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Mol Reprod Dev. Author manuscript; available in PMC 2018 February 19.

Published in final edited form as:

Author manuscript

Mol Reprod Dev. 2016 August ; 83(8): 675–683. doi:10.1002/mrd.22682.

## **Keeping stem cells under control: new insights into the mechanisms that limit niche-stem cell signaling within the reproductive system**

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## **Summary**

Adult stem cells reside in specialized microenvironments called niches that maintain stem cells in an undifferentiated and self-renewing state. Despite extensive studies on the signaling pathways that operate within stem cells and their niches, the mechanisms that restrict niche signal exclusively to stem cells remained elusive: such a mechanism is crucially important to ensure that stem cells undergo self-renewal while their progeny, often located just one cell diameter away from the niche, differentiate. Here, we review recent progress on the characterization of nichestem cell interactions with special focus on emerging mechanisms that spatially restrict niche signaling.

Adult stem cells reside in specialized microenvironments called niches that maintain stem cells in an undifferentiated and self-renewing state. Despite extensive studies on the signaling pathways that operate within stem cells and their niches, the mechanisms that restrict niche signal exclusively to stem cells remained elusive: such a mechanism is crucially important to ensure that stem cells undergo self-renewal while their progeny, often located just one cell diameter away from the niche, differentiate. Here, we review recent progress on the characterization of nichestem cell interactions with special focus on emerging mechanisms that spatially restrict niche signaling.

## **Keywords**

Germline stem cells; Niche; Short range signaling; MT-nanotubes; Spermatogonial stem cells

## **Introduction**

Cells need to communicate with their neighbors in the correct manner and at the right time to build and maintain functional tissues and organs. Only a handful of signaling pathways

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appear to mediate the majority of cell-to-cell communication within complex tissues. While much has been learned about the molecular mechanics of these pathways, how signal transduction is spatially and temporally regulated in such a precise manner in vivo remains less well understood. Adult tissue homeostasis, in particular, depends on the correct spatiotemporal regulation of signaling between stem cells and their cellular neighbors. Improper signaling can lead to maladaptive increases or decreases in stem cell numbers. Such changes can result in cancer or tissue degeneration. Mechanisms that adjust stem cell signaling in the face of ever changing conditions ensure the proper balance of stem cell self-renewal and differentiation needed for normal tissue function (reviewed in Morrison and Kimble, 2006Rando 2006)). In this review, we will highlight recent insights into the mechanisms that fine-tune stem cell signaling in vivo, with a particular focus on the reproductive system. The general underlying mechanisms involved in regulating stem cell-niche signaling in the ovary and testis are likely used in other stem cell systems as well.

## **Stem cell niches and signaling**

The "niche" hypothesis, first proposed by Schofield in 1978 (Schofield 1978), posits that local environments determine whether stem cells remain in an undifferentiated state in vivo. Since this original publication, numerous cellular and non-cellular niches have been described in the literature (reviewed in (Morrison and Spradling 2008) (Wagers 2012) (Scadden 2014)). Niches have also been found to regulate stem cell division rates and survival (reviewed in (Li and Xie 2005), (Morrison and Spradling 2008)). In "cellular niches", dedicated niche cells form specialized microenvironments that promote stem cell self-renewal and/or prevent stem cell differentiation. Niche cells influence stem cell behavior by producing various signaling molecules, such as Delta, Hedgehog (Hh), bone morphologic proteins (BMPs), Wnt/Wingless (Wg), cytokines, chemokines and other growth factors (reviewed in (Li and Xie 2005), (Morrison and Spradling 2008)). In "non-cellular niches", extracellular molecules, such as extracellular matrix (ECM) proteins, provide essential signals to create the niche. The ECM can also function to concentrate self-renewing signaling molecules that might come from distant sources, thus creating a specialized microenvironment for stem cells. Additional variables influence stem cell behavior. For example, pH, oxygen, ions, mechanical force and electrical stimuli can all modulate stem cell activity, adding to the complexity in niche-mediated stem cell regulation (reviewed in (Wagers 2012)).

