

## NIH Public Access

**Author Manuscript** 

Genesis. Author manuscript; available in PMC 2014 February 01

#### Published in final edited form as:

Genesis. 2013 February ; 51(2): 128–134. doi:10.1002/dvg.22357.

# *Musashi1*-CreER<sup>T2</sup>: A New Cre Line for Conditional Mutagenesis in Neural Stem Cells

Haruna Takeda<sup>1,2,\*</sup>, Hideto Koso<sup>1,3,\*</sup>, Lino Tessarollo<sup>4</sup>, Neal G. Copeland<sup>1,5</sup>, and Nancy A. Jenkins<sup>1,5,§</sup>

<sup>1</sup>Division of Genomics and Genetics, Institute of Molecular and Cell Biology, Agency for Science, Technology and Research, Singapore 138673

<sup>2</sup>Department of Microbiology, Graduate School of Medicine and Faculty of Medicine, The University of Tokyo, Tokyo, 113-0033, Japan

<sup>3</sup>Division of Molecular and Developmental Biology, The Institute of Medical Science, The University of Tokyo, Tokyo, 108-8639, Japan

<sup>4</sup>Neural Development Group, Mouse Cancer Genetics Program, Center for Cancer Research, National Cancer Institute, Frederick, Maryland, 21702, USA

<sup>5</sup>Cancer Research Program, The Methodist Hospital Research Institute, Houston, TX 77030, USA

### Abstract

The RNA-binding protein Musashi1 (Msi1) is one of two mammalian homologues of *Drosophila* Musashi, which is required for the asymmetric cell division of sensory organ precursor cells. In the mouse central nervous system (CNS) Msi1 is preferentially expressed in mitotically active progenitor cells in the ventricular zone (VZ) of the neural tube during embryonic development and in the subventricular zone (SVZ) of the postnatal brain. Previous studies showed that cells in the SVZ can contribute to long-term neurogenesis in the olfactory bulb (OB) but it remains unclear whether Msi1-expressing cells have self-renewing potential and can contribute to neurogenesis in the adult. Here we describe the generation of *Msi1-CreER<sup>T2</sup>* knock-in mice and show by cell lineage tracing that *Msi1-CreER<sup>T2</sup>*-expressing cells mark neural stem cells (NSCs) in both the embryonic and adult brain. *Msi1-CreER<sup>T2</sup>* mice thus represent a new tool in our arsenal for genetically manipulating NSCs, which will be essential for understanding the molecular mechanisms underlying neural development.

#### Keywords

Msi1; NSCs; SVZ; Olfactory epithelium; Knock-in mice

In order to generate the  $Msi1-CreER^{T2}$  knock-in mice, a targeting vector was created by inserting a  $CreER^{T2}$  expression cassette downstream of the first ATG in Msi1 exon 1 using recombineering, along with a PGKneo selection cassette (Copeland *et al.*, 2001) (Fig. 1A).  $CreER^{T2}$  is an inducible form of the *Cre* recombinase that can be activated by tamoxifen administration (Feil *et al.*, 1997). The targeting cassette was subsequently electroporated into ES cells, and the cells screened for homologous recombination by Southern blotting (Fig. 1A, 1B). Correctly targeted cells were subsequently injected into mouse blastocysts to

Correspondence to: Nancy A. Jenkins, njenkins2@tmhs.org, Tel: 713-441-6623.
\*These authors contributed equally to this work

The authors declare no conflict of interests.

generate Msi1- $CreER^{T2}$  knock-in mice, and the selection cassette was then removed by crossing the mice to  $\beta$ -*actin-Flp* mice. RT-PCR confirmed that the knock-in allele produces no Msi1 transcript containing exon 1 (Fig 2A, see Msi1 F1+R). We examined the coding potential of the shorter transcript in the eye by using an antibody that recognizes an epitope coded by the exon 10 (LAPGYTYQFP) (Kaneko *et al.*, 2000). Msi1 expression was detected in Muller glia and photoreceptors in the wild type mouse retina as previously reported (Nickerson *et al.*, 2011) (Fig. 2B). However, the signal was not detected in the retina of homozygous Msi1- $CreER^{T2}$  knock-in mice (Fig. 2B), indicating that the shorter transcript is not translated into an N-terminal truncated Msi1 protein in the eye. While heterozygous Msi1- $CreER^{T2}$  mice were healthy and fertile, 50% of homozygous Msi1- $CreER^{T2}$  mice (6 of 12) developed hydrocephalus (Fig. 2C), a phenotype described previously for the Msi1 mutant mice that lack exons 1–4 (Sakakibara *et al.*, 2002), indicating that the inactivation of a transcript containing exon 1 is responsible for the phenotype. Northern blot analysis showed that the 1.4 kb Msi1- $CreER^{T2}$  transcript was expressed in retina, brain and testis (Fig. 3), consistent with the known expression pattern of Msi1 (Sakakibara *et al.*, 1996).

