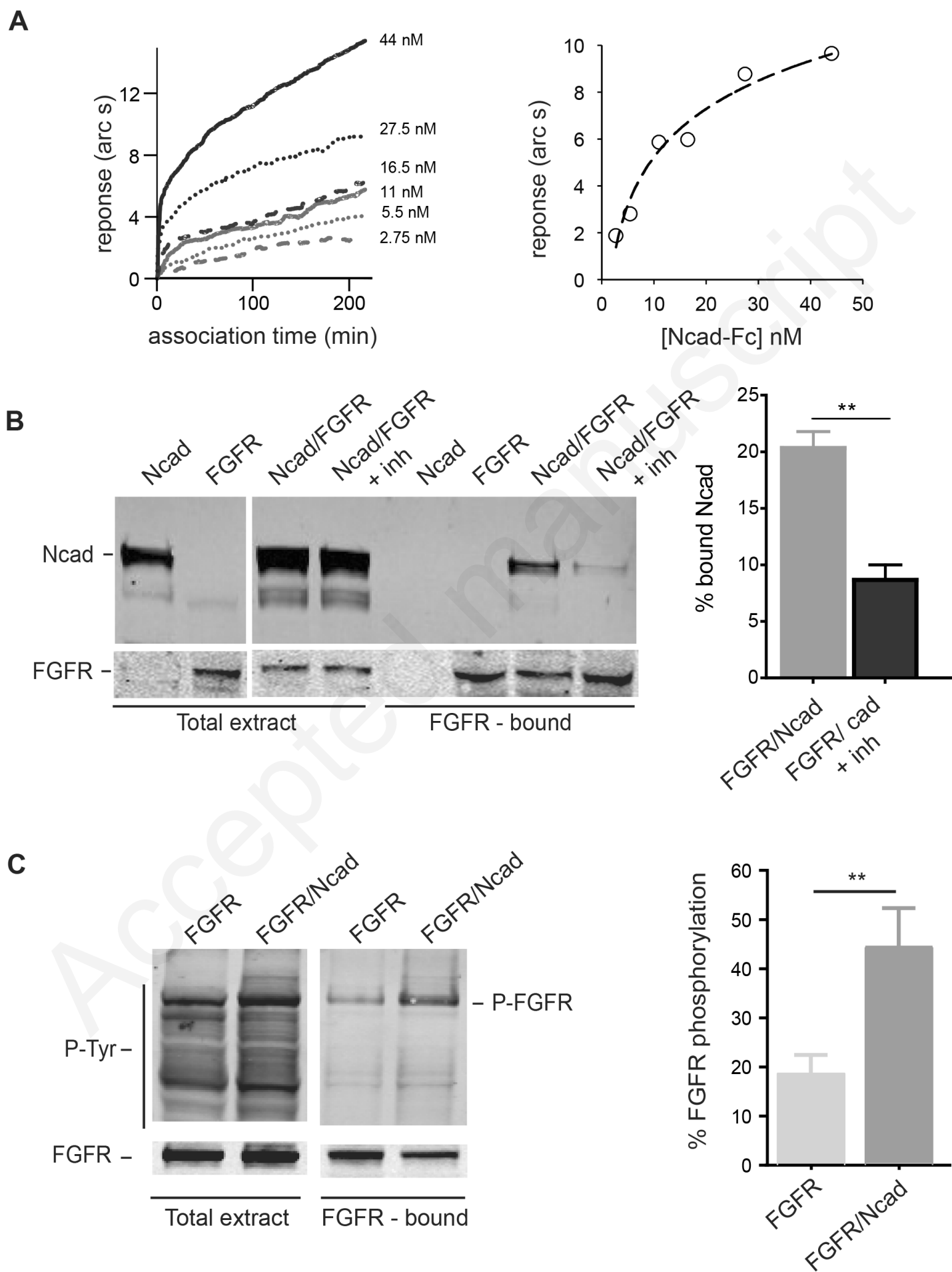
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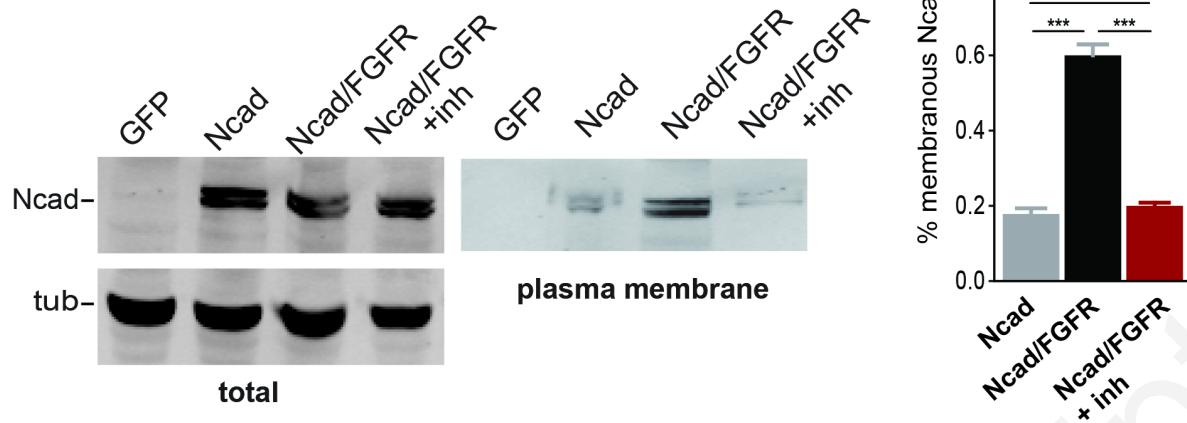
**Figure 4**



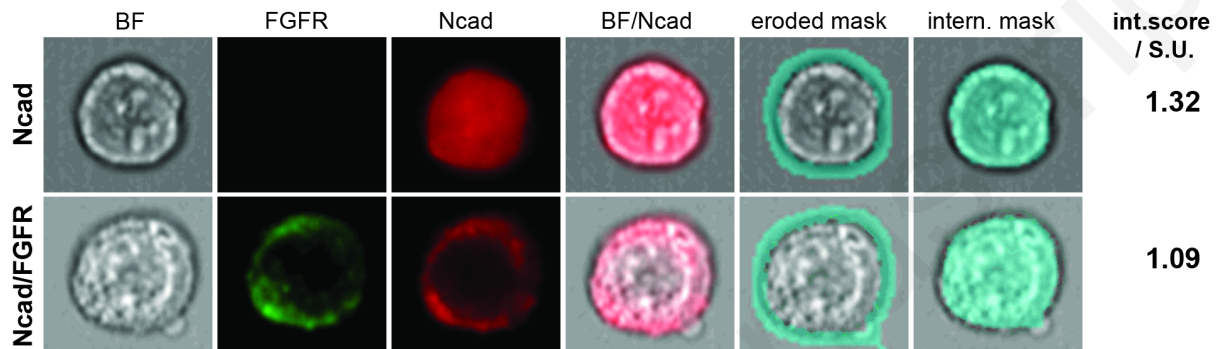


# Figure 5

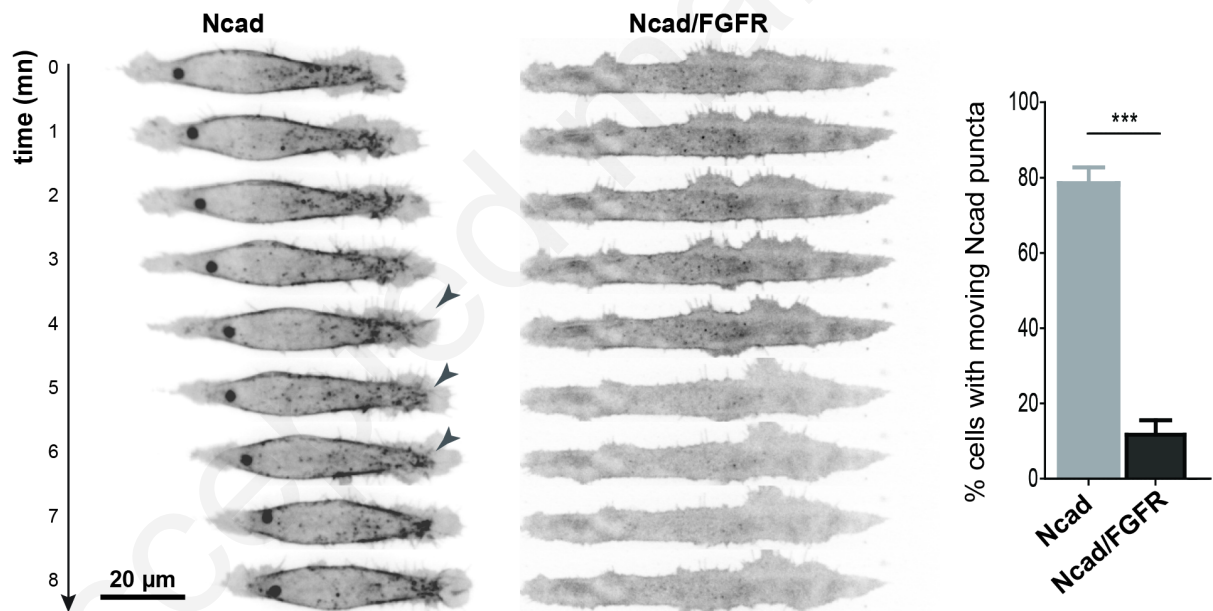
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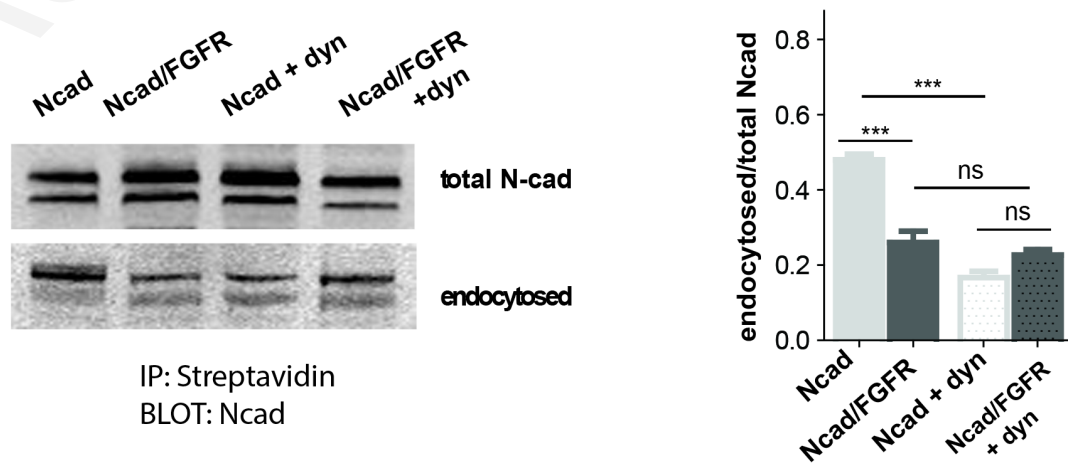
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**C**