While significant progress has been made in understanding which niche signals foster stem cell self-renewal, there is a considerable lack of understanding regarding the mechanisms that prevent inappropriate delivery of self-renewing signals to stem cell progeny that have left the niche. Further insights into these mechanisms will have important implications for our understanding of tissue homeostasis and disease.

#### **Germline stem cell systems in invertebrate model organisms**

The germline stem cells (GSCs) of C. elegans and Drosophila have long served as useful models for studying stem cell niches. The simplicity and accessibility of worm and fly gonads, combined with the availability of robust and sophisticated genetic tools have greatly

accelerated the characterization of the in vivo cellular niches that help to maintain these GSCs. The *C. elegans* gonad represents perhaps one of the simplest examples of a cell based stem cell niche. A distal tip cell (DTC), located at the tip of each gonad arm, extends a number of cellular projections that make contact with a small group of undifferentiated and mitotically active germ cells (Figure 1A). Ablation of the DTC causes germ cells at the tip of the gonad to exit mitosis and initiate the meiotic program. Further work has shown that the DTC acts to prevent undifferentiated germ cells from entering meiosis via Notch signaling pathway (see below, reviewed in (Byrd and Kimble 2009)(Kimble 2014)).

Drosophila gonads house slightly more complex cellular niches. In male Drosophila, a cluster of hub cells located at the apical tip of each testis provides a niche for GSCs. In the ovary, a small group of 5–7 cap cells help form the female GSC niche. Hub cells and cap cells both produce a number of ligands essential for germline stem cell (GSC) self-renewal. In males, hub cells produce Unpaired (Upd), a ligand in the Jak/Stat signaling pathway, and Decapentaplegic (Dpp) and Glass bottom boat (Gbb), ligands in the BMP pathway. BMP signaling also promotes GSC maintenance in the ovary (Michel et al. 2012) (Amoyel et al. 2013) (Luo et al. 2015). In both male and female Drosophila gonads, ectopic expression of niche ligands leads to expansion of GSC-like cells outside of the normal niche, and/or delays in the differentiation of GSC progeny, demonstrating that niche-produced factors play a major role in stem cell fate determination (Tulina and Matunis 2001) (Kiger et al. 2001)(Xie and Spradling 1998).

#### **Mammalian SSC (spermatogonial stem cell) niche**

Recent work has also begun to cast light on the complex nature of niche-stem cell interactions within the mammalian testis. Spermatogonia reside within the basal compartment of the seminiferous tubules and are classified as  $A_{single}$ ,  $A_{paired}$ ,  $A_{aligned}$ , Intermediate and B-subtypes, based on morphological and molecular markers (J. M. Oatley and Brinster 2012)(S.-R. Chen and Liu 2015). Recent work using lineage tracing has shown that a Pax7+ subset of the  $A_{single}$  population contains bona fide spermatogonial stem cells (SSCs) (Aloisio et al. 2014). Pax7+  $A<sub>single</sub> SSCs$  are fast cycling stem cells and have longterm self-renewal capacity. Further quantitative analysis will provide insights into what percentage of bona fide stem cells are Pax7+  $A_{single}$  cells and what percentage of Pax7+ Asingle cells are stem cells. For example, in addition to Pax7, Id4 also marks a rare subset of A<sub>single</sub> spermatogonia potentially enriched for stem cells (Chan et al. 2014)(Sun et al. 2015). The relationship between Pax7+  $A_{single}$  cells and Id4+  $A_{single}$  cells remains unknown.