To more thoroughly characterize Msi1- $CreER^{T2}$  expression, we crossed Msi1- $CreER^{T2}$ mice with Rosa26-lsl-LacZ reporter (R26R3) mice (Soriano, 1999). In Msi1- $CreER^{T2}$ ; R26R3 mice, LacZ expression is blocked by the presence of the transcriptional floxed-stop (lsl) cassette. However, once Msi1- $CreER^{T2}$  is activated by tamoxifen, the stop cassette is removed by Cre recombination and LacZ expression can now be visualized by X-Gal staining. We focused on the eye, brain, olfactory bulb and testis since these are all known sites of Msi1 expression. Msi1- $CreER^{T2}$ ; R26R3 mice were treated with tamoxifen at 4 weeks of age (Fig. 4A) and euthanized 5 days later. Whole mount staining of the eye showed that the entire retina was stained with X-Gal (Fig. 4B), while paraffin sections showed strong expression in the photoreceptor cell layer (Fig. 4C) where Msi1 expression has previously been observed (Susaki *et al.*, 2009). In the brain, a small number of LacZ+ cells were observed in the wall of the lateral ventricle but not in the rostral migratory stream (RMS) and the OB (Fig. 4D, 4E, 4G and 4H), indicating the presence of Msi1- $CreER^{T2}$ expressing cells in the SVZ. LacZ expression was tamoxifen dependent since no LacZ+ cells were detected in the absence of tamoxifen (Fig 4F).

Next, we traced the lineage of *Msi1-CreER<sup>T2</sup>*-expressing cells by performing X-Gal staining 60-days after tamoxifen treatment. LacZ+ cells were found in the OB, in addition to the RMS and SVZ (Fig. 4I–K), demonstrating that recombination took place in NSCs in the SVZ rather than transit-amplifying cells. Lineage-tracing experiments have shown that labeled transit-amplifying cells and their progeny complete migration to the OB within one month, with no labeled cells remaining in the RMS and SVZ after that time (Kim *et al.*, 2007).

Green fluorescent protein and its variants are useful in lineage tracing, because they allow detection of labeled cells without chemical substrates, and the morphology of labeled cells can be more easily analyzed in tissue sections. Thus, we next generated a second reporter line, *Msi1-CreER<sup>T2</sup>; Rosa26<sup>YFP</sup>*. Analysis of frozen sections of *Msi1-CreER<sup>T2</sup>; Rosa26<sup>YFP</sup>* mice at 2 months after tamoxifen treatment showed that YFP+ cells were present in the SVZ (Fig. 4L) where endogenous *Msi1* is expressed (Figure 4M), indicating that *Msi1-CreER<sup>T2</sup>* expression faithfully recapitulates endogenous *Msi1* expression in the SVZ. Immunohistochemical analysis showed that YFP+ cells in the SVZ expressed GFAP, a marker for adult NSCs (Ming and Song, 2011) (Fig. 4N). NSCs in the SVZ give rise to different subtypes of interneurons in the OB (Ming and Song, 2011). Immunohistochemical analysis showed that express markers for interneurons, such as Calbindin and Calretinin (Ming and Song, 2011) (Fig. 4O and 4P). Together, these data demonstrate that *Msi1-CreER<sup>T2</sup>* is expressed in NSCs of the SVZ.

In the subgranular zone (SGZ), putative multipotent neural stem or progenitor cells, which will give rise to dentate granule cells in the hippocampus, have also been identified (Bonaguidi *et al.*, 2011; Zhao *et al.*, 2008). We detected clusters of YFP+ cells in the dentate gyrus in the hippocampus of *Msi1-CreER<sup>T2</sup>; Rosa26<sup>YFP</sup>* mice at 2 months after tamoxifen addition (Fig. 4Q). YFP+ cells express a marker for dentate granule cell, Calbindin (Ming and Song, 2011) (Fig. 4R), suggesting that Msi1-CreER<sup>T2</sup> is expressed in the precursor cells for dentate granule cells.