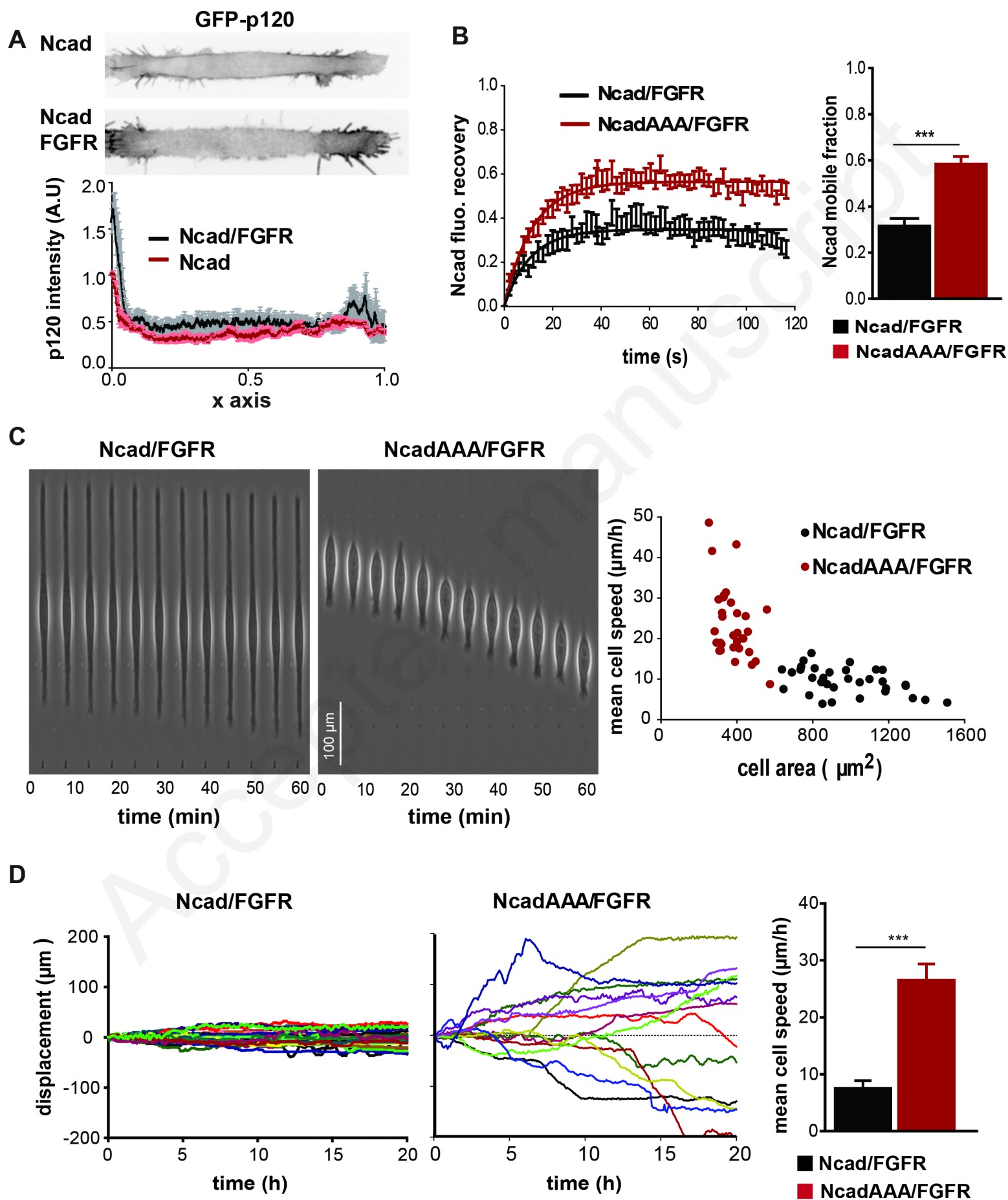


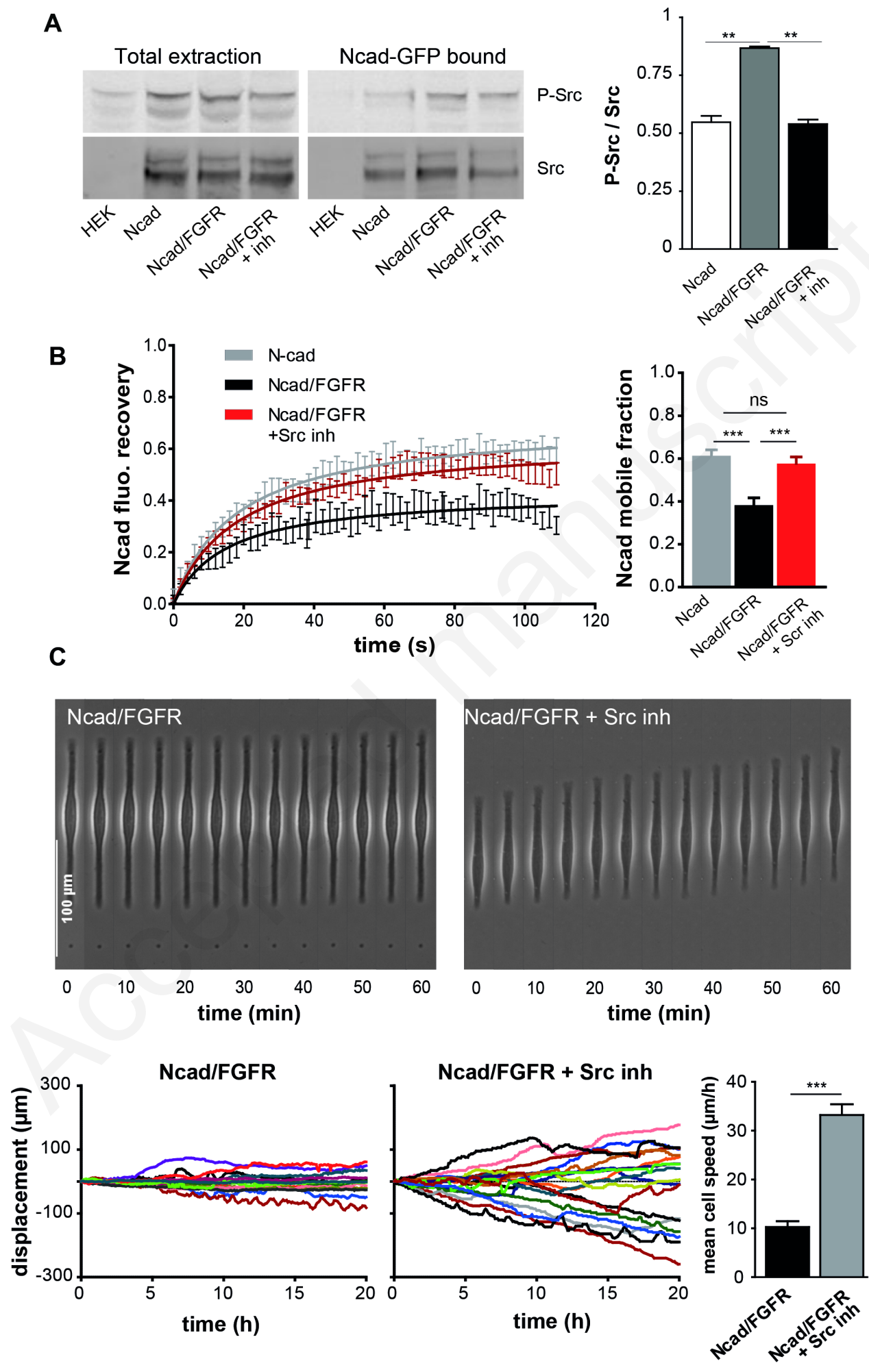
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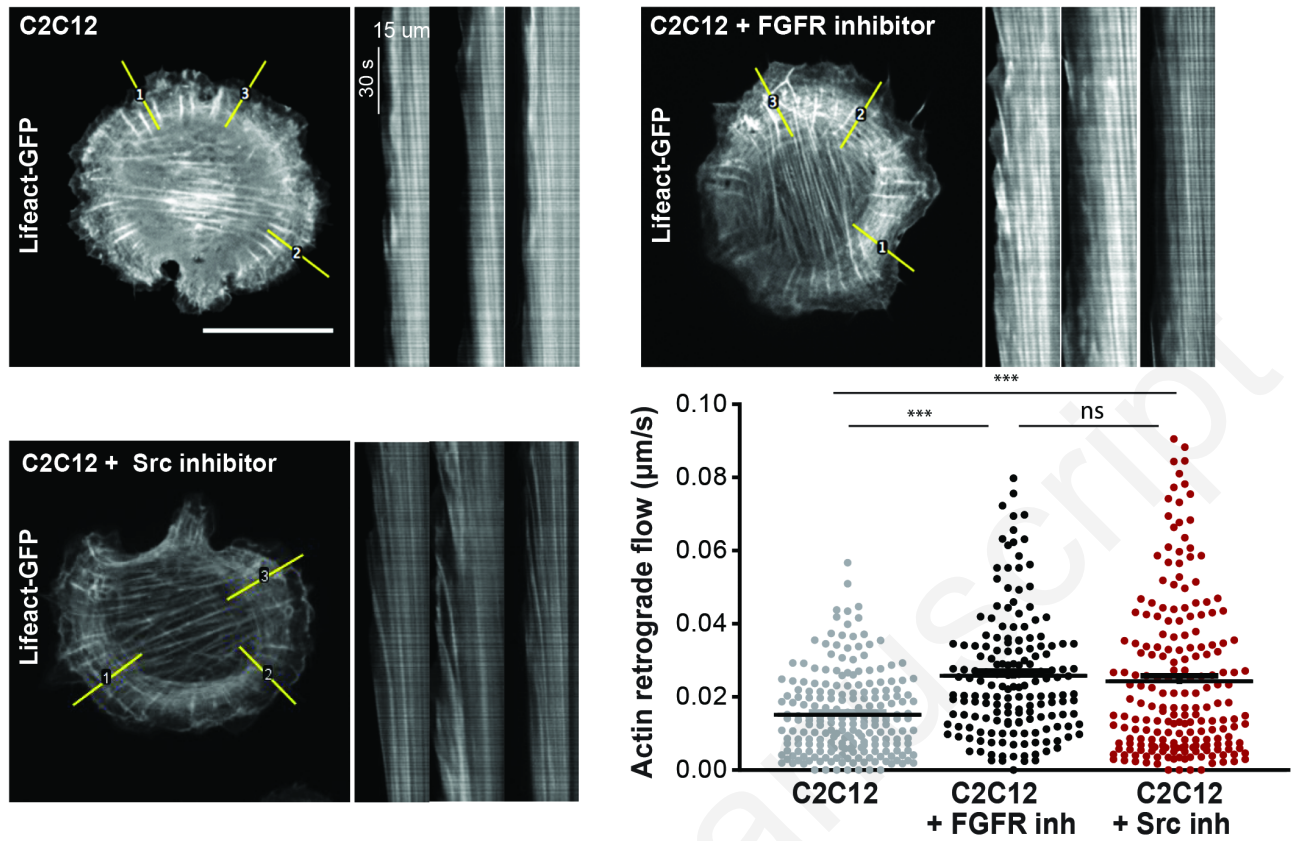
**Figure 6**



