# nature portfolio

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## **Reporting Summary**

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

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| For all statistical a | nalyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.  |  |  |  |
|-----------------------|---|--|--|--|
| n/a Confirmed         |   |  |  |  |
| ☐ ☐ The exac          | exact sample size $(n)$ for each experimental group/condition, given as a discrete number and unit of measurement   |  |  |  |
| A statem              | ent on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly   |  |  |  |
| The stati             | stical test(s) used AND whether they are one- or two-sided mon tests should be described solely by name; describe more complex techniques in the Methods section.   |  |  |  |
| A descrip             | tion of all covariates tested   |  |  |  |
| A descrip             | tion of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons  |  |  |  |
| A full des            | A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)  |  |  |  |
|                       | hypothesis testing, the test statistic (e.g. $F$ , $t$ , $r$ ) with confidence intervals, effect sizes, degrees of freedom and $P$ value noted uses as exact values whenever suitable.  |  |  |  |
| For Baye              | sian analysis, information on the choice of priors and Markov chain Monte Carlo settings  |  |  |  |
| For hiera             | rchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes   |  |  |  |
| Estimate              | s of effect sizes (e.g. Cohen's $d$ , Pearson's $r$ ), indicating how they were calculated  |  |  |  |
|                       | Our web collection on statistics for biologists contains articles on many of the points above.  |  |  |  |
| Software ar           | nd code   |  |  |  |
| Policy information    | about <u>availability of computer code</u>  |  |  |  |
| Data collection       | Micromanager, LightCycler480 Software, Olympus FV10-ASW, Leica Application Suite X  |  |  |  |
| Data analysis         | ImageJ, R, Kyplot 6.0, Microsoft Excel, MEGA 11, GeneDoc 2.7.000  |  |  |  |
|                       | g custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information. |  |  |  |

#### Data

Policy information about availability of data

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our  $\underline{\text{policy}}$

The authors confirm that all relevant data are provided in this paper, in its Supolementary Information files, or public data stroage shown in the text.

| Research inv | olving hu | man participants, their data, or biological material |
|--------------|-----------|--|
|              |           | with   |

## **Antibodies**

Antibodies used

Anti-DIG-POD, anti-DIG-AP, anti-fluorescein, Anti-Cholecystokinin (CCK8) antibody (C2581,Sigma), DDDDK-tag antibody (PM020, MLB)

Authentication

Immunohistochemistry of ccka and cckb double KO validated the specificity of C2581.DDDDK-tag antibody was confirmed to label no cell in the WT sample.

Sense and antisense probes validated the specificity of antibodies used in ISH.

### Eukaryotic cell lines

Policy information about cell lines and Sex and Gender in Research

Cell line source(s) HeLa cells from Riken BRC Cell Bank (Cell number: RCB0007) were used in this study. HeLa cells were established from the

cervix cells of a female human.

HeLa cells were authenticated by short tandem repeat (STR) profiling (National Institute of Biomedical Innovation, Osaka, Japan)

Mycoplasma contamination HeLa cells were confirmed to be mycoplasma free (Biotherapy Institute of Japan, Tokyo, Japan).

Commonly misidentified lines (See ICLAC register)

No commonly misidentified cell lines were used in the study.

### Animals and other research organisms

Policy information about <u>studies involving animals</u>; <u>ARRIVE guidelines</u> recommended for reporting animal research, and <u>Sex and Gender in Research</u>

Laboratory animals

d-rR strain medaka (Oryzias latipes), himedaka (Oryzias latipes) and Japanese eel (Anguilla japonica)

Wild animals

No wild animals are used.

Sex was determined by the color and fin shapes of the sample

Field-collected samples No field-collected samples are used

Ethics oversight

Experiments were performed at the Atmosphere and Ocean Research Institute and Department of Aquatic Bioscience of Graduate school of agricultural and life sciences of the university of the Tokyo in accordance with the protocols approved by the animal care and use committee of Atmosphere and Ocean Research Institute of the University of Tokyo (Permission number, P22-6, P22-7).

Note that full information on the approval of the study protocol must also be provided in the manuscript.

#### **Plants**

| Seed stocks           | N/A |
|-----------------------|-----|
| Novel plant genotypes | N/A |
|                       |     |
| Authentication        | N/A |