## **Supplemental Information**

## **Figure S1**



Figure S1. Overexpression of the other 7 miRNAs doesn't reduce the formation of both granulocytes and macrophages in zebrafish primitive myelopoiesis together. (A-L) Whole mount *in situ* hybridized embryos showing the expressions of mpx (A, D, G, J), mfap4 (B, E, H, K), and lcp (C, F, I, L) in the 26 hpf embryos microinjected with NC (A-C) or inhibitor NC (G-I) were similar to the ones microinjected with miR-129-5p mimic (D-F) or miR-722 inhibitor (J-L). Embryos were viewed laterally and positioned head left. The images in panels (A, B, C) were also used in Figure 1 (B, D, F) because these results were from one ISH experiment and shared the same negative controls. (M-N) qRT–PCR results showing the expressions of mpx (M) in the embryos microinjected with mimics of miR-7b, miR-30a, miR-125c, and miR-129-5p, or inhibitors of miR-738 were similar to their control embryos, and the expressions of mfap4 (N) in the embryos microinjected with mimics of miR-7b, miR-125c, and miR-129-5p, or miR-135a inhibitor were similar to the embryos microinjected with mimics of mix-7b, mix-125c, mix-125c, mix-125c, mix-125c, mix-125c, mix-135a inhibitor were similar to the embryos microinjected with mix-129-5p.

expressions of *mfap4* (N) in the embryos microinjected with miR-30a mimic, miR-738 inhibitor or miR-153a inhibitor were all reduced compared to their controls. The gene expression levels in NC group was normalized as 1.0 and the fold changes of gene expression levels in the embryos microinjected mimics or inhibitors were shown relative to NC group (shown in Y-axis). NC: mimic negtaive control; inhNC: inhibitor negative control; \*: P<0.05, \*\*: P<0.01.

## Figure S2



Figure S2. The amount of mature miR-210-5p was significantly increased in the embryos microinjected with miR-210-5p mimic. (A-D) Whole mount in situ hybridization results showing the expression of miR-210-5p was significantly increased in the embryos microinjected with miR-210-5p mimic (C, D) compared to the ones microinjected with NC (A, B) at 14 hpf (A, C) and 26 hpf (B, D), respectively. (E) In vitro assay showing the miR-210-5p mimic was stable after overnight incubation with RNase A at 65°C. The *in vitro* reaction was performed by incubating the dsRNA (miR-210-5p mimic)) with RNase A under the same conditions (without adding yeast tRNA) as the experiment of whole mount in situ hybridization. The 1st lane (in the leftmost) shows the DNA maker. The 2<sup>nd</sup> lane (labeling with 1) shows the miR-210-5p mimc was stable in the solution without RNase A after incubation overnight at 65°C. The 3<sup>rd</sup> lane (labeling with 2) shows the miR-210-5p mimc was stable in the solution with RNase A (final concentration: 50 µg/ml) after incubation overnight at 65°C. The 4<sup>th</sup> lane (labeling with 3) shows the miR-210-5p mimc was stable in the solution with RNase A (final concentration: 50 µg/ml) after incubation overnight at room temperature. The smaller size of miR-210-5p occurred in the 3<sup>rd</sup> and 4<sup>th</sup> lanes (labeling with 3 and 4) compared with the  $2^{nd}$  lane (labeling with 2) was due to the removal of the 2 nt 3'-overhange in each strand of the dsRNA (see Table S1) by RNase A (4 nt shorter than the "wild type" miR-210-5p mimic). (F) qRT-PCR results showing the amount of mature miR-210-5p was significantly increased in the embryos microinjected with miR-210-5p mimic compared to the ones microinjected with NC at 14 hpf and 26 hpf, respectively. The expression level of mature miR-210-5p in NC group was normalized as 1.0 and the fold change of miR-210-5p expression level in miR-210-5p mimic microinjection group was shown relative to NC group (shown in Y-axis), respectively. miR-210-5p: miR-210-5p mimic; RNase: RNase A; O/N: overnight; NC: mimic negative control; \*\*: P<0.01.

## Figure S3



Figure S3. The expressions of *foxj1b* and *slc3a2a* were significantly increased in miR-210 knockout embryos and the increased expressions were inhibited in the *gata5/6* morphants. (A-B) qRT-PCR results showing the expressions of *foxj1b* and *slc3a2a* were significantly increased in the miR-210 knockout embryos. The expression level of genes in  $miR-210^{+/+}$  embryos was normalized as 1.0 and the fold change of the gene expression levels in  $miR-210^{+/+}$  embryos was shown relative to  $miR-210^{+/+}$  embryos (shown in Y-axis), respectively. \*\*: P<0.01. (C-J) The expressions of *foxj1b* and *slc3a2a* were reduced in the *gata5/6* morphants (D, H) compared with the embryos microinjected with control MO (C, G). However, knocking out miR-210 dramatically increased the expressions of *foxj1b* and *slc3a2a* (E, I) compared to the wild type embryos (C, G). The reduced expressions of *foxj1b* and *slc3a2a* in the *gata5/6* morphants (D, H) were effectively rescued in the miR-210 knockout embryos (F, J). Embryos were *in situ* hybridized at 14 hpf and viewed dorsally (C-F) or laterally (G-J). Embryos were positioned head top (C, F) or left (G-J).