The olfactory neuroepithelium (OE) undergoes continual neurogenesis throughout life. *Msi1-CreER<sup>T2</sup>; Rosa26<sup>YFP</sup>* mice were treated with tamoxifen at 4 weeks and the OE was analyzed 8 months later. Clusters of YFP+ cells were occasionally observed in the OE. Most of YFP+ cells appeared to be olfactory receptor neurons (ORNs) as judged by their position in the OE and their cellular morphology (Fig. 4O). It is possible that these clusters of YFP+ cells were derived from a single *Msi1-CreER<sup>T2</sup>*-expressing stem or progenitor cell, because YFP+ cells were usually visible as cell clusters or single cells. We also detected YFP+ cells whose morphology and position seemed to correspond to sustentacular cells (Fig. 4O). Genetic experiments have reported that ORNs and sustentacular cells may share a common progenitor (Beites *et al.*, 2005). Our data suggest that *Msi1-CreER<sup>T2</sup>* is expressed in the common progenitor cells of the OE in adult mice.

Northern blot analysis showed the expression of CreER<sup>T2</sup> in the testis, however we could not detect any cells that express reporter genes in either *Msi1-CreER<sup>T2</sup>; R26R3* or *Msi1-CreER<sup>T2</sup>; Rosa26<sup>YFP</sup>* mice at five days after tamoxifen injection (Fig. 4A), raising a possibility that the expression level of CreER<sup>T2</sup> might not be strong enough to induce recombination in the testis.

Next, we examined *Msi1-CreER*<sup>T2</sup> expression during embryonic development. Pregnant *Msi1-CreER*<sup>T2</sup>; *R26R3* mice were treated with tamoxifen or oil at E10.5 and the embryos stained for LacZ at E12.5 (Fig. 5A). No LacZ staining was observed in oil-treated embryos at E12.5 (Fig. 5B). LacZ+ cells were, however, observed within the entire nervous system at E12.5 in tamoxifen-treated embryos (Fig. 5C and 5D), indicating that *Msi1-CreER*<sup>T2</sup> is expressed in the entire nervous system at E10.5. Coronal sections of embryonic brains showed strong X-Gal staining in the entire neuroepithelium (Fig. 5E), indicating that *Msi1-CreER*<sup>T2</sup> is expressed in either NSCs or differentiating precursor cells in the developing cortex.

During neural development, the cortical layer is formed by an inside-out pattern of migration of differentiating cells, confining mitotic NSCs to the ventricular surface of the developing cortex, whereas differentiated cells occupy the peripheral region (Nadarajah and Parnavelas, 2002). To determine whether *Msi1-CreER<sup>T2</sup>*-expressing cells are NSCs or differentiating precursor cells in the developing cortex, we induced recombination in Msil-CreER<sup>T2</sup>; R26R3 embryos at E10.5 and then examined the brain at E17.5. No LacZ+ cells were seen in oil-treated embryos (Fig. 5O), but in tamoxifen treated embryos, LacZ was expressed within the entire cortex in the forebrain, cerebellum and brain stem (Fig. 5I-L), showing that *Msi1-CreER<sup>T2</sup>*-expressing cells can give rise to differentiated cells in these regions. Importantly, intense LacZ staining was observed in the ventricular surface of the striatum (Fig. 5J and 5N), suggesting that *Msi1-CreER*<sup>T2</sup>-expressing cells have permanently labeled self-renewing stem cells in the VZ. These data indicate that Msi1-CreER<sup>T2</sup> is also expressed in NSCs during embryonic development. Forni and colleagues showed that the activation of *nestin-CreER*<sup>T2</sup> transgene induces brain abnormalities in a dose-dependent manner (Forni et al., 2006). However, the administration of 1 mg of tamoxifen did not induce any brain defects in the Msi1-CreER<sup>T2</sup> line (Fig 5I-L and 5O), suggesting that the

expression level of CreER<sup>T2</sup> is not as strong as the *nestin-CreER<sup>T2</sup>* line (Forni *et al.*, 2006; Imayoshi *et al.*, 2006).