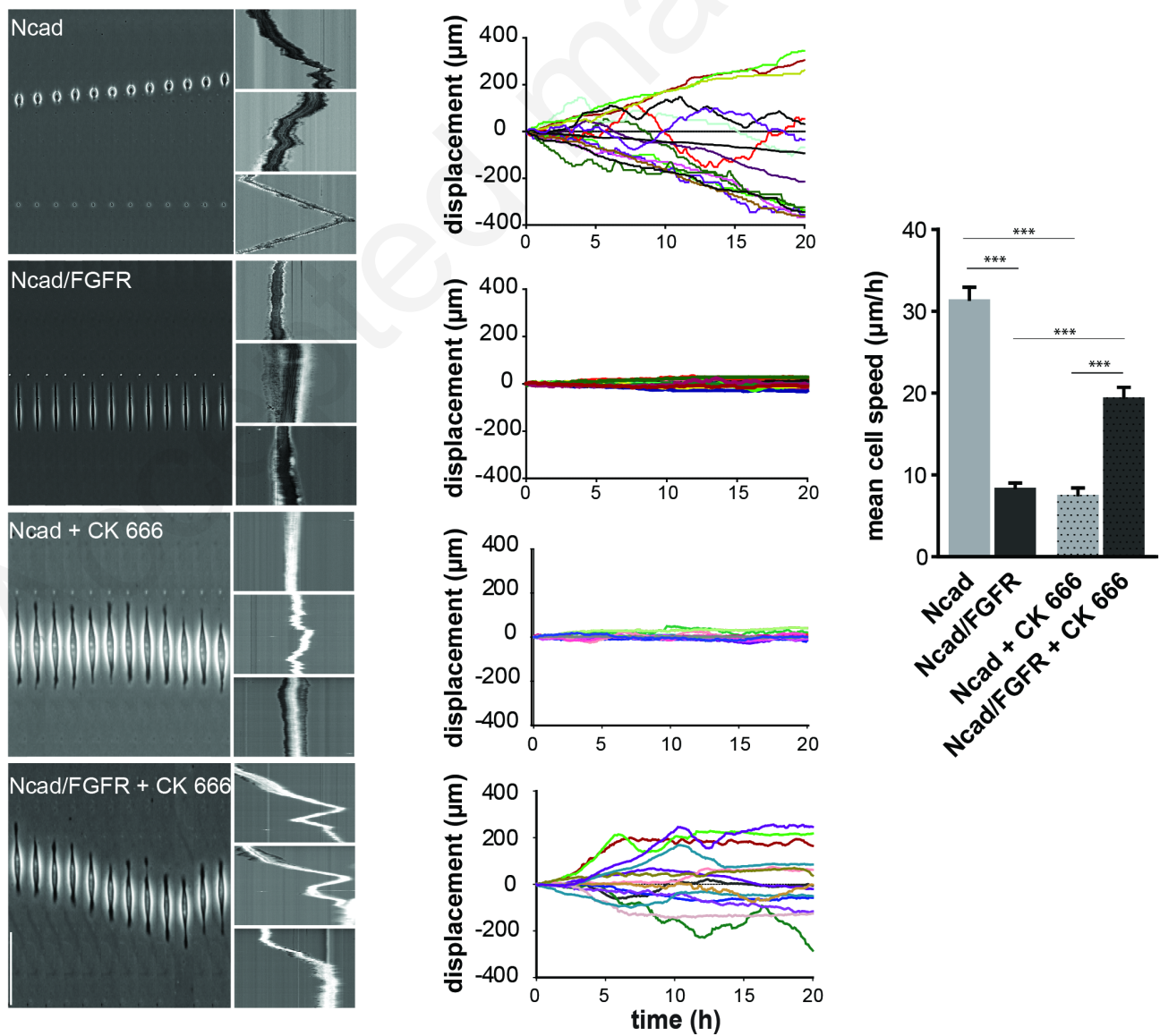
**Figure 7**

# Figure 8

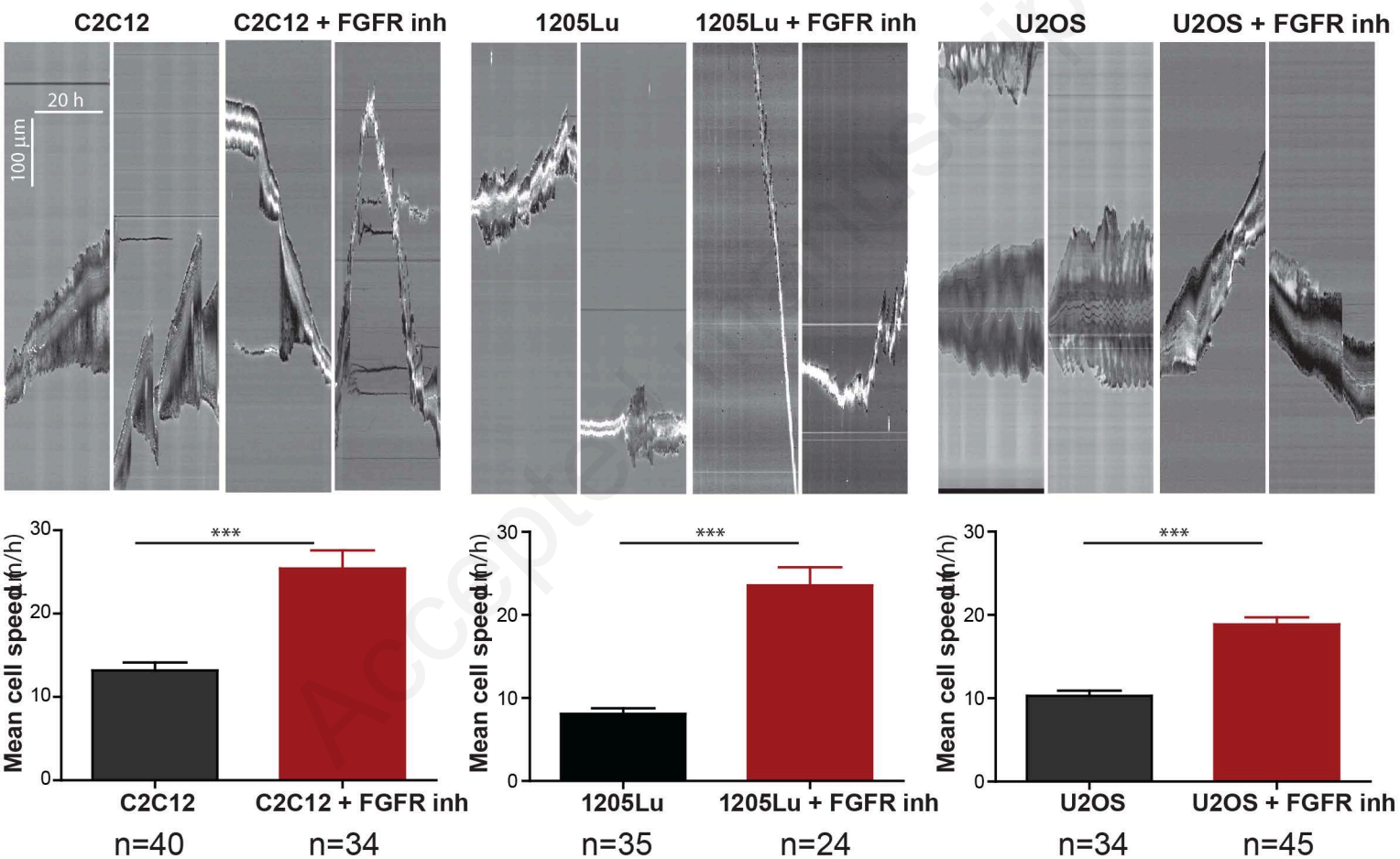
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**B**



