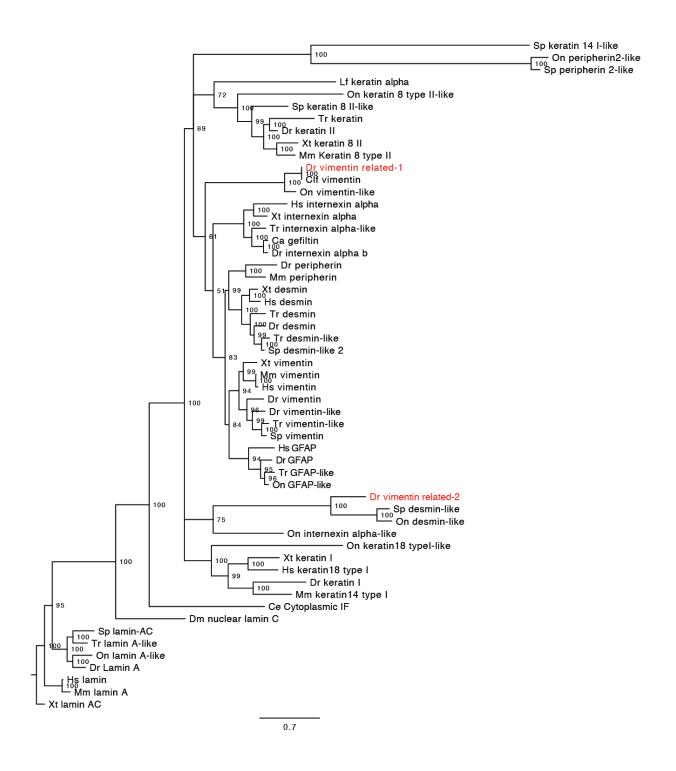


Goto et al. Supplemental Figure 1

Supplemental Figure 1. Time course expression of vimentin related genes

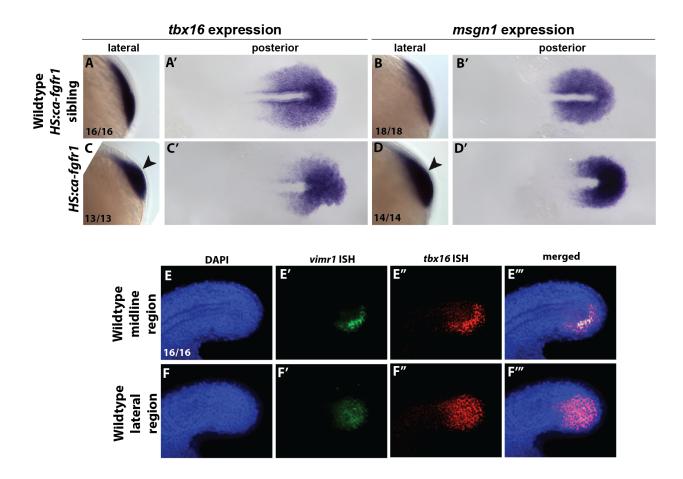
Vimr1 (A-J) and Vimr2 (K-T) expression in wildtype embryos fixed at different stages. The beginning of notochord expression at the 18-somite stage is indicated by arrowheads (G and Q). Shield staged embryos were taken laterally with the shield towards the right. Embryos from 8-somite stage and onward were taken laterally with anterior towards the bottom.



Goto et al. Supplemental Figure 2

Supplemental Figure 2. Phylogenetic analysis of intermediate filaments

Bayesian phylogenetic analysis of zebrafish vimentin-like genes with other intermediate filaments with laminin genes serving as an outgroup. The numbers at the node specify Bayesian posterior probabilities. Species abbreviations with common names in parentheses: Ca, Carassius auratus (goldfish); Ce, Caenorhabditis elegans (nematode); Clf, Canis lupus familiaris (Dog); Dm, Drosophila melanogaster (fruit fly); Dr, Danio rerio (Zebrafish); Hs, Homo sapiens (human); Lf, Lampetra fluviatilis (Lamprey); Mm, Mus musculus (mouse); On, Oreochromis niloticus (Nile tilapia); Sp, Stegastes partitus (bicolor damselfish); Tr, Takifugu rubripes (pufferfish); Xt, Xenopus tropicalis (frog)