In the adult OE we showed that  $Msi1-CreER^{T2}$  was likely expressed in progenitor cells for ORNs and sustentacular cells. Next we examined whether  $Msi1-CreER^{T2}$  is expressed in embryonic OE progenitor cells. During embryonic development of the OE, expansion of stem cells takes place by E12.5. Around E13.5-E14.5, the OE becomes organized into its mature pattern and established neurogenesis is initiated (Beites *et al.*, 2005). We treated pregnant  $Msi1-CreER^{T2}$ ; R26R3 reporter mice with tamoxifen or oil at E10.5 and analyzed LacZ staining in embryos at E17.5. Clusters of LacZ+ cells were observed in the entire OE (Figure 5M), showing that  $Msi1-CreER^{T2}$  is expressed in stem or progenitor cells of the OE as early as E10.5.

Inducible recombination in NSCs has also been reported using Nestin-CreER<sup>T2</sup> transgenes (Chen et al., 2009; Imayoshi et al., 2006; Lagace et al., 2007). Although we could not determine the efficiency of recombination due to the poor quality of CreER<sup>T2</sup> immunostaining, X-Gal staining of the adult SVZ and the embryonic cortex in Msi1-CreER<sup>T2</sup>; R26R3 mice showed comparable patterns with those reported in Nestin-CreER<sup>T2</sup> transgenic mice (Chen et al., 2009; Imayoshi et al., 2006). However, one striking difference was expression in the OE. The Nestin regulatory element directed reporter expression to a subpopulation of embryonic OE progenitor cells that are located in the dorsal medial zone (Murdoch and Roskams, 2008). In contrast, we did not detect regional restriction of CreER<sup>T2</sup> expression in the OE. Tamoxifen administration of Msi1-CreER<sup>T2</sup>; Rosa26<sup>YFP</sup> adult mice induced clusters of LacZ+ cells in ORN and sustentacular cells in the OE. suggesting that CreER<sup>T2</sup> is expressed in the common progenitor for these lineages. Moreover, tamoxifen treatment in the embryos as early as E10.5 induced clusters of LacZ+ cells in the OE, indicating that CreER<sup>T2</sup> is expressed in expanding stem cells in the OE. Detailed lineage tracing of CreER<sup>T2</sup> expressing cells will be needed to reveal the features of neurogenesis in the OE.

*Msi1* is also reported to be a marker for intestinal stem cells (Potten *et al.*, 2003), however by Northern blotting or lineage tracing, we failed to detect *Msi1-CreER<sup>T2</sup>* expression in the intestine (data not shown). *Msi1* is known to have several transcriptional variants (Susaki *et al.*, 2009) (Ensemble Genome Browser, http://www.ensembl.org/). In *Msi1-CreER<sup>T2</sup>* knockin mice, *Msi1-CreER<sup>T2</sup>* expression initiates from the first ATG in exon 1. The *Msi1* transcript initiating from this ATG was not detected in the intestine (Fig. 2A, see Msi1 F1+R), whereas a shorter transcript lacking exon 1 is expressed in the intestine (Fig. 2A, see Msi1 F2+R), suggesting that the alternative transcript is abundant in the intestine. Our *Msi1-CreER<sup>T2</sup>* knock-in mice should therefore represent a useful new tool for manipulating NSCs, which is not confounded by expression in the intestine.

#### Methods

#### Generation of Msi1-CreER<sup>T2</sup> knock-in mice

A mouse BAC clone (RP23-268H8) containing the entire *Msi1* genomic locus was obtained from the Children's Hospital Oakland Research Institute (CHORI) and electroporated into the recombineering strain SW105 (Warming *et al.*, 2005). The entire 10.0 kb *Msi1* genomic locus in addition to 5.0 kb of upstream and 5.0 kb of downstream sequence, was then subcloned into pBlight (Warming *et al.*, 2006) by recombineering. A CreER<sup>T2</sup> fragment was subsequently excised from pCAG-CreERT2 (Addgene) and inserted into a plasmid derivative of pL451 (Liu *et al.*, 2003) that has a *Neomycin* selection cassette flanked between two FRT sites. Sequences of the first exon located downstream of the first ATG were then replaced by the CreER<sup>T2</sup>-polyA, FRT-PGKneo-FRT cassette using