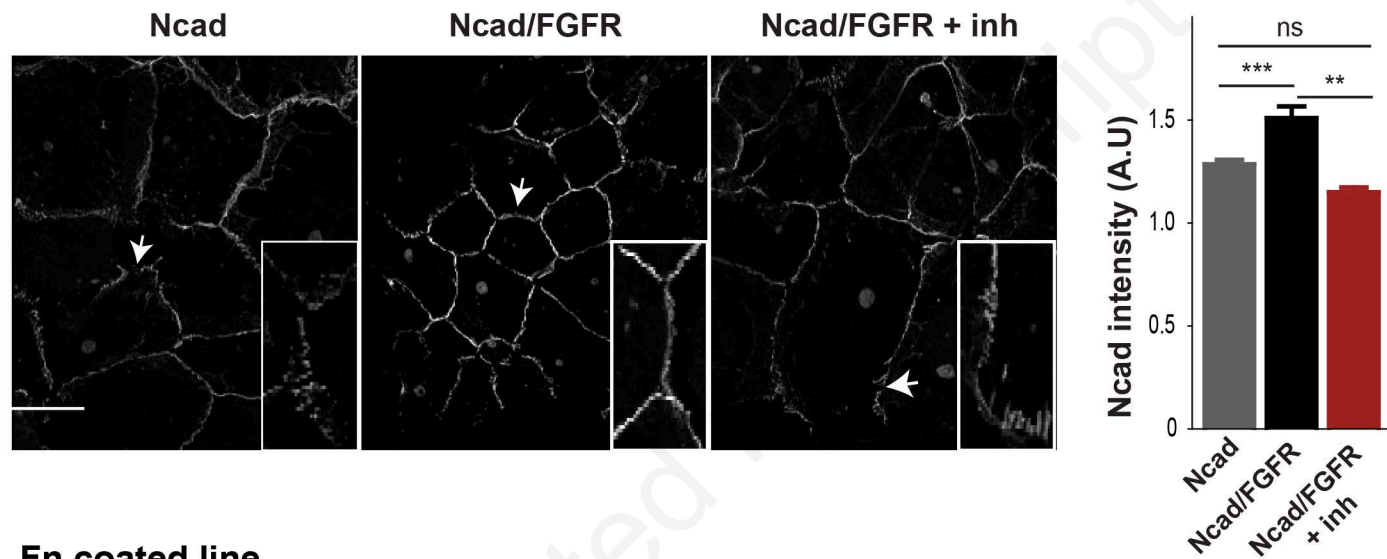
**Figure S1**





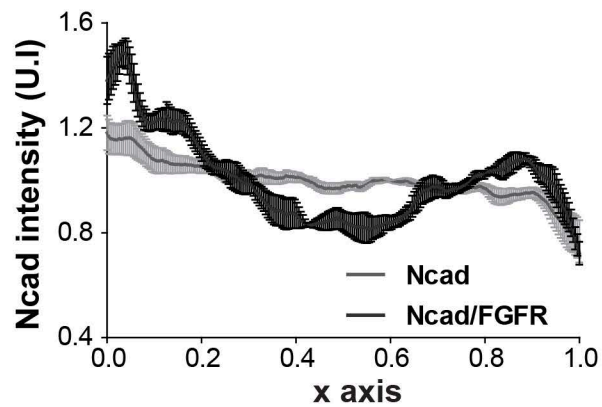
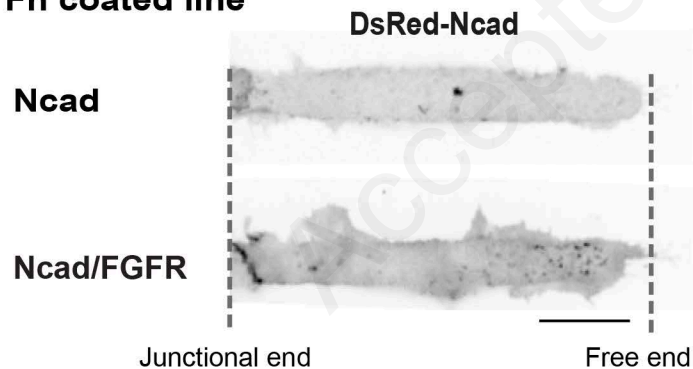
**Figure S2**

**A**



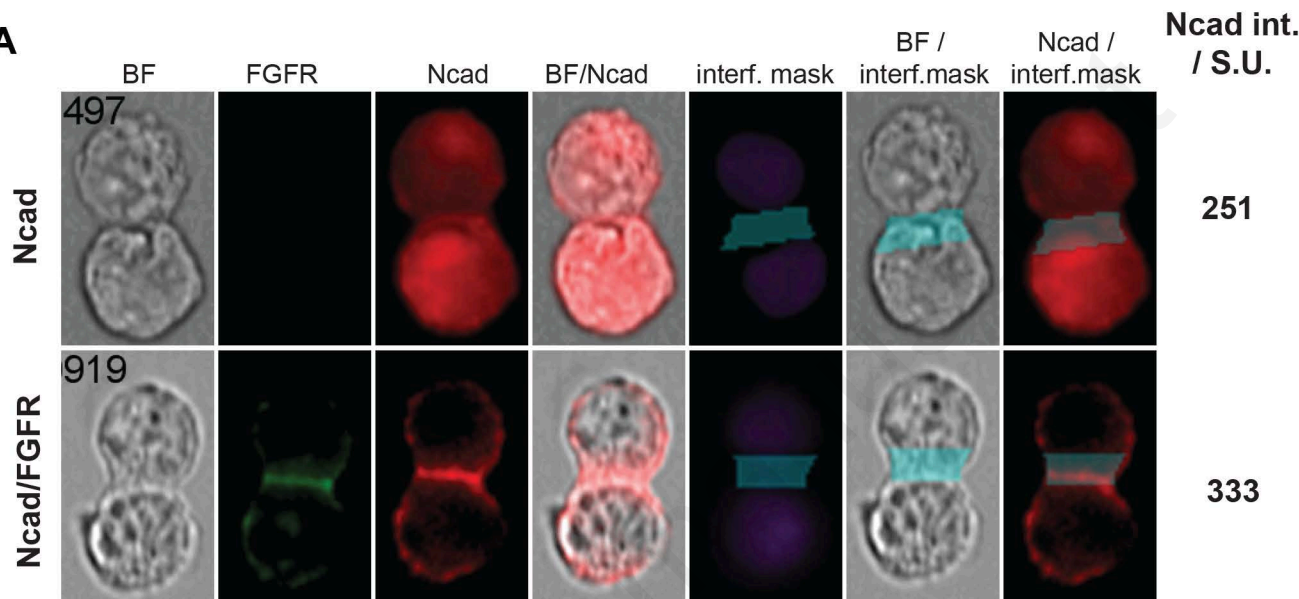
**B**

**Fn coated line**

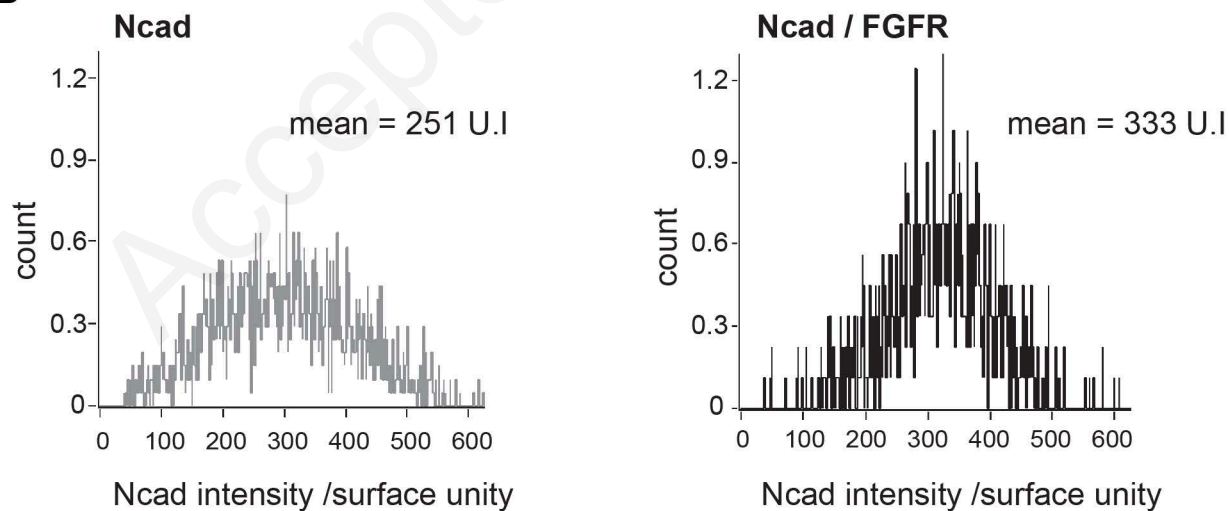


**Figure S3**

**A**



**B**

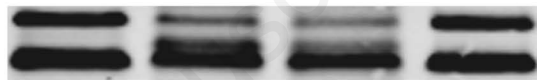


**Figure S4**

**A**

+ FGF2	—	—	+	+
+ FGFR inh.	—	+	—	+

Erk1/2



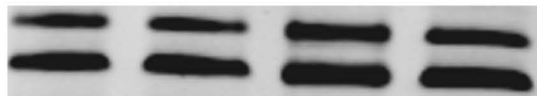
P-Erk1/2



**B**

+ Ca <sup>2+</sup>	—	—	+	+
+ FGFR inh.	—	+	—	+

Erk1/2

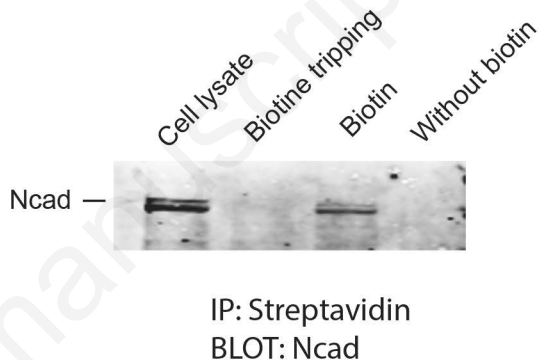
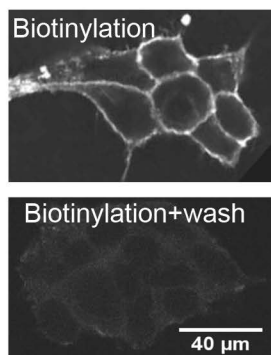