Glial cell line-derived neutrophic factor (GDNF), a member of the TGFβ super family of signaling molecules, and its receptor GDNF-family receptor-α1 (GFR-α1) comprise a core SSC self-renewal signaling pathway (J. M. Oatley and Brinster 2012)(S.-R. Chen and Liu 2015). GFR-α1 is expressed in subsets of Asingle, Apaired, Aaligned cells (Grasso et al. 2012), while Sertoli cells express GDNF (Meng et al. 2000). Interestingly, A<sub>paired</sub> and A<sub>aligned</sub> cells have the ability to fragment into single cells under certain conditions, perhaps suggesting these different cell types can dedifferentiate in response to niche signals (Nakagawa et al. 2010). Transplantation experiments show that Sertoli cells support the maintenance of SSCs (M. J. Oatley et al. 2011). GDNF heterozygous mutants display premature differentiation of

SSCs (Meng et al. 2000) and decreases in GDNF expression correlate with decreases in functional SSCs during the course of aging (Ryu et al. 2006). Moreover, over-expression of GDNF appears to block germ cell differentiation, giving rise to an expansion of undifferentiated stem cell-like germ cells (Meng et al. 2000). These findings together point to GDNF produced by Sertoli cells as being a critical factor in the maintenance of SSCs.

While Pax7 and Id4 mark a rare subset of undifferentiated spermatogonia compared to the larger GFR-α1+ population, it remains unclear whether niche signaling directly influences the expression of these two transcription factors. This is in contrast to the clear niche-stem cell signaling relationship that exists between Sertoli-derived GDNF and GFR-α1 expressing spermatogonial cells. Whether a bona fide SSC niche specifically regulates Id4 and/or Pax7 expression in neighboring germ cells, conferring SSC identity, represents a point of significant interest.

While it is clear that GDNF represents a principle niche signal that regulates SSC selfrenewal, several fundamental questions arise:

- **1.** GDNF-GFR-α1 signaling likely occurs in a population of cells broader than bona fide SSCs (including A<sub>aligned</sub> populations). Does this indicate that the bona fide niche (e.g. potentially a subset of Sertoli cells) provides additional unidentified SSC-specifying signals?
- **2.** Sertoli cells are present throughout the seminiferous tubules, but the number and position of potential SSCs appears more limited. Whether only a subset of Sertoli cells serve as niches remains elusive.
- **3.** Furthermore, Sertoli cells encapsulate not only SSCs but also all stages of differentiating germ cells. Sertoli cells are large cells that occupy space from the basement membrane to the lumen of the seminiferous tubules, thus contacting all stages of germ cells from SSCs to differentiating spermatids. This begs the question of how Sertoli cells can specify SSC identity, while encapsulating (and likely regulating) spermatid differentiation at the same time.

A few possibilities exist in regards to these questions. Oatley et al. (2009) showed that Leydig cells and select peritubular myoid cells express colony-stimulating factor 1 (CSF1). A SSC enriched Thy+ population of germ cells express the receptor for CSF1 and recombinant CSF1 appeared to enhance SSC self-renewal. A more recent study showed that interstitial macrophages also express CSF1, in addition to enzymes involved in retinoic acid enzyme biosynthesis (DeFalco et al. 2015). The phenotypes resulting from depletion of macrophages within the testis remain somewhat controversial: early studies suggest loss of macrophages disrupt meiotic progression within germ cells while more recent findings indicate that ablation of macrophages results in reduced numbers of A<sub>aligned</sub> cells (Cohen et al. 1996)(Cohen, Hardy, and Pollard 1997)(Pollard et al. 1997)(DeFalco et al. 2015).

Additional factors may influence niche size. For example, Follicle Stimulating Hormone and Testosterone for different sources influence the activity of Sertoli cells (J. M. Oatley and Brinster 2012)(Smith and Walker 2014). Other studies have shown that the basement membrane that lines seminiferous tubules and Peritubular Myoid (PM) cells may also help

to promote the maintenance of SSCs. Recently obtained data show that PM cells express GDNF and can support SSC self-renewal in culture (L.-Y. Chen et al. 2014). The vasculature of the testis also appears to play an important role in stem cell renewal (Yoshida et al. 2007). Careful analysis using live cell imaging of mouse gonads showed that Asingle cells tend to reside close to the vascular network, while their differentiating daughters move away from these regions and disperse through the basal compartment of the testis. However, a more recent study on Id4+ SSCs showed that this population of SSCs does not associate with the vasculature (Chan et al. 2014). Therefore, caution should be taken in regarding the vasculature as a possible niche component. In addition, a valve-like terminal segment of the seminiferous tubules also supported SSC maintenance in hamster testis, suggesting niches may come in different varieties (Aiyama et al. 2015). Thus contributions of various different cell types might explain why not all Sertoli cells function as the SSC niche: the combination of signals from Sertoli cells, Leydig cells, macrophages and possibly additional cells may be required to fully define the functional niche.