recombineering. The Msi1 targeting vector was then linearized and transfected into B6/129 F1 hybrid embryonic stem (ES) cells (Southon and Tessarollo, 2009). G418 and FIAU double-resistant ES cell clones were selected and analyzed for homologous recombination using Southern blot hybridization of BamHI-digested genomic DNA. Probes were PCR amplified with the following primers. Probe C F: 5'-ccc agg act cca ccc atgc-3', R: 5'-agg caa ctc att aac aaa ag-3'; Probe2 F: 5'-cct ctt gac tct agg cca tc-3'. R: 5'-cag cag gac ctg gca gcc tt-3'. Two independently targeted ES cell clones were injected into C57BL/6 blastocysts to produce chimeric mice (Reid and Tessarollo, 2009). The chimeras were then mated to C57BL/6 mice and germline transmission in agouti offspring confirmed by PCR. The neomycin selection cassette was later excised in vivo by crossing the mice with b-actin-FLPe mice (The Jackson Laboratory). *Msi1-CreER<sup>T2</sup>* heterozygous mice were then intercrossed to obtain homozygous animals. Homozygous mice showing signs of hydrocephalus (an enlarged and domed head) were sent to necropsy, and the brains fixed in 4% PFA in PBS. All animal experiments were performed with institutional animal care and use committee (IACUC) approval. The mice will be made available to the scientific community upon request.

#### Reporter strains and tamoxifen treatment

The *R26R3* reporter strain was obtained from Philippe Soriano (Mount Sinai School of Medicine, New York, NY) (Soriano, 1999), while the *Rosa26*<sup>YFP</sup> reporter strain (Srinivas *et al.*, 2001) was purchased from the Jackson Laboratory. Tamoxifen (Sigma) was prepared as a 20 mg/ml stock solution in corn oil (Sigma). At 4 weeks of age, *Msi1-CreER*<sup>T2</sup>;*R26R3 or Msi1-CreER*<sup>T2</sup>;*R0sa26*<sup>YFP</sup> mice were injected intraperitoneally with 2 mg of tamoxifen for 3 consecutive days. Tissues were collected at 5 days, 2 months or 8 months after tamoxifen treatment. For embryonic induction, 1 mg of tamoxifen was injected intraperitoneally into pregnant mothers at E10.5, and embryos subsequently collected at E12.5 or E17.5.

#### Northern blotting and RT-PCR

Total RNA was isolated using Trizol (Invitrogen). Northern blotting was performed according to standard protocols. Ten µg of total RNA was run on a 1% agarose gel in MOPS buffer (0.4M of 3-morpholinopropanesulfonic acid, 0.1M of sodium acetate and 20mM of EDTA). The coding regions of *Cre* and *Gapdh* were used as probes. Superscript III (Invitrogen) was used to synthesize cDNA. PCR was performed using the following primers. Msi1 F1: 5'-CAC GAC CCC TGC AAG ATG-3', F2: 5'-TCG TCA CTT TCA TGG ACC AG-3', R: 5'-TGA AGG CAT CCA TTC CGT AG-3'; b-actin F: 5'-TAG ACT TCG AGC AGG AGA TG-3', R: 5'-ACT CAT CGT ACT CCT GCT TG-3'.

#### X-Gal staining and histology

For whole-mount X-Gal staining, embryos and adult tissues were fixed in 1% formaldehyde, 0.2% glutaraldehyde, 0.02% IGEPAL CA-630 (Sigma) in phosphate buffered saline (PBS) for 10 min at room temperature (RT), rinsed in PBS three times, and stained at 37°C overnight in X-Gal staining buffer (1 mg/ml X-Gal, 2 mM MgCl<sub>2</sub>, 0.02% IGEPAL CA-630, 1 mM K<sub>3</sub>Fe(CN)<sub>6</sub>, 1 mM K<sub>4</sub>Fe(CN)<sub>6</sub> in PBS). For X-Gal staining of tissue sections, embryos and adult brains were fixed and sectioned using a vibratome (100 mm). Then, free-floating sections were stained in X-Gal staining buffer as described above. For retinal sections, X-Gal stained eyes were fixed with 4% paraformaldehyde (PFA) in PBS, washed and embedded in paraffin wax prior to sectioning. Sections were counterstained with eosin. For visualization of YFP+ cells, brains were fixed with 4% PFA in PBS overnight at 4°C, washed in PBS, cryoprotected in 25% sucrose in PBS and embedded in Tissue-Tek (Sakura). Frozen sections were made using a cryostat (Leica CM3050 S) at 30 mm thickness. Sections were counter stained with 4',6-diamidino-2-phenylindole dihydrochloride (DAPI) (Sigma) and analyzed using an upright microscope (Zeiss

