



**FIG. 7.** The proportion of neurons from neural stem cells derived from D3m-Mock1 and D3m-shM6A1 cells. Cells were differentiated with ACM for 2 weeks, fixed, and immunostained with Tubb3, MAP2, DCX, ChAT, GADs, GABA, TH, and serotonin antibodies. The number of total cells was counted by the number of DAPI staining nuclei. Values are expressed as the mean  $\pm$  SE ( $n = 4$ ). \* $p < 0.01$  and \*\* $p < 0.001$  compared with neurons from D3m-Mock1 cells. All experiments were independently carried out at least three times, and almost the same results were obtained each time.

results imply that the proportion of NSC in differentiated cells derived from D3m-shM6A1 cells was smaller than those in D3m-Mock1 cells. Therefore, it was suggested that suppression of GPM6A transcript caused NSC generation from undifferentiated ES cells. There seems to be a correlation between the ratio of suppressed expression of mouse GPM6A transcripts and the ratio of differentiation of NSC from undifferentiated mouse ES cells. Moreover, the expression level of neuroectodermal marker genes, which are associated with the development of the brain [14–24], was reduced by the suppression of mouse GPM6A transcripts using shRNA against mouse GPM6A in neural-differentiated mouse ES cells (Fig. 2B). On the other hand, expression levels of mesodermal marker genes (*Nkx2.5*, *GATA-4*, *BMP-4*, *Fli1* etc. [32,33]) were clearly increased in differentiated cells derived from D3m-shM6A1 cells compared with those in D3m-Mock1 cells (data not shown). In addition, we also obtained results showing that expression level of undifferentiated ES marker gene (*OCT3/4*) was barely detected in differentiated cells derived from both D3m-Mock1 and D3m-shM6A1 cells (data not shown), and the doubling time of undifferentiated D3m-shM6A1 cells was equivalent to that of undifferentiated D3m-Mock1 cells (data not shown). A recent study has shown that transfection of small interference RNA in hippocampal neurons leads to decreased filopodium/spine formation but not to cell death [13]. As a result of these findings, we speculated that reduction of NSC generation by suppression of neural differentiation by GPM6A knockdown might lead to increase of other germ lineages, or ES cells remaining undifferentiated, but not to cell death.

Mouse GPM6A is known to be expressed in mouse neurons [8,9]. It is also reported that expression of mouse GPM6A is observed from embryonic day 10 on cells in the marginal zone of the neural tube and is widespread throughout the brain and spinal cord, where it remains throughout adulthood [8,9]. In addition, it has been shown that treatment with anti-GPM6A antibody interferes with extension of neurites in cultured neurons [8]. A recent study has shown that GPM6A is related to neurite outgrowth, filopodium/spine formation, and synaptic formation in cultured neurons [13]. In this study, the inhibition of mouse

GPM6A expression decreased the generation of NSC. In addition, expression levels of cholinergic neuron marker (ChAT), catecholaminergic neuron markers (TH and DCC), and GABAergic neuron marker (GADs) [29–31] decreased in neurons derived from D3m-shM6A cells (Fig. 5). Moreover, our results revealed a suppression of differentiation into various neuronal cells, such as cholinergic (ChAT), catecholaminergic (TH and serotonin), and GABAergic (GABA and GADs) neurons, derived from D3m-shM6A1 cells (Figs 6 and 7). Furthermore, there was no difference in differentiation efficiency by neuronal cell type. Taken together, these findings suggest that mouse GPM6A is necessary for differentiation into NSC from mouse ES cells, and is one of the factors that trigger differentiation of NSC to various neuronal cells. However, the effect of shRNAs against mouse GPM6A, in which the level of expression of mouse GPM6A transcripts was maximally suppressed by about 80% in neural-differentiated cells derived from mouse ES cells, was extant (Figs 2A and 5). Therefore, it is uncertain how the remaining mouse GPM6A transcripts influence neural differentiation from NSC.

In the present study, we showed that inhibition of mouse GPM6A transcripts expression reduced NSC generation and neuron proportion derived from mouse ES cells. Further investigation, such as analysis of knockout mice, may lead to a better understanding of the relationship between neural differentiation and GPM6A signal transduction.

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