Alternatively, it is possible that cell intrinsic fate determinant(s) are segregated during SSC divisions, conferring SSC identity to those that inherit the determinants. In this scenario, GDNF-expressing Sertoli cells could be the only population that provides SSC niche function, which determines SSC identity in combination with cell-intrinsic fate determinants within SSCs themselves.

The most challenging question is how a single population of Sertoli cells can simultaneously regulate SSC and differentiating germ cells. Because tight junctions have been observed between Sertoli cells and germ cells, it is tempting to speculate that at each stage, germ cells are subjected to spatially segregated and distinct signaling events. In combination with this type of compartmentalization, secretion from Sertoli cells may also be polarized, i.e. GDNF is only secreted toward the SSC area, whereas other factor(s) are secreted toward a different domain of the Sertoli cell surface. Alternatively, germ cells and/or Sertoli cells may extend distinct set of nanotubes/cytonemes to mediate specific signaling (see the section "Protrusion mediated access to ligand source"). Considering that *Drosophila* trachea air sac primordium extend distinct sets of cytonemes (FGF-specific cytonemes and Dpp-specific cytonenes) toward different target cells (Roy et al. 2014), this remains a distinct possibility.

## **What restricts niche signaling?**

As described above, many signaling pathways have been shown to function in GSC systems. However, the mechanisms that restrict niche signaling to foster the appropriate number of stem cells needed for tissue homeostasis under different environmental conditions remain poorly understood. Recent work using simple model systems may provide important clues as to the types of mechanisms that may be utilized to limit signaling in different contexts. Below, we describe several biological processes that can modulate the range of the niche signaling within model systems.

#### **i) Tissue geometry**

Tissue architecture, and more specifically the exact spatial positioning of cells relative to one another, can dictate cell fate. As illustrated in Figure1, GSCs in model systems directly

adhere to their niche cells. Drosophila GSCs typically align their spindles perpendicularly toward the hub or cap cells, placing one daughter cell in direct contact with the niche while displacing the other daughter away from the niche (Yamashita, Jones, and Fuller 2003). This positioning ensures an asymmetric outcome of the stem cell division, i.e. self-renewal and differentiation (Figure 1B). Given the proximity of GSCs and their differentiating daughter cells, the effective range of niche signals has to be tightly restricted. Considering that many niche ligands, including Dpp, act over a long range  $(\sim 100 \,\mu m)$  in other contexts, such as developing imaginal discs, mechanisms that limit the effective range of these ligands within the niche to 1 cell diameter  $(\sim 7 \mu m)$  must be in place.