AxioImager Z1). Antibodies used for immunohistochemical analysis were rabbit polyclonal anti-Musashi1 (Millipore, AB5977), rat monoclonal anti-Musashi1 (MBL, D270-3) for the detection of an epitope coded by the exon 10 (LAPGYTYQFP) and Alexa594-conjugated goat polyclonal anti-rabbit IgG (Molecular Probes).

#### Acknowledgments

We thank Keith Rogers, Susan M. Rogers and Vanessa Shiyun Tay for technical support. This work was supported by the Agency for Science, Technology and Research (A-STAR), Singapore, and the Cancer Prevention Research Institute of Texas. N.A.J. and N.G.C. are both CPRIT Scholars in Cancer Research. LT was supported by the Intramural Research Program of the NCI, Center for Cancer Research, NIH.

#### References

- Beites CL, Kawauchi S, Crocker CE, Calof AL. Identification and molecular regulation of neural stem cells in the olfactory epithelium. Experimental cell research. 2005; 306:309–316. [PubMed: 15925585]
- Bonaguidi MA, Wheeler MA, Shapiro JS, Stadel RP, Sun GJ, Ming GL, Song H. In vivo clonal analysis reveals self-renewing and multipotent adult neural stem cell characteristics. Cell. 2011; 145:1142–1155. [PubMed: 21664664]
- Chen J, Kwon CH, Lin L, Li Y, Parada LF. Inducible site-specific recombination in neural stem/ progenitor cells. Genesis. 2009; 47:122–131. [PubMed: 19117051]
- Copeland NG, Jenkins NA, Court DL. Recombineering: a powerful new tool for mouse functional genomics. Nature reviews. Genetics. 2001; 2:769–779. [PubMed: 11584293]
- Feil R, Wagner J, Metzger D, Chambon P. Regulation of Cre recombinase activity by mutated estrogen receptor ligand-binding domains. Biochemical and biophysical research communications. 1997; 237:752–757. [PubMed: 9299439]
- Forni PE, Scuoppo C, Imayoshi I, Taulli R, Dastru W, Sala V, Betz UA, Muzzi P, Martinuzzi D, Vercelli AE, Kageyama R, Ponzetto C. High levels of Cre expression in neuronal progenitors cause defects in brain development leading to microencephaly and hydrocephaly. The Journal of neuroscience : the official journal of the Society for Neuroscience. 2006; 26:9593–9602. [PubMed: 16971543]
- Imayoshi I, Ohtsuka T, Metzger D, Chambon P, Kageyama R. Temporal regulation of Cre recombinase activity in neural stem cells. Genesis. 2006; 44:233–238. [PubMed: 16652364]
- Kaneko Y, Sakakibara S, Imai T, Suzuki A, Nakamura Y, Sawamoto K, Ogawa Y, Toyama Y, Miyata T, Okano H. Musashi1: an evolutionally conserved marker for CNS progenitor cells including neural stem cells. Developmental neuroscience. 2000; 22:139–153. [PubMed: 10657706]
- Kim EJ, Leung CT, Reed RR, Johnson JE. In Vivo Analysis of Ascl1 Defined Progenitors Reveals Distinct Developmental Dynamics during Adult Neurogenesis and Gliogenesis. JNeurosci. 2007; 27:12764–12774. [PubMed: 18032648]
- Lagace DC, Whitman MC, Noonan MA, Ables JL, DeCarolis NA, Arguello AA, Donovan MH, Fischer SJ, Farnbauch LA, Beech RD, DiLeone RJ, Greer CA, Mandyam CD, Eisch AJ. Dynamic contribution of nestin-expressing stem cells to adult neurogenesis. The Journal of neuroscience : the official journal of the Society for Neuroscience. 2007; 27:12623–12629. [PubMed: 18003841]
- Liu P, Jenkins NA, Copeland NG. A highly efficient recombineering-based method for generating conditional knockout mutations. Genome research. 2003; 13:476–484. [PubMed: 12618378]
- Ming GL, Song H. Adult neurogenesis in the mammalian brain: significant answers and significant questions. Neuron. 2011; 70:687–702. [PubMed: 21609825]
- Murdoch B, Roskams AJ. A novel embryonic nestin-expressing radial glia-like progenitor gives rise to zonally restricted olfactory and vomeronasal neurons. The Journal of neuroscience : the official journal of the Society for Neuroscience. 2008; 28:4271–4282. [PubMed: 18417707]
- Nadarajah B, Parnavelas JG. Modes of neuronal migration in the developing cerebral cortex. Nature reviews. Neuroscience. 2002; 3:423–432. [PubMed: 12042877]