P-Erk1/2

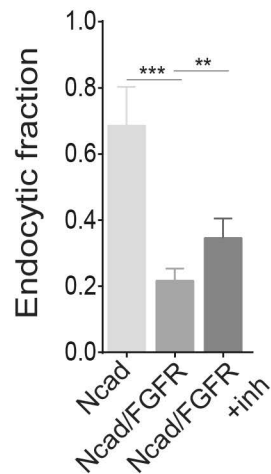
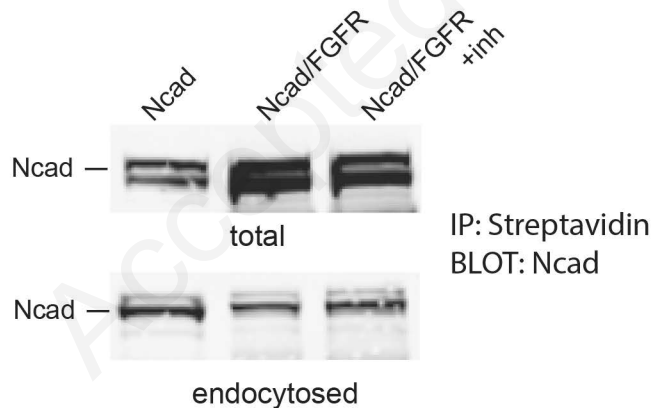


**Figure S5**

**A**

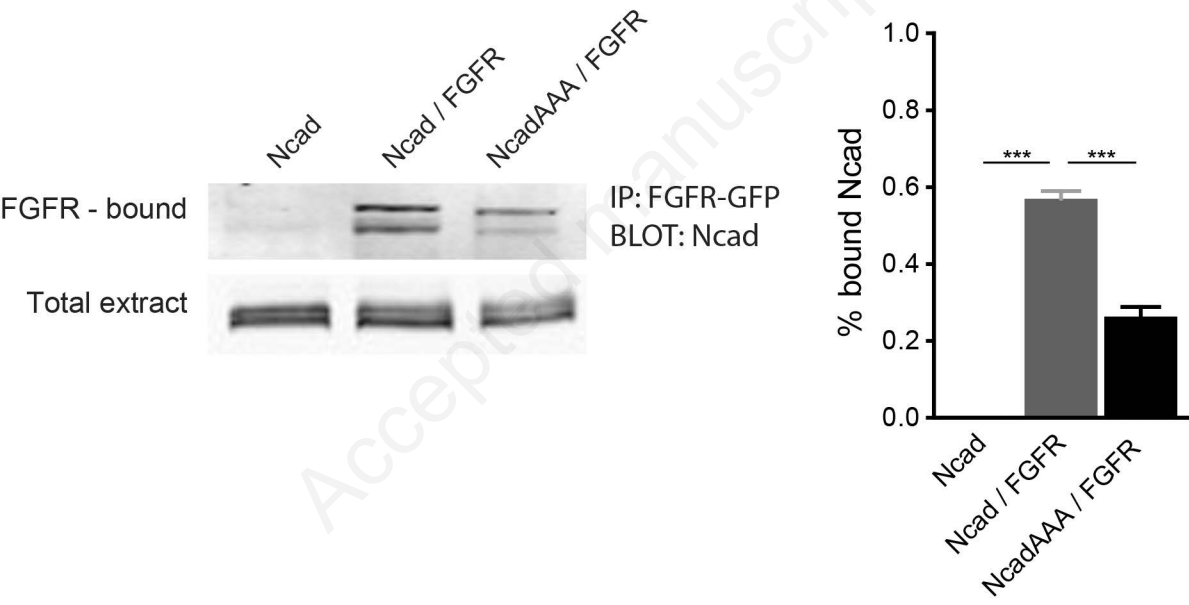


**B**

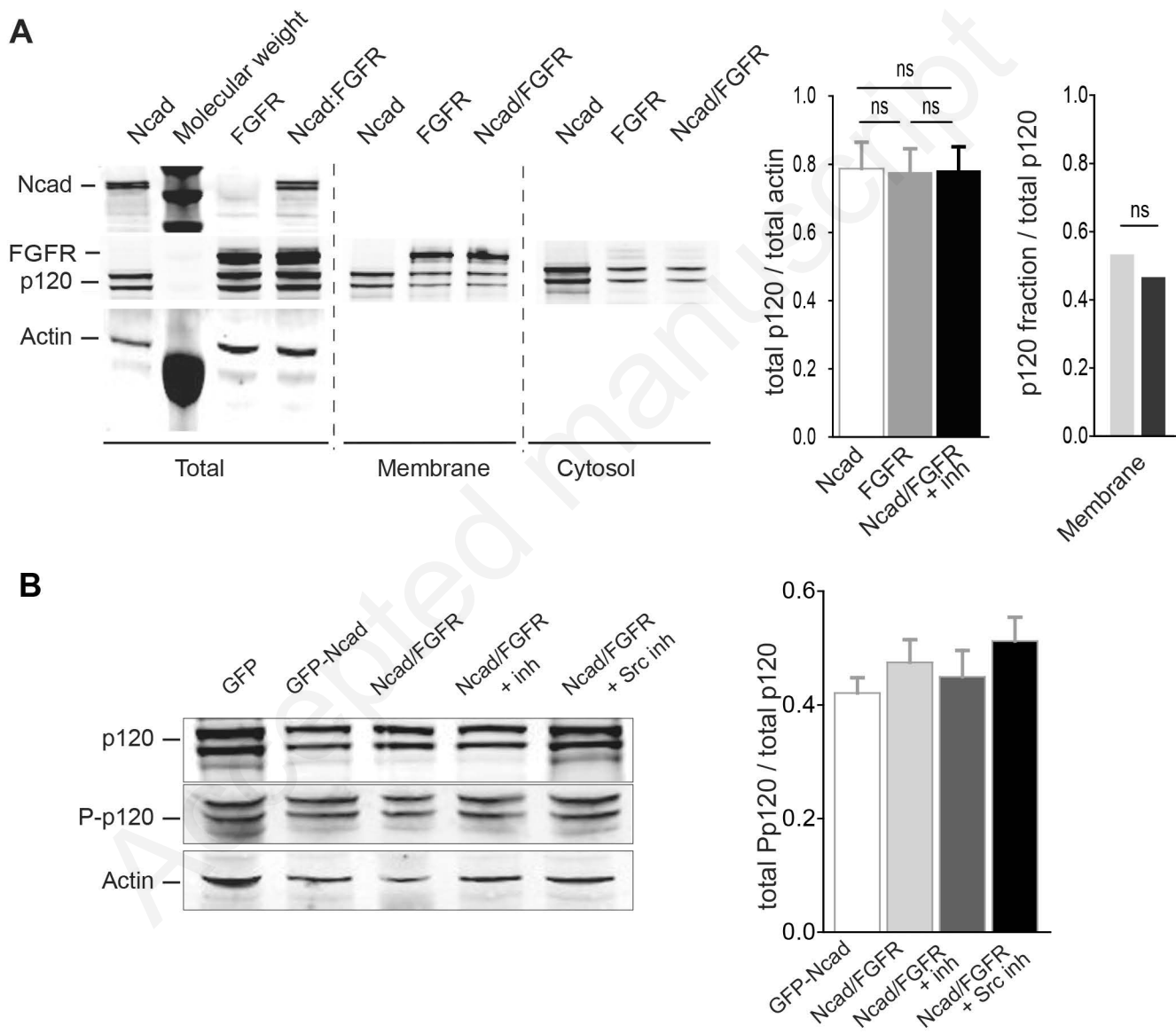




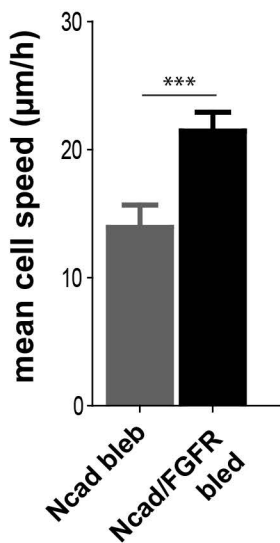
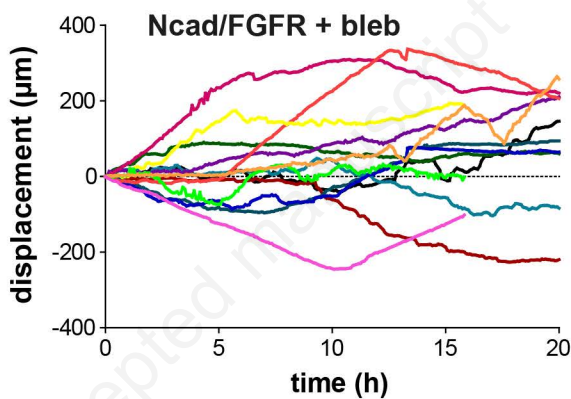
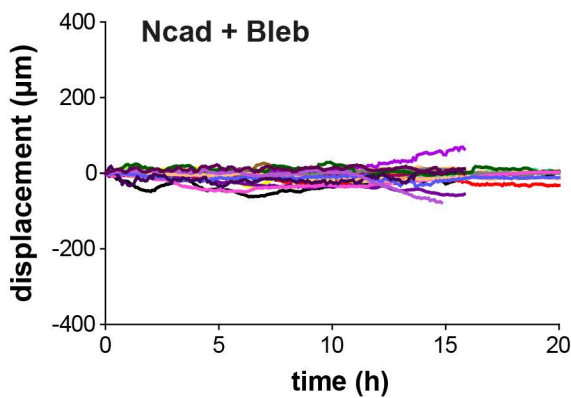
# Figure S6