#### **ii) Juxtacrine or contact dependent signaling**

By its very nature, contact-dependent or "juxtacrine" signaling allows for highly selective cell-to-cell communication. The Notch pathway (reviewed in (Kopan and Ilagan 2009)) represents one of the best studied examples of juxtacrine signaling. Notch and its ligands (Delta and Serrate in Drosophila) are both transmembrane proteins. Thus, activation of the pathway occurs only when communicating cells are in direct contact with one another. These molecules are not released into the extracellular space, minimizing the possibility of ectopic signal activation. The Notch pathway functions in a number of stem cell niches (Liu et al. 2010). For example, within the C. elegans gonad, Notch signaling keeps GSCs in an undifferentiated state. Within this system, maintenance of undifferentiated germ cells depends on the repression of three pathways: gld-1, gld-2 and a third meiotic entry pathway that remains poorly understood (Kadyk and Kimble 1998)(Eckmann et al. 2004)(Hansen et al. 2004)(Fox et al. 2011). The DTC expresses the Notch ligand LAG-2, while the germline expresses the receptor GLP-1. Notch pathway activation within germ cells induces the transcription of a number of target genes that act in concert with additional factors to repress the ability of germ cells from entering meiosis through the aforementioned pathways (Brenner and Schedl 2016). Thus, by using the Notch pathway, the DTC directly and precisely regulates the size of the GSC population (Byrd and Kimble 2009)(Kimble 2014). Notch signaling also acts in Drosophila gonads to control GSC numbers. In this case, the Notch pathway does not mediate direct communication between niche cells and GSCs, but rather acts during the formation of the niche itself. In the developing *Drosophila* ovary, expression of the Notch ligand Delta by developing terminal filament cells induces pathway activation in immediately adjacent somatic cells, specifying them to become cap cells (Song et al. 2007). Limited cell-cell communication is essential for the formation of properly sized niches. For example, ectopic activation of the Notch pathway in a greater number of cells within the developing gonad leads to the formation of ectopic niches and the inappropriate expansion of the GSC population in adults (Ward et al. 2006)(Song et al. 2007). Similarly, during male gonad development, Notch pathway regulates differentiation of somatic gonadal precursors (SGPs), precursors of hub cells. Notch pathway within SGPs acts together with EGF pathway to signal from primordial germ cells (PGCs) to determine appropriate niche size (Kitadate and Kobayashi 2010).

Notably, Notch is not the only example of a juxtacrine signal: several ligands once thought to function as secreted factors were later found to act in a juxtacrine manner. For example, pro-TGF-α, tethered to the plasma membrane of a mouse bone marrow stromal cell binds to

EGFR on an adjacent hematopoietic progenitor cell (Anklesaria et al. 1990). In addition, bone marrow niche cells lacking membrane-bound Steel factor (ligand for c-Kit), despite retaining the expression of a secreted form, failed to maintain HSCs (Barker 1997)(Ding et al. 2012). Several other molecules including cytokines and growth factors (e.g. c-Kit, TGF-α and Amphiregulin) also act in a juxtacrine manner in specific contexts (reviewed in (Singh and Harris 2005)). In all these examples, juxtacrine signaling guarantees that cell-cell communication will be spatially restricted to those neighbors that immediately contact one another, thus making them an ideal candidate to participate in the type of spatially-limited signaling observed in most stem cells niches.

#### **iii) Limit the amount of ligand production and/or secretion**

Several potential mechanisms may control the range of niche signaling despite relying on the secreted ligands. For example, the simple modulation of ligand production at the level of transcription or translation could influence the local signaling gradient. In addition, the availability of niche ligands can also be regulated at the level of secretion. Exocytosis represents a highly regulated process, both in terms of the amount and the subcellular location of the molecules targeted for secretion. Polarized exocytosis play an important role in most eukaryotic cells (He and Guo 2009). A multi-protein complex, the exocyst, is required for this polarized exocytosis. Exocyst localizes to sites of active exocytosis and mediates the targeting and tethering of post-Golgi vesicles to the plasma membrane prior to membrane fusion. In the Drosophila male GSC niche, Michel et. al. showed that BMP secretion and E-cadherin membrane targeting require exocytosis and recycling endosomes in the GSC niche (Michel et al. 2011). They observed the colocalization of E-Cadherin and BMP ligand at adherens junctions within the niche cell membrane. It is still unclear if adherens junctions are functionally required for BMP ligand secretion or not. Regardless, this study suggests that specific mechanisms regulate the secretion of ligands, potentially allowing for the precise control of signal availability to stem cells within the niche.