- Nickerson PE, Myers T, Clarke DB, Chow RL. Changes in Musashi-1 subcellular localization correlate with cell cycle exit during postnatal retinal development. Experimental eye research. 2011; 92:344–352. [PubMed: 21320487]
- Potten CS, Booth C, Tudor GL, Booth D, Brady G, Hurley P, Ashton G, Clarke R, Sakakibara S, Okano H. Identification of a putative intestinal stem cell and early lineage marker; musashi-1. Differentiation; research in biological diversity. 2003; 71:28–41.
- Reid SW, Tessarollo L. Isolation, microinjection and transfer of mouse blastocysts. Methods Mol Biol. 2009; 530:269–285. [PubMed: 19266343]
- Sakakibara S, Imai T, Hamaguchi K, Okabe M, Aruga J, Nakajima K, Yasutomi D, Nagata T, Kurihara Y, Uesugi S, Miyata T, Ogawa M, Mikoshiba K, Okano H. Mouse-Musashi-1, a neural RNA-binding protein highly enriched in the mammalian CNS stem cell. Developmental biology. 1996; 176:230–242. [PubMed: 8660864]
- Sakakibara S, Nakamura Y, Yoshida T, Shibata S, Koike M, Takano H, Ueda S, Uchiyama Y, Noda T, Okano H. RNA-binding protein Musahi familiy: Roles for CNS stem cells and a subpopulation of ependymal cells revealed by targeted disruption and antisense ablation. PNAS. 2002; 99:15194– 15199. [PubMed: 12407178]
- Soriano P. Generalized lacZ expression with the ROSA26 Cre reporter strain. Nat. Genet. 1999; 21:70–71. [PubMed: 9916792]
- Southon E, Tessarollo L. Manipulating mouse embryonic stem cells. Methods Mol Biol. 2009; 530:165–185. [PubMed: 19266324]
- Srinivas S, Watanabe T, Lin CS, William CM, Tanabe Y, Jessell TM, Costantini F. Cre reporter strains produced by targeted insertion of EYFP and ECFP into the ROSA26 locus. BMC developmental biology. 2001; 1:4. [PubMed: 11299042]
- Susaki K, Kaneko J, Yamano Y, Nakamura K, Inami W, Yoshikawa T, Ozawa Y, Shibata S, Matsuzaki O, Okano H, Chiba C. Musashi-1, an RNA-binding protein, is indispensable for survival of photoreceptors. Experimental eye research. 2009; 88:347–355. [PubMed: 18662689]
- Warming S, Costantino N, Court DL, Jenkins NA, Copeland NG. Simple and highly efficient BAC recombineering using galK selection. Nucleic acids research. 2005; 33:e36. [PubMed: 15731329]
- Warming S, Rachel RA, Jenkins NA, Copeland NG. Zfp423 is required for normal cerebellar development. Molecular and cellular biology. 2006; 26:6913–6922. [PubMed: 16943432]
- Zhao C, Deng W, Gage FH. Mechanisms and functional implications of adult neurogenesis. Cell. 2008; 132:645–660. [PubMed: 18295581]

Takeda et al.