**Figure S7**



**Figure S8**



## Supplemental Figure Legends

**Figure S1: FGFR inhibitor treatment increases the migration of C2C12, 1205Lu and U2 OS cells on N-cadherin coated lines.** cells were seeded at low density on 10  $\mu$ m-width Ncad-Fc coated lines in the absence or in the presence of FGFR inhibitor (+ FGFR inh) and imaged in phase contrast for 24 hours. Upper panels: Representative kymographs of the displacement of two cells for each condition. Lower panels: Histograms representing the mean cell speed C2C12, 1205Lu and U2OS cells, in the absence and in the presence of the inhibitor, respectively. \*\*\*  $p \leq 0.0001$ , ANOVA multi-comparison test, Newman-Keuls post-test).

**Figure S2: FGFR1 expression promotes N-cadherin recruitment and strengthens cell-cell contacts.** (A) DsRed-Ncad distribution in fixed monolayers of Ncad, Ncad/FGFR cells and Ncad/FGFR+inh cells grown overnight on glass coverslips in the presence of serum (fibronectin/vitronectin coating). Scale bar: 20  $\mu$ m. Boxes show zoomed views of cell-cell contacts indicated by arrows. Cell-cell contacts in Ncad/FGFR cells appear straighter than those in Ncad cells and Ncad/FGFR+inh cells. Histograms on the right show DsRed-Ncad intensities measured at the cell-cell contacts in the three conditions thanks to Imaris. \*\*\*  $p \leq 0.0001$ ; \*\*  $p < 0.01$ ; ns: non-significant, ANOVA multi-comparison test, ( $n \geq 50$ ). (B) Ncad and Ncad/FGFR cells were seeded on 10  $\mu$ m width fibronectin-coated stripes and fixed after 2 hours. Left: cell doublets were imaged (scale bar: 20  $\mu$ m). Right: the graphs show the mean distribution of DeRed-N-cad intensity along the cell width (z axis) normalised along the x axis of the cell with 0 value defined as the junctional edge of the cell and 1 value as the free edge, for Ncad and Ncad/FGFR cells ( $n = 30$  and 27 doublets, respectively, SD = 0.177)).

**Figure S3: FGFR enhances the recruitment and the clustering of N-cadherin at cell-cell contacts.** (A) Analysis by flow cytometry imaging of Ncad recruitment at cell-cell interface in cell doublets. DsRed-Ncad recruitment at cell-cell contacts was quantified as the average normalised Ncad fluorescence intensity per surface unit in cell-cell areas (Ncad int./S.U.). FGFR expression significantly increased Ncad recruitment at cell-cell contacts. Data acquisition was performed for  $1.5 \times 10^5$  cells for each condition and repeated 4 times. (B) Distribution plot of values obtained for the normalised Ncad fluorescence intensity per surface unit at cell-cell areas (Ncad int./S.U.) presented in Figure S2A.

**Figure S4: Enhanced N-cadherin engagement sustains activation of FGFR downstream pathway.** C2C12 cells were treated with FGF2 (5 nM, 15 minutes) or/and with FGFR inhibitor (10 nM, 1 hour) (**A**); or preincubated with EGTA (2 mM, 30 minutes), then switch back to medium containing 2 mM  $\text{Ca}^{2+}$  for 10 minutes in the absence or in the presence of FGFR inhibitor (**B**). Cells were lysed and total extractions were subjected to electrophoresis and Western blotting using anti-Erk1/2 and anti-P-Erk1/2 antibodies.

**Figure S5: FGFR decreases N-cadherin internalisation.** **A.** Left: Fluorescent imaging of cell surface protein biotinylation; Cy5-conjugated streptavidin labelling of freshly biotinylated cells and of biotinylated cells following reducing wash. Surface biotinylated Ncad cells were lysed and incubated with streptavidin. Total lysate of biotinylated cells (Cell lysate), streptavidin precipitated lysates of biotinylated cells submitted immediately after to a reducing wash at 4°C (Biotin + stripping), of biotinylated cells (Biotin) and of non-biotinylated cells (Without biotin) were resolved by SDS-PAGE and immunoblotted using anti-Ncad antibodies. **B.** Analysis of N-cad endocytic fraction following cell surface biotinylation for Ncad, Ncad/FGFR and Ncad/FGFR + inh cells. Experiments were performed similarly than in Figure 6B except that the hydroxy-dynasore treatment was replaced by FGFR inhibitor treatment. (Left) Western blot detection of N-cad in anti-N-cadherin (total) and streptavidin pooled (endocytosed) proteins. (Right) Quantification of the N-cad endocytosed/total ratio obtained over 3 independent western blots. \*\* $p \leq 0.01$ ; \*\*\*  $p \leq 0.001$ ; ns: non-significant, ANOVA multiple comparison test.

**Figure S6: p120 is involved in the binding of N-cadherin to FGFR.** Analysis of N-cad binding to FGFR for Ncad, Ncad/FGFR and NcadAAA/FGFR cells. Experiments were performed similarly than in Figure 4B. (Left) Western blot detection of N-cad using anti-N-cadherin antibodies in total cell lysate (total) and FGFR-GFP immunoprecipitated proteins (FGFR-bound). (Right) Quantification of the FGFR-bound/total ratio obtained over 3 independent western blots. \*\* $p \leq 0.01$ ; \*\*\*  $p \leq 0.001$ ; ns: non-significant, ANOVA multiple comparison test.

**Figure S7: FGFR expression reduces the total amount and cytosolic fraction of N-cadherin and p120.** (A) FGFR, Ncad and Ncad/FGFR HEK cells were collected without detergent. Then total, cytosolic and membranous fractions were then separated as detailed in Material and Methods, the proteins extracted and immunoblotted for N-cad, FGFR and p120 for total extracts and FGFR and p120 only for the subcellular fractions. Histograms present the quantification of total level of p120 reported to actin, and membrane-associated p120 in membranous fractions versus total p120 cellular content, respectively. ns: non-significant, unpaired t test, n = 3 experiments. (B) Total protein extracts of GFP-HEK, Ncad, Ncad/FGFR and Ncad/FGFR cells treated with Src or FGFR inhibitor were separated and immunoblotted using anti-Pp120 and anti-p120 antibodies. Actin was used for protein loading control. The histogram shows the ratio of Pp120 over p120 in total extract, determined from the quantification of 3 independent immunoblots then converted to percentage.

**Figure S8: Effect of blebbistatin treatment on the migration of cells on N-cadherin lines.** Tracked displacements over 20 hours of Ncad (n = 12), Ncad/FGFR cells (n = 13). Histogram represents the mean cell speed.

## **Video legends**

**Video 1** (1mn55s): Ncad, Ncad/FGFR, Ncad/FGFR + FGFR inh cells migration on Ncad coated lines.

**Video 2** (12s): Magnetic tweezer experiments on Ncad, Ncad/FGFR, Ncad/FGFR + FGFR inh cells.

**Video 3** (48s): Ncad trafficking at the leading edge of Ncad, Ncad/FGFR migrating cells.

**Video 4** (1mn55s): Ncad/FGFR and NcadAAA/FGFR cells migration on Ncad coated lines.

**Video 5** (1mn55s): Ncad/FGFR and Ncad/FGFR+ Src inhibitor migration on Ncad coated lines.

**Video 6** (43s): Measurement of Ncad/actin mechanocoupling in C2C12 cells.

**Video 7** (1mn55s): Ncad/FGFR and Ncad/FGFR+ CK666 migration cells on Ncad coated lines.

## Supplemental Material and Methods

### *Plasmid constructions*

The construct encoding GFP-FGFR1 (FGFR1 tagged with GFP at its carboxy-terminal extremity) was obtained using as a template pMIRB-FGFR1-Myc plasmid (gift from D. Ornitz, University of Washington), which encodes for the mouse fgfr1-IIIc full length sequence. By performing polymerase chain reactions (PCR) using sets of appropriate primers, (i) HindIII restriction site was introduced at 5' extremity (5'-GCGAAGCTTACCATGTGGGGCTGGAAGTGCC-3') while, (ii) stop codon was abolished, and AgeI restriction site was introduced at 3' extremity (5'-GCGACCGGTGGGCGCCGTTTGAGTCCACTGTT-3') of the FGFR1 encoding sequence. Resulting PCR product was subcloned into the pEGFP-N1 vector using the HindIII and AgeI restriction sites. The constructs encoding Flag-FGFR1 (FGFR1 tagged with a flag tag at its carboxy-terminal extremity) was obtained using also as a template pMIRB-FGFR1-Myc plasmid, by performing polymerase chain reactions (PCR). Using sets of appropriate primers, (i) NheI restriction site was introduced at 5' extremity (5'-GCGGCTAGCACCATGTGGGGCTGGAAGTGCC-3') while, (ii) a Flag tag encoding sequence and PmeI restriction site was introduced at 3' extremity (5'-CGCGTTTAAACTCACTTATCGTCGTCATCCTTGTAATCGGCGGCCCCGCGCCGTTTGAGTCCACTGTT-3') of the FGFR1 encoding sequence. Resulting PCR product was subcloned into the pcDNA3.1 hygro(-) vector using the NheI and PmeI restriction sites.