Goto et al. Supplemental Figure 3

Supplemental Figure 3. Expression of mesodermal maturation genes in FGF overactivated tailbud and expression domains of *vimr1* and *tbx16* in the wildtype tailbud. HS:*cafgfr1* and wildtype embryos were heat-shocked at bud stage and fixed at 8-somite stage to examine the effect of FGF overactivation on maturation gene expression. Overactivation of FGF leads to strong expression of *tbx16* and *msgn1*, which expands into the NMP region (C, C', D, D') compared to the control (A, A', B, B'). Double fluorescent *in situ* hybridization of *tbx16* and *vimr1* ina wild-type 20-somite stage embryo. As the mesodermal progenitors move from the dorsal region to ventral region, the cells turn on *vimr1* and *tbx16* (E, E', E'', E'''). However, as the cells enter the maturation zone in the lateral regions, the cells turn of *vimr1* (F, F', F'', F''').

Table S1. Statistics for time lapse tracking. P values were calculated by Student's t-test. S indicates significant (p<0.05), while NS implies nonsignificant (p>0.05).

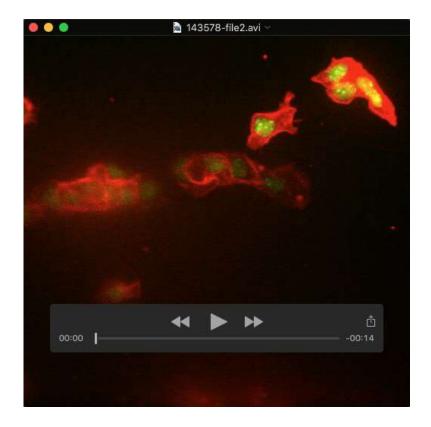
			Displace	ment Mean Track Speed		Track straightness		
Embryo	number of embryos	Total tracks	P val compared to wildtype- DMSO	P val compared to HStbx16- SU5402	P val compared to wildtype- DMSO	P val compared to HStbx16- SU5402	P val compared to wildtype- DMSO	P val compared to HStbx16- SU5402
wildtype-DMSO	4	268						
HS:tbx16-DMSO	3	177	p=0.7425; NS	p=0.433; NS	p=0.0323; S	p=0.2539; NS	p=0.1601; NS	p=0.9717; NS
wildtype-SU5402	3	143	p=0.0323; S	p=0.0073; S	p=0.0073; S	p=0.0151; S	p=0.1886; NS	p=0.1116; NS
HS:tbx16-SU5402	3	223						
HS:caFGFr1-DMSO	3	169	p=0.5225; NS		p=0.0065; S		p=0.3332; NS	

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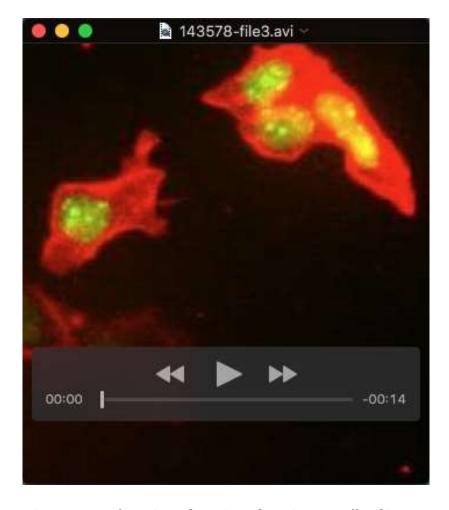
Table S2. Raw data of transplanted lineage analysis and statistics. P values were calculated by

Fisher's exact test. S indicates significant (p<0.05), while NS denotes nonsignificant (p>0.05).

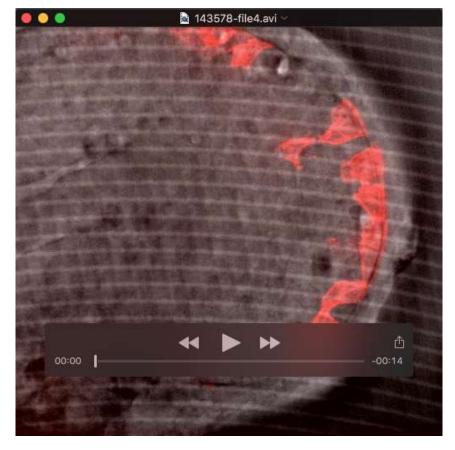
	number of	Total	Total	% .	% .	P val compared to	P val compared to
Embryo	embryos	muscle cells	nonmuscle cells	muscle	nonmuscle	wildtype	dnfgf
wildtype	10	167	68	71.064	28.936		
HS:dnfgfr1	10	33	127	20.625	79.375	p<0.0001; S	
pmsgn1/wildtype	9	234	88	72.671	27.329	p=0.7028; NS	
pmsgn1/HS:dnfgfr1	10	216	191	53.071	46.929	p<0.0001; S	p<0.0001; S
ptbx16/wildtype	8	177	96	64.835	35.165	p=0.153; NS	
ptbx16/HS:dnfgfr1	9	97	81	54.494	45.506	p=0.0006; S	p<0.0001; S