(A) A *CreER<sup>T2</sup>-PGKneo* expression cassette was inserted into the first ATG located in exon 1 of *Msi1*. The *pBlight-Msi1-CreER<sup>T2</sup>-PGKneo* targeting vector was used for homologous recombination to generate the *Msi1-CreER<sup>T2</sup>* targeted allele. The *PGKneo* cassette was subsequently excised by flip-mediated recombination. B; *Bam*HI, prbC (5' probe), prb2 (3' probe). (B) Southern blot analysis to screen ES cells for correctly targeted alleles. For the knock-in (KI) allele, *Bam*HI digestion produces 14.5 kb and 5 kb bands for prbC and prb2, respectively. For the wild type (+) allele, *Bam*HI digestion produces 19.5 kb bands for both probes.

Takeda et al.

Page 9



#### FIG. 2. Expression of *Msi1-CreER<sup>T2</sup>* in homozygous knock-in mice

(A) RT-PCR analysis of *Msi1* mRNA expression in wild type (wt) and *Msi1-CreER*<sup>T2</sup> homozygous knock-in mice (*KI/KI*). Red arrows, PCR primers; Br, brain; Ey, eye; SI, small intestine; Co, colon; Te, testis. (B) Msi1 expression was detected in Muller glial cells in the INL and photoreceptors in the ONL of the wild type (wt) retina, but not in the retina of homozygous knock-in mice (*KI/KI*). GCL, ganglion cell layer; INL, inner nuclear layer; ONL, outer nuclear layer. The nucleus is counterstained with DAPI. Bars represent 50 µm. (C) Hydrocephalus was often observed in Msi1-CreER<sup>T2</sup> homozygous knock-in mice (*KI/KI*).

## Re Br St Si Co Li Ki Mu Te Ov Ut







FIG. 3. Tissue distribution of  $CreER^{T2}$  transcripts in  $Msil-CreER^{T2}$  knock-in mice The expression levels of  $CreER^{T2}$  mRNA in adult tissues were determined by Northern blot analysis. The membrane was then reprobed with *Gapdh* as a loading control. Re, retina; Br, brain; St, stomach; Si, small intestine; Co, colon; Li, liver; Ki, kidney; Mu, muscle; Te, testis; Ov, ovary; Ut, uterus Takeda et al.



FIG. 4. Lineage tracing of *Msi1-CreER*<sup>T2</sup>-expressing cells *Msi1-CreER*<sup>T2</sup>; *R26R3 or Msi1-CreER*<sup>T2</sup>; *Rosa26*<sup>YFP</sup> mice were treated for three consecutive days, with 2 mg of tamoxifen or corn oil, beginning at 4 weeks of age (A). X-Gal staining of the eye (B), retina (C) and brain (D-H) 5 days after tamoxifen or corn oil treatment. X-Gal staining of sagittal (I) and coronal (J, K) sections of the brain at 2 months after tamoxifen treatment. LacZ+ cells were observed in the RMS (I, arrow heads). YFP expression in the SVZ (L, N), OB (O, P) and SGZ (Q, R) at 2 months after tamoxifen treatment, or 8 months after tamoxifen treatment for the OE (S). Endogenous Msi1 expression was observed in the SVZ (M). In the SVZ, YFP+ cells and Msi1+ cells showed a

similar pattern of localization (**L**, **M**). YFP+ cells in the SVZ expressed GFAP, a marker for NSCs (**N**, arrow heads). YFP+ cells in the OB expressed markers for interneurons, Calbindin (**O**, arrow heads) and Calretinin (**P**, arrow heads). YFP+ cells in the SGZ expressed Calbindin, a marker for dentate granule neuron (**R**, arrow head). In the OE, YFP+ cells with morphological characteristics of sustentacular cells were observed (**S**, arrow head). Br, brain; Le, lens; LV, lateral ventricle; Nc, nasal cavity; Re, retina; Tam, tamoxifen; Te, testis; Bars: 100 µm.

Takeda et al.



FIG. 5. Lineage tracing of Msi1- $CreER^{T2}$ -expressing cells in the embryonic neural system Time course of tamoxifen injection and X-Gal staining (A and F). Pregnant mice were treated with 1 mg of tamoxifen (C–E, H–N) or corn oil (B and O) at E10.5. Msi1- $CreER^{T2}$ ; R26R3 embryos were stained with X-Gal in whole mount at E12.5 (A–E), or embryonic brains were sectioned at various levels using a vibratome (G), and stained with X-Gal at E17.5 (F–O). Strong X-Gal staining was observed in cells lining the lateral ventricle (N, arrowheads).