### *Cell culture and transfection*

HEK 293 (Human Embryonic Kidney) and C2C12 mouse myoblastic, U2OS human osteosarcoma and 1205Lu human metastatic melanoma (gift of Lionel Larue, Inst. Curie, Paris) cells were grown in DMEM medium supplemented with 10% foetal bovine serum (FBS), 2 mM L-Glutamine, 100 IU of penicillin, 100 µg/mL streptomycin at 37°C in the presence of 5% CO<sub>2</sub>. HEK cells were transiently electroporated with the plasmids encoding for dsRed-fused wild type N-cadherin (dsRed-Ncad), or N-cadherin 3A mutated in the p120 binding site (dsRed-NcadAAA)<sup>1, 4</sup> and/or with a plasmid coding for GFP- or Flag- FGFR1. Electroporation was performed with the Amaxa Cell Line Nucleofector (kit V, program X-032). To generate dsRed-Ncad, GFP-FGFR, dsRed-Ncad/GFP-FGFR dsR-Ncad/Flag-FGFR stable HEK cell lines, transfected cells were grown under a selection pressure of 200 µg/mL of Hygromycin B, 1 mg/mL of Geneticin or both. Drug resistant cells were then sorted out by



FACS (Influx 500 Cytopeia/BD-Biosciences) subcloned and further maintained with half of concentration of antibiotic pressure.

### ***Protein extraction and co-immunoprecipitation***

Proteins were extracted from  $5-6 \times 10^6$  cells. Cell cultures were rinsed in ice-cold PBS, detached with non-enzymatic detaching solution (Cell Dissociation Solution Non-enzymatic 1x, Sigma) and centrifuged at 200 rcf for 7 minutes. Cell pellets were suspended in lysis buffer (10 mM Tris HCl pH 7.5; 150 mM NaCl; 0.5 mM EDTA; 0.5% Triton) on ice. Cells were then passed slowly 10 times through a 26 gauge needle and left on ice for 1 hour with extensive pipetting every 10 minutes. Lysates were cleared by centrifugation at 20000 rcf for 10 min at 4°C. GFP-tagged proteins were then immunoprecipitated using magnetic GFP-Trap®-M beads accordingly to manufacturer instructions (Chromotek). Briefly, 25 µl of GFP-Trap®-M beads were washed 3 times with the wash buffer (10 mM Tris HCl pH 7.5, 150 mM NaCl, 0.5 mM EDTA). Fifteen µl of beads were added to 300 µl of protein extracts diluted in lysis buffer and tumbled end-over-end for 1 hour at 4°C. Beads were then magnetically separated, washed 3 times with the wash buffer, suspended in 100 µl 2x sample buffer plus reducing agent (NuPAGE, Invitrogen) and boiled at 95°C for 10 minutes to recover bound proteins. Proteins from the input and bound proteins were subjected to SDS-PAGE electrophoresis using Bis-Trisacrylamide 4%-12% NuPAGE gels (Invitrogen) then transferred on nitrocellulose membranes (0.45 µm, GE Healthcare). Membranes were blocked with 5% nonfat milk and incubated with the adequate primary antibodies and then with IRDye-coupled secondary antibodies (Rockland). The membranes were scanned using Odyssey Imaging System (LY-COR Biosciences).

### ***Surface biotinylation and endocytosis***

Cell cultures were chilled down to 4°C by three washes with cold PBS/Mg<sup>2+</sup>/Ca<sup>2+</sup> then labelled with 1 mg/ml of NHS-SS biotin (Pierce) in PBS/Mg<sup>2+</sup>/Ca<sup>2+</sup> for 12 minutes at 4°C under gentle rocking. Biotinylation was stopped by adding PBS/Mg<sup>2+</sup>/Ca<sup>2+</sup>, 50 mM glycine, 0.5% BSA at 4°C. Cells were washed twice in cold PBS/Mg<sup>2+</sup>/Ca<sup>2+</sup> and lysed. Biotinylated plasma membrane proteins were then separated by precipitation with streptavidin-coated magnetic beads (Pierce).

For endocytosis analysis, biotinylated cells were returned at 37°C for 40 minutes. Cells were then chilled down with cold PBS/Mg<sup>2+</sup>/Ca<sup>2+</sup> and bound biotin remaining at the cell surface was cleaved by incubating with 50 mM Glutathione in 75 mM NaCl, 10 mM EDTA,

pH 7.5 for 15 minutes under gentle agitation. Cells were lysed and protein extracts were subjected to precipitation with streptavidin-coated beads or with GFP-Trap®-M beads (Chromotek).

### ***Cell Fractionation***

Cells cultures were washed with ice-cold PBS then scraped in 500 µl of detergent-free buffer (20 mM Tris-Cl pH 7.4, 100 mM NaCl, 2 mM MgOAc, 5 mM KCl, 10 mM GTP + protease inhibitor cocktail- both added fresh). Cells were dounced 35 times with a 26-gauge needle on ice and centrifuged at 450 rcf, 10mn. Supernatants were collected and centrifuged at 20000 rcf for 30 minutes. The supernatants, (cytosolic fraction) were collected. The pellets (membrane fraction) were rinsed with 1ml of lysis buffer then spun at 20000 rcf for 30 minutes. The pellets were re-suspended in 180 µl of detergent-free buffer supplemented with 1% NP-40 and incubated on ice for 1 hour while mixing every 10 minutes. The fractions were resolved by SDS-PAGE electrophoresis and immunoblotting.

### ***Fixed cell imaging***

Cell cultures were fixed at room temperature in PBS 4 % formaldehyde for 15 minutes. Preparations were then mounted in Mowiol, 90% glycerol. Images were acquired with a Leica TCS SP5 inverted confocal microscope AOBS tandem, equipped with a 63x oil objective (N.A=1.4), controlled by LAS AF (Leica System).

### ***Cell-cell contact disruption assay***

Glass or plastic surfaces were microcontact printed with silicon stamps bearing 200 µm fibronectin-coated squares as described in <sup>6</sup>. Cells were seeded on the patterned surfaces in the presence of 10 µg/mL mitomycin. After 1 h, unattached cells and mitomycin were washed out and preparations were returned to the incubator overnight. Preparation were processed for live image directly after addition of 5 mM of EGTA solution or fixed after 15 minutes of EGTA treatment. Live images were acquired at 20 x objective, every 30 seconds for 30 minutes under a controlled environment (37°C, 5% CO<sub>2</sub>, type Inverted Olympus IX81, camera CoolSnap HQ<sup>2</sup>) using MetaMorph. Fixed samples were acquired with the same microscope and camera, using 20 x and 60 x objectives.

### ***Cell doublets on fibronectin-coated line assay***

Stable dsR-Ncad and dsR-Ncad/Flag-FGFR HEK cell lines were transiently transfected with GFP-p120 or lifeact-GFP and seeded on fibronectine-coated-10  $\mu\text{m}$  width lines<sup>6</sup>, and let adhere for 1-2 hours. Samples were gently washed and returned to the incubator overnight. cell doublets were chosen to image in red-phenol-free DMEM supplemented with 10% serum every minutes during 1 hour under a controlled temperature and CO<sub>2</sub> environment (37°C, 5% CO<sub>2</sub>, 40x oil objectives (N.A = 1.4), Spinning disk CSU22). Localisations of fluorescent proteins relative to the junction end were analysed using ImageJ for mask creating and Matlab for intensity calculation. All the fluorescence images were background-subtracted before quantification. The cells shape were detected by segmenting the fluorescence intensity image using Otsu method and converted into binary mask images with values outside the cell set to zero. The cell lengths were normalised to unity in the strip direction (x direction). For each individual cell, the fluorescence intensities within the cell mask along the x direction were averaged in the y direction (perpendicular to the strip direction) and projected in the x direction. The average intensity curves were normalised by the whole cell intensity and plotted against the normalised distance to the junction end. The average intensities in the x direction from multiple cells with the same experimental condition were calculated and an average curve was then created using Matlab function smooth by filtering with locally quadratic regression using a moving window of size 5. The overall behaviour of each group of multiple cells was then represented by one single curve.

### ***Ncad-Fc line guided cell migration***

Patterned silicon microcontact stamps bearing 10  $\mu\text{m}$  width lines spaced of 70  $\mu\text{m}$  were prepared by soft lithography according to a protocol derived from <sup>6</sup>. Patterned stamps were incubated with 1  $\mu\text{g}/\text{cm}^2$  anti-human IgG (Jackson ImmunoResearch), pressed on non-culture treated petri dishes or on cleaned glass coverslip previously activated by deep UV (Jelight, 4 X 60W, 15 minutes). Microcontact printed surfaces were then passivated by incubation for 1 hour with 1% Pluronic F-127 (Sigma) diluted in distilled water, followed by 3 washes with PBS. Surfaces were incubated with 1 $\mu\text{g}/\text{cm}^2$  hNcad-Fc (R&D) for 2 hours at room temperature then washed three times with PBS. Cells in culture were then dissociated on non-enzymatic detaching solution (Cell Dissociation Solution Non-enzymatic 1x, Sigma), seeded (10<sup>5</sup> cells/200 $\mu\text{l}/\text{cm}^2$ ) on these arrays of Ncad-Fc-coated lines and allowed to adhere for 1-2 hours in culture medium containing 1  $\mu\text{g}/\text{mL}$  of mitomycin, before non-adhesive cells were gently washed off. Cells were imaged live or fixed 18 hours after seeding. For live imaging, Images were acquired with a 10 X objective, every 6 minutes during 24 hours under

controlled environment (37° C, 5% CO<sub>2</sub>, Biostation Nikon). Manual tracking of individual cells was performed with the MTrackJ plugin. Individual trajectories were positioned on an orthonormal axis with the coordinates of the cells at  $t_0 = (0:0)$ . The displacements and mean cell speed were then extracted for each condition and plotted versus time and cell area, respectively.