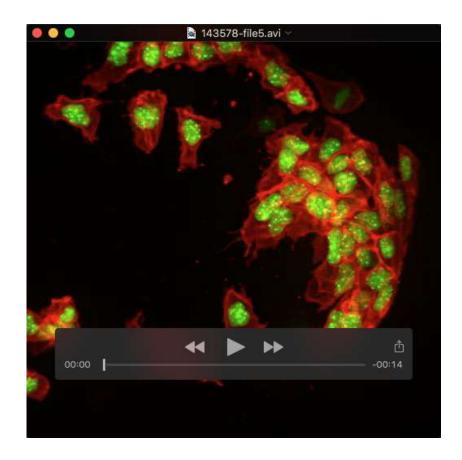
Movie 1. Time lapse movie of transplanted *HS:(CAAX)mCherry-p2a-(NLS)kikume* during the 1st EMT step. Cells of interest are in the upper right of the movie. Images were acquired every 5 minutes for 8 hours.



Movie 2. Cropped version of movie 1, focusing on cells of interest.



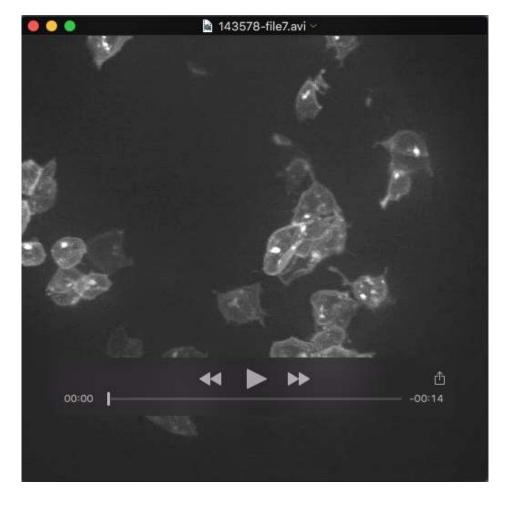
Movie 3. Time lapse movie of transplanted *HS:TCF∆C* cells. . Images were acquired every 5 minutes for 8 hours.



Movie 4. Time lapse movie of transplanted *HS:(CAAX)mCherry-p2a-(NLS)kikume* during the **2**nd **EMT step.** The cell of interest is in the lower left of the movie. . Images were acquired every 5 minutes for 8 hours.



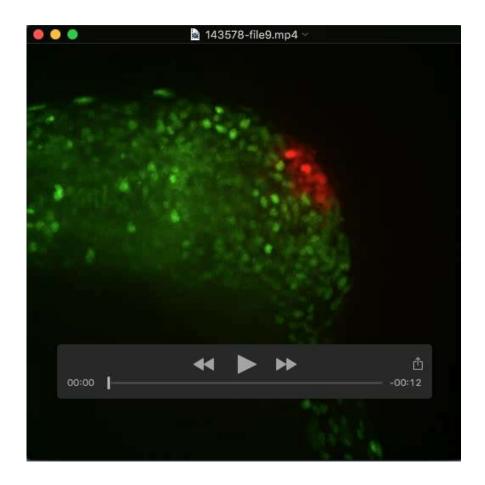
Movie 5. Cropped version of movie 4, focusing on the cell of interest.



Movie 6. Time lapse movie of transplanted *HS:dnfgfr1* **cells.** The cell of interest is in the lower left of the movie. . Images were acquired every 5 minutes for 8 hours.



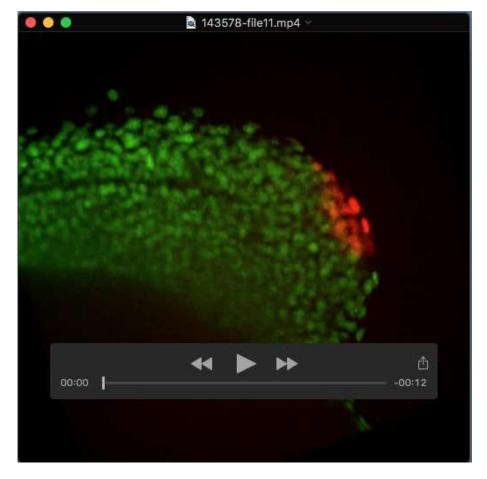
Movie 7. Cropped version of movie 6, focusing on the cell of interest.