#### **iv) Modulation of the ligand diffusion outside the ligand-producing cells**

Regulating how a ligand diffuses through a tissue represents another potential mechanism by which to modulate the range of signaling. The extracellular matrix (ECM) can influence how far a ligand can travel from its source. In addition, several ECM components function as reservoirs of ligands: ECM components bind to soluble ligands to regulate their diffusion characteristics, thus regulating local availability of the ligands (reviewed in (Hynes 2009)). For example, fibronectin, vitronectin, collagens and proteoglycans are known to bind BMPs and other growth factors including FGFs and HGFs. ECM molecules can influence the solubility and activity of these ligands. Enzymes, such as metalloproteinases (Matrix metalloproteinases: MMP), induce the remodeling of ECM components and permit the release of factors as necessary (Hynes 2009). The *Drosophila* HSPG (heparan sulfate proteoglycan) protein Dally is essential for concentrating Dpp molecules on the surface of cells in wing discs (Akiyama et al. 2008). In the female GSC niche, Dally is specifically expressed in niche cells to ensure a high-level of BMP signaling, and thus promotes GSC identity (Guo and Wang 2009). HSPG is thought to function as a co-receptor to activate the receptor on the surface of target cells. In this manner, HSPG enhances the specificity between ligand-producing and –receiving cells. Strikingly, ectopic expression of Dally in the

Drosophila ovary expands the number of undifferentiated germline stem cells, suggesting that Dally influences the range of niche signaling (Hayashi, Kobayashi, and Nakato 2009). Similarly, in the male GSC niche, the secreted ECM protein Magu/Pentagone (Pent) is specifically expressed in hub cells and modulates Dpp signal activation exclusively in the GSC population (Zheng et al. 2011). Notably, ECM proteins do not always restrict ligand availability, but can also increase the distance over which signals act. Type IV collagens bind to Dpp and regulate BMP signaling in both the *Drosophila* embryo and ovary (Akiyama et al. 2008)(Guo and Wang 2009). Interaction between Dpp and type IV collagen appears to promote long-range gradient formation in the embryo, while it restricts the range of BMP pathway activation in the ovary through sequestration of the Dpp ligand.

#### **v) Protrusion mediated access to ligand source**

Our recent discovery of MT (microtubule based)-nanotubes suggests another potential mechanism that influences which cells can respond to niche signals (Figure 2A) (Inaba et al. 2015). MT-nanotubes are MT-based protrusions that extend from GSCs into the hub cell area. MT-nanotubes are sensitive to fixation similar to other thin protrusions reported to date, such as cytonemes and tunneling nanotubes, explaining why they have escaped detection in previous studies. 3D reconstitution of confocal stacks revealed that the MTnanotubes invaginate into the hub cell cluster but do not breach the membranes of hub cells. Double plasma membranes from both cells appeared to wrap around the core microtubule bundle extending from the GSC. Tkv receptor protein expressed within GSCs translocates to the tips of MT-nanotubes, where it interacts with Dpp ligand expressed by hub cells (Figure 2A, B). Dpp ligand fused to mCherry expressed within hub cells exhibits a punctate pattern within hub cell-hub cell junctions, likely representing the point where MT-nanotubes foster efficient signal reception (Inaba et al. 2015, Figure 2B). IFT (intraflagellar transport) proteins appeared to participate in MT-nanotube formation and/or maintenance. Perturbation of MT-nanotubes compromises activation of Dpp signaling within GSCs, leading to GSC loss, indicating that MT-nanotubes promote signal reception (Inaba et al. 2015). Similar to the cytonemes whose formation and/or stabilization requires ligand-receptor interactions (Roy et al. 2014), MT-nanotube formation depends on interactions between Dpp and Tkv. Taken together, these data suggest that GSCs sense niche produced ligands and extend/ stabilize MT-nanotubes towards the source of these signals. MT-nanotube formation in turn allows GSCs around the hub to experience the necessary threshold of signal activation needed for their self-renewal. MT-nanotube-mediated signaling represents one of the first examples in which cells utilize cellular protrusions to foster short-range signaling. These, and other similar structures, may promote efficient signal transduction in response to a limited amount of ligand produced by a local source, such as niche cells. By contrast, MTnanotubes do not promote stem cell self-renewal via Jak/Stat signaling between hub cells and GSCs. The specificity of MT-nanotubes for BMP signaling suggests that stem cells likely use multiple mechanisms to receive signals from the niche.