### ***Fluorescence Recovery after Photobleaching***

Dual wavelength fluorescence recoveries after photobleaching (FRAP) was performed at 37°C on stably transfected Ncad, FGFR or Ncad/FGFR expressing cells, as well as on transiently transfected Ecad and Ecad/FGFR expressing cells. FRAP was performed using a Leica TCS SP5 confocal microscope equipped with a 40 X immersion objective (N.A=1.4) and carried out by setting the double scanning mode at 560 nm for dsRed and 480 nm for GFP and the image format to 256 x 256 pixels. After 3 prebleach scans (0.347 sec), a 20 x 40 µm ROI over the cell-cell contact was bleached with laser at full power by performing repeated scans. Recovery was recorded by imaging with low laser power every 0.347 sec (20 scans) then every 2 sec (20 scans) and finally every 10 sec (20 scans). The normalised recovery of fluorescence was expressed as a ratio of prebleach fluorescence after correction for photobleaching as reported previously<sup>3</sup>. Normalised fluorescence recovery in function of time curves were fitted with a one-term exponential equation using GraphPad Prism 5.01 software (one-phase decay non-linear regression functions), allowing to extract a plateau value representing the fraction of diffusion-limited molecules (mobile fraction) and a recovery half-time ( $t_{1/2}$ ) as a proxy of the apparent diffusion coefficient of diffusion-limited molecules<sup>5</sup>.

### ***Flow Cytometry***

Cells were detached using non-enzymatic detaching solution (Cell Dissociation Solution Non-enzymatic 1x, Sigma), centrifuged at 200 rcf during 4 minutes, resuspended in culture medium and returned to the incubator for 10 minutes favouring moderate cell-cell adhesion in suspension. Cells were centrifuged again at 200 rcf for 4 minutes, fixed in 4% formaldehyde for 15 minutes, washed 3 times in PBS, incubated in the presence of 0.1% Dapi in PBS-BSA 0.1 % for 5 minutes, washed again then imaged under flow using ImageStream X (Amnis, Proteigene) set with the 405, 488, 560-nm laser and 480-560 filter. Data were analysed using the IDEAS software (Amnis, Proteigene) focusing on singlets for the quantifications of internal pool and on doublets for cell-cell accumulation of DsRed-Ncad fluorescence. For internal fluorescence quantification, regions corresponding to the total cell

surface, the internal cell area and the cell membrane area were extracted from bright field images of singlets. Briefly, the morphology mask was applied to bright field images. Then, 4 pixels were evenly eroded from the border of the mask in order to exclude the cell membrane from the mask. The resulting mask was applied to the fluorescence channel. The internalisation feature was then applied to the final mask in order to calculate the internalisation score. Internalisation score per surface unit was defined as the ratio of internal fluorescence intensity per surface unit over the overall intensity per surface unit in the whole cell expressed on a logarithmic range.

The N-cadherin recruitment at the cell-cell interface was determined on doublet populations. Regions corresponding to the total cell surface and the cell-cell interface were extracted from bright field images and Dapi staining, respectively. The 4 pixels interface mask was determined as a region centred at the dimmest pixel between the 2 nuclei (Dapi). The interface mask was applied to the bright field channel to determine the surface area of the cell-cell contacts in the doublets, then to the fluorescence channel to count the intensity of Ncad staining at the cell-cell contact. Results were expressed as fluorescence intensity per surface unit.

### ***Production and biotinylation of soluble FGFR1 extracellular domain***

The soluble FGFR1 extracellular domain (FGFR1-ST) used here corresponds to amino acids 120 to 368 of the FGFR1IIIc comprising the acid box and immunoglobulin loops D2 and D3 with a poly-histidine-tail sequence followed by a thrombin cleavage site at the N-terminus and a Factor Xa cleavage site followed by a Strep-Tag II sequence at the C-terminus and is termed FGFR1-ST. This protein was produced in CHO cells and is heavily N-glycosylated <sup>2</sup>. For biotinylation, 2  $\mu\text{L}$  of sodium periodate at 0.5 M freshly resuspended in water was added to 28  $\mu\text{L}$  of purified FGFR1-ST at 16  $\mu\text{M}$  in PBS supplemented by 0.01% of tween (PBS-T). The mixture was left incubated 1 hour in the dark. Excess sodium periodate was then removed using nanosep 30 kDa centrifugal device following the manufacturer recommendation and using PBS-T as exchange buffer. Purified periodated FGFR1-ST was recovered in 50  $\mu\text{L}$  of PBS-T and 1.5  $\mu\text{L}$  of biotin-LC-hydrazide at 135 mM in DMSO was then added. The reaction was left incubated overnight at RT and then kept 6 hours at 4°C. Excess biotin-Lc-Hydrazide was removed using nanosep 30 kDa centrifugal device following the manufacturer recommendation and PBS-T as exchange buffer. Biotinylated FGFR1-ST was recovered in 200  $\mu\text{L}$  of PBS-T and kept at -20° c. Its concentration was estimated at 1.14  $\mu\text{M}$  by measurement of the absorbance at 280 nm ( $\epsilon_{280\text{ nm}} = 47424 \text{ mole}^{-1}.\text{L}.\text{cm}^{-1}$ ).

### ***Optical biosensor experiments***

Streptavidin (Sigma, 50  $\mu$ l at 2.5 mg/mL) was immobilised on aminosilane surfaces using bisulfosuccinimidyl suberate (BS<sup>3</sup>, Perbio, 1 mM) as the cross linker following the manufacturer's recommendations (NeoSensors, Sedgefield, UK). Surface was washed 5 times with 80  $\mu$ L of Pi buffer (10 mM sodium phosphate buffer pH 7.2) and then incubated 3 min in 2M Tris-HCl, pH 8 to stop the cross-linking reaction. Three washes with 80  $\mu$ L of Pi buffer and then of PBST were then performed before the addition of 25  $\mu$ L of biotinylated FGFR1-ST at 1.14  $\mu$ M. The reaction was left incubated for 4 to 6 hrs at RT and then overnight at 4°C. Cuvette was then cleaned by washing 3 times with 50  $\mu$ L of PBST, NaCl 2 M in Pi buffer, PBST, HCl 20 mM and then PBST again.

Binding assays were carried out in TNC buffer (Tris-HCl 20 mM pH 7.5, NaCl 150 mM, CaCl<sub>2</sub> 2 mM and Tween-20 0.02%). A single binding assay consisted of adding 0.5 to 6  $\mu$ L of purified Ncad-Fc to a cuvette containing 19 to 24.5  $\mu$ L or 24 to 29.5  $\mu$ L of TNC. The association reaction was followed until binding was at least 90% of the calculated equilibrium value, usually between 150 s and 230 s. The cuvette was then washed three times with 50  $\mu$ L TNC to initiate the dissociation of bound Ncad-Fc. Regeneration of the surface between each binding assay was performed by washing 3 times with 50  $\mu$ L 2 M NaCl in Pi buffer, TNC, 20 mM HCl, and then TNC, which removed 98% to 100% of bound Ncad-Fc. Binding parameters were calculated using the non-linear curve fitting program FASTFit (NeoSensors). Each binding assay yielded four binding parameters, which are the slope of initial rate of association, the on-rate constant ( $k_{on}$ ) and the extent of binding, all calculated from the association phase, and the off-rate constant ( $k_{off}$ , equivalent to the dissociation rate constant,  $k_{diss}$ ), calculated from the dissociation phase. Biosensor experiments were carried out 4 times on 3 different FGFR1-ST-derivatised surfaces. The determination of binding kinetics in optical biosensors was prone to second phase binding sites at high concentration of Ncad-Fc. Thus, limiting amounts of ligand were immobilised on the sensor surface, whereas the slope of initial rate,  $k_{on}$  and the extent of binding were only determined at low concentrations of ligate (3 different experiments), and  $k_{off}$  was measured at higher concentrations of ligate (2 different experiments), to avoid steric hindrance and rebinding artefacts. A single site model was used to calculate all binding parameters. The dissociation constant ( $K_D$ ) was calculated both from the ratio of the  $k_{diss}$  and  $k_{ass}$  and from the extent of binding, to provide an estimate of the self-consistency of the results.

### ***Magnetic tweezers***

The magnetic microneedle device was made of a 5 cm long stainless steel sewing needle glued to the top of permanent neodymium iron boron (NeFeB) surface surrounded by an aluminium rod. The montage was assembled on a micromanipulator (MP-285, Sutter Instrument) at a 30°C vertical angle, and the tip initially aligned at 600 µm from the centre of the observation field. The whole device was mounted on an inverted microscope (Olympus IX 81) equipped with a 40 x phase contrast air objective and a CCD camera (CoolSnap HQ<sup>2</sup>) operating in the burst mode (frequency of 15 frames/s for 2 minutes). The micromanipulator allowed translational movement across all three axes with nanometer precision to position the magnetic field in the vicinity of beads. The distance between the tip of needle and detached bead was measured with imageJ.

The force applied to the bead is decreasing exponentially with the distance to the magnetic rod. To calibrate the magnetic force with is a function of the distance between the needle tip and the bead, 4.5 µm beads were placed in a 100% polyethylene glycol solution (Mn 700, Sigma) at various distances of the needle and the bead motion was tracked by video microscopy. The instantaneous horizontal bead velocity ( $v$ ) was extracted using ImageJ tracking. The force applied on the bead ( $F$ ) was calculated respecting Stokes equation:  $v = F/6\pi\eta r$ , where  $\eta$  is the dynamic fluid viscosity (for PEG Mn700:  $\eta = 25$  Pa.s at 25°C) and  $r$  is the radius of the bead. The calibration was performed ten times and the forces versus distance data were regressed to an exponential equation.

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