**Table S1.** Sequences of miRNA mimics, miRNA inhibitors and primers used for sgRNA synthesis, RT-PCR and qRT-PCR

Name	Sequence
miR-129-5p mimic	Guide strand: CUUUUUGCGGUCUGGGCUUGCU
	Passenger strand:CAAGCCCAGACCGCAAAAAGUU
miR-210-5p mimic	Guide strand: AGCCACUGACUAACGCACAUUG
	Passenger strand:AUGUGCGUUAGUCAGUGGCUUU
miR-7b mimic	Guide strand: UGGAAGACUUGUGAUUUUGUU
	Passenger strand:CAAAAUCACAAGUCUUCCAUU
miR-30a mimic	Guide strand: UGUAAACAUUCCCGACUGGAAG
	Passenger strand: UCCAGUCGGGAAUGUUUACAUU
miR-125c mimic	Guide strand: UCCCUGAGACCCUAACUCGUGA
	Passenger strand:ACGAGUUAGGGUCUCAGGGAUU
miR-722 inhibitor	AAUCUGAAACGUUUCUGCAAAAAA
miR-153a inhibitor	GAUCACUUUUGUGACUAUGCAA
mimic negative control	Guide strand: UUCUCCGAACGUGUCACGUTT
	Passenger strand: ACGUGACACGUUCGGAGAATT
inhibitor negative control	CAGUACUUUUGUGUAGUACAA
sgRNA1F	TAATACGACTCACTATAAGTTCCGATGGCACAGGAAAGTTTTAGAGCTA
	GAAATAGC
sgRNA2F	TAATACGACTCACTATAGACAGCGGCTAACCAGCTTTGTTTTAGAGCTA
	GAAATAGC
sgRNAR	AAAAAAGCACCGACTCGGTGCC AC
gata4-FL-F	GCTCGTGGAGAATAATCGC
gata4-FL-R	TGCGTTTATGCCAGAATCAG
gata6-FL-F	CCTCATTGTGGACCCTACC
gata6-FL-R	AATTCAGCCTCAAGATCACC
<i>foxj1b</i> -FL-F	CCGCTCAAACATACGAGTTA
foxj1b-FL-R	CGTGTATATGTCATGCGTTC
<i>slc3a2a-</i> FL-F	GTCCGCCGGCTTGAAACAGA
<i>slc3a2a-</i> FL-R	TGTGTCTAGCCCGGGAATGG
stard3-FL-F	GCTGATTCTGGGATGTGCTC
stard3-FL-R	CGATAGGTCACCGGTACGAG
<i>mpx</i> -qRT-F	GGGGCAGAAGAAGAAGTCC
<i>mpx</i> -qRT-R	CCCTTGCTAAACTCTCATCTCG
<i>lcp1-</i> qRT-F	GAAGCTCTGATCGCTCTGCT
<i>lcp1-</i> qRT-R	CCCTTGCTAAAC TCTCATCTCG
<i>mfap4-</i> qRT-F	ATGAAGAGAACGGAGGATGG
mfap4-qRT-R	CACATTCCCGAATCCTCTCT
actb1-qRT-F	TTCCTTCCTGGGTATGGAATC
actb1-qRT-R	GCACTGTGTTGGCATACAGG
tal1-qRT-F	GGAGATGCGGAACAGTATGG

tal1-qRT-R	GAAGGCACCGTTCACATTCT
lmo2-qRT-F	GGACGCAGGCTTTACTACAAAC
lmo2-qRT-R	CCGGATCCTCTTTCACAGGAA
spi1b-qRT-F	GGGCAGTTTTAACCAAAGATCA
spi1b-qRT-R	CCCAAGAGTGATCGTTCTGAC
<i>foxj1b-</i> qRT-F	TTATAGCAACGAGGACGAGCAG
foxj1b-qRT-R	ATCCCTCACAAAACGCATAGCC
rnasel3-qRT-F	TGTCAAGAAGAGCAAGTCATTTC
rnasel3-qRT-R	TTCTGCTGGTTGACCGTAAG
<i>slc3a2a-</i> qRT-F	TAAAGGTCAGTGCCGGAGA
<i>slc3a2a-</i> qRT-R	AACCAGTAAAGCCCAACGAG
stard3-qRT-F	GCTGATTCTGGGATGTGCT
stard3-qRT-R	GGCGTTTAAAGAAGCAATGG
miR-210-5P-qRT	AGCCACTGACTAACGCACATTG
foxj1b-3utr-F	ACGCATGACATATACACGCA
foxj1b-3utr-R	CTGGAATTGATTTTGCGGAG
slc3a2a-3utr-F	GTCATTATTTACTCGGCTCC
slc3a2a-3utr-R	TCTGTACAGAGATCAGCTTC
<i>slc3a2a-</i> 3UTR-F	agatcgccgtgtaattctagaGTCATTATTTACTCGGCTCC
<i>slc3a2a-</i> 3UTR-R	gccggccgccccgactctagaTCTGTACAGAGATCAGCTTC
foxj1b-3UTR-F	agatcgccgtgtaattctagaACGCATGACATATACACGCA
foxj1b-3UTR-R	gccggccgccccgactctagaCTGGAATTGATTTTGCGGAG
F-foxj1b-3UTR-MT	GCGGAAAGATCGCCGTGTAATTCTAGAacgcatgacatatacacgca
R-foxj1b-3UTR-MT	AAGCGGCCGGCCGCCCGACTCTAGAgtacagattgatcttttat
F-slc3a2a-3UTR-MT	GCGGAAAGATCGCCGTGTAATTCTAGAacacacagacactgttta
R-slc3a2a-3UTR-MT	AAGCGGCCGGCCGGCCCGACTCTAGAatcctcatttgccatttat