Other examples of protrusion mediated niche-stem cell regulation have been reported. For example, cap cells express Hedgehog protein and use short filopodia (cytonemes) to transport this ligand to escort cells (Rojas-Ríos, Guerrero, and González- Reyes 2012). In addition co-cultured osteoblast and human hematopoietic progenitor cells form long distance

cytoplasmic connections (tunneling/membrane nanotubes; TNTs), which mediate trafficking of SARA endosomes between cells to regulate Smad signaling (Gillette et al. 2009). These observations give rise to an intriguing possibility that cytoplasmic contents may be directly transported between niche cells and stem cells. However, it remains unclear whether TNTs actually function to foster *in vivo* niche-stem cell interactions.

#### **vi) Ligand diffusion vs. contact-dependent/protrusion-mediated signaling?**

Integration of protrusion-mediated signaling and ECM-mediated diffusion may serve to finetune the delivery of niche ligands to stem cells. For example, the ECM may help to increase the local concentration of ligands specifically around protrusions, making the protrusionmediated restriction of ligand delivery even tighter. Whether both mechanisms function together in the same niche or whether these mechanisms play distinct roles in the regulation of signaling remains an open question.

Many niches rely on signaling molecules that are presumably secreted into the extracellular space. A potential benefit of using diffusible ligands within niches includes the ability to adaptively adjust stem cell numbers in response to physiological change. For example, if niche signaling solely depended on juxtacrine signaling, re-establishment of the stem cell population after stem cell loss would be difficult. However, by using diffusible ligands, niches can influence cell fate at a distance, potentially allowing for the dedifferentiation of otherwise differentiating cells. Dedifferentiation has been observed in a number of systems including the Drosophila ovary and testis (Kai and Spradling 2004)(Brawley and Matunis 2004). Therefore, diffusible ligands may allow adaptable stem cell regeneration, if controlled properly. In such a context, ECM-mediated control of ligand diffusion/concentration may complement protrusion-dependent restriction of the niche signaling to maintain long-term tissue homeostasis.

#### **vii) Intrinsic factors that mediate the ability of a cell to respond to a signal**

Intrinsic factors within individual cells also help to sharpen the boundary between which cells experience signal transduction in response to ligands and which do not. For example, in the differentiating daughters of female GSCs (Xia et al. 2010) and male GSCs (Chang et al. 2013), BMP signaling is actively repressed by the HECT-domain ubiquitin E3 ligase SMAD ubiquitination regulatory factor (Smurf), which targets Tkv for degradation. smurf mutant ovaries exhibit an expansion of GSC-like cells outside of the niche (Xia et al. 2010), indicating that the degradation of Tkv promotes female germ cell differentiation. Likewise, smurf mutant testes also show increased/expanded Mad phosphorylation and an increased number of GSCs and transit-amplifying cell divisions (Chang et al. 2013). These results indicate that prompt inactivation of Dpp signaling is essential for timely differentiation of germ cells. However, in both cases, how the degradation activity is regulated differently between stem cells and their differentiating daughters remains unclear, and thus there must be additional mechanism(s) for decoding a cells' location within the tissue.

## **Concluding remarks**

Our understanding of potential regulatory mechanisms that control communication between niche cells and stem cells has greatly improved with the study of model organisms including Drosophila GSCs. These studies revealed remarkable complexity and precision in signaling mechanism regulating stem cell identity, differentiation and asymmetric divisions. At the same time, these studies have raised many interesting questions that will be addressed in future work. How are multiple mechanisms that control cell-cell signaling integrated into a single asymmetric event? Which event happens first? How flexible is the system? Does the effective range of signaling change to adapt to developmental and physiological changes? Are these mechanisms mutually dependent or do they provide redundancy