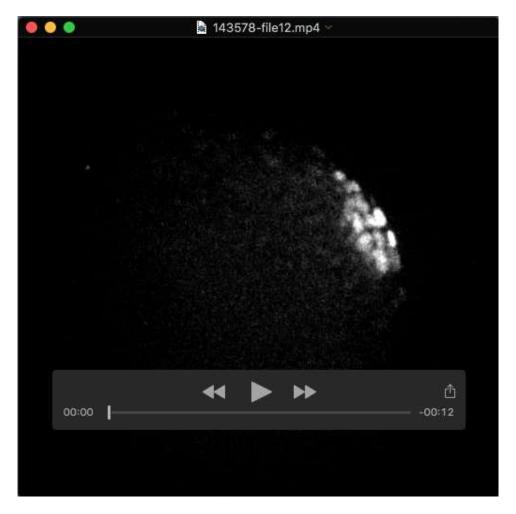
Movie 8. Time lapse movie of photoconverted cells in a wildtype tailbud (merged) A movie of wildtype photoconverted red tailbud cell with Z projection from 5um Z-stacks merged with a projection of nonconverted cells in green. The embryo was treated with DMSO (vehicle). Images were acquired every 5 minutes for 5 hours. (5 frames per second).



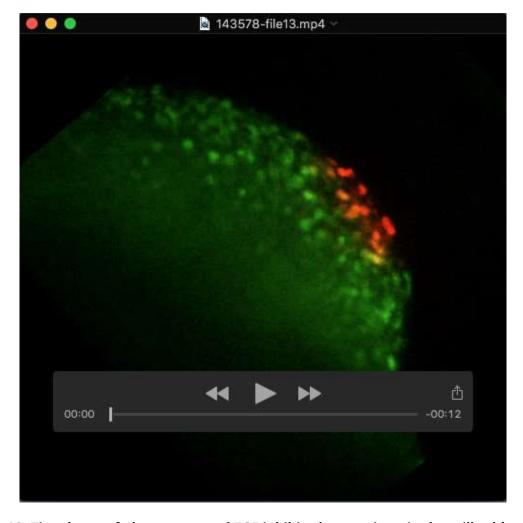
Movie 9. Time lapse movie of photoconverted cells in a wildtype tailbud (photoconverted cells only) A movie of wildtype photoconverted tailbud cell with Z projection from 5um Z-stacks. The embryo was treated with DMSO (vehicle). Images were acquired every 5 minutes for 5 hours. (5 frames per second).



Movie 10. Time lapse movie of photoconverted tbx16 overexpressed progenitors in the tailbud (merged) A movie of HS:tbx16 photoconverted tailbud cells with Z projection from 5um Z-stacks merged with unconverted green field projected plane. The embryo was treated with DMSO (vehicle). Images were acquired every 5 minutes for 5 hours. (5 frames per second).

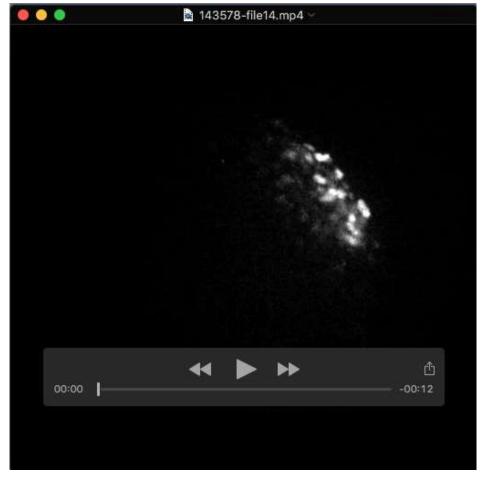


Movie 11. Time lapse movie of photoconverted tbx16 overexpressed progenitors in the tailbud (photoconverted cells only) A movie of DMSO treated HS:tbx16 photoconverted tailbud cells with Z projection from 5um Z-stacks. Images were acquired every 5 minutes for 5 hours. (5 frames per second).

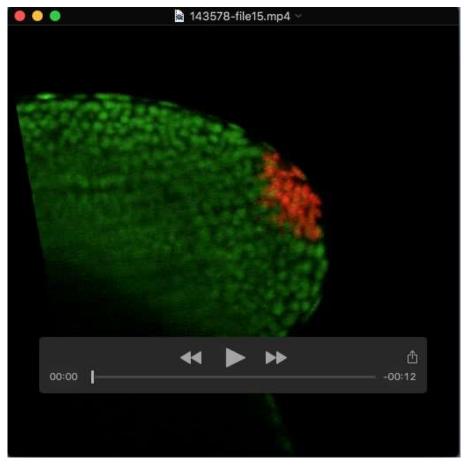


Movie 12. Time lapse of photoconverted FGF inhibited progenitors in the tailbud (merged)

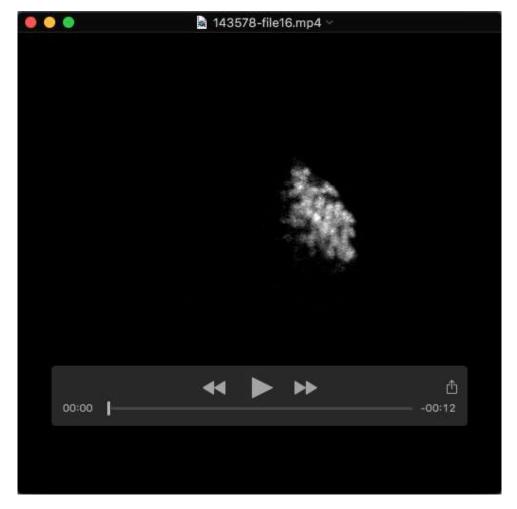
Time lapse movie of SU5402 treated photoconverted tailbud cells with Z projection from 5um Z-stacks merged with a projection of nonconverted cells in green. Images were acquired every 5 minutes for 5 hours. (5 frames per second).



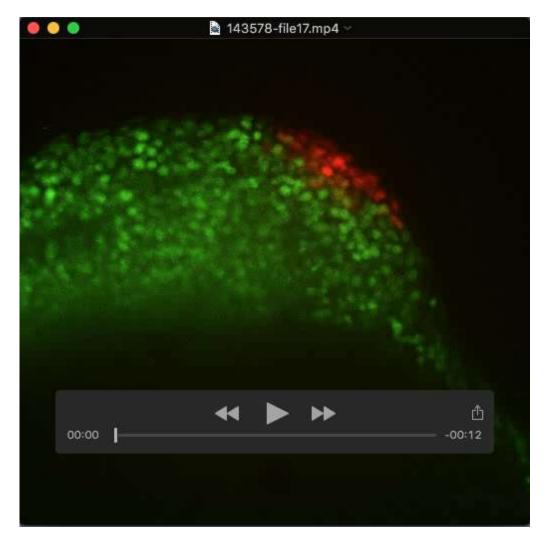
Movie 13. Time lapse of photoconverted FGF inhibited progenitors in the tailbud (photoconverted cells only) Time lapse movie of SU5402 treated photoconverted tailbud cells with Z projection from 5um Z-stacks. Images were acquired every 5 minutes for 5 hours. (5 frames per second).



Movie 14. Time lapse of photoconverted tbx16 overexpressing progenitors in the SU5402 treated tailbud (merged) Time lapse movie of SU5402 treated photoconverted *HS:tbx16* tailbud cells with Z projection from 5um Z-stacks merged with a projection of unconverted green field plane. Images were acquired every 5 minutes for 5 hours. (5 frames per second).



Movie 15. Time lapse of photoconverted tbx16 overexpressing progenitors in the SU5402 treated tailbud (photoconverted cells only) Time lapse movie of SU5402 treated photoconverted nuclei of *HS:tbx16* tailbud cells with Z projection from 5um Z-stacks. Images were acquired every 5 minutes for 5 hours. (5 frames per second).



Movie 16. Time lapse of photoconverted FGF overactivating progenitors in the tailbud (merged) Time lapse movie of DMSO treated photoconverted *HS:caFGFr1* tailbud cells with Z projection from 5um Z-stacks merged with a projection of unconverted nuclei in green. Images were acquired every 5 minutes for 5 hours. (5 frames per second).



Movie 17. Time lapse of photoconverted FGF overactivating progenitors in the tailbud (photoconverted cells only) Time lapse movie of DMSO treated photoconverted *HS:caFGFr1* tailbud cells with Z projection from 5um Z-stacks. Images were acquired every 5 minutes for 5 hours. (5 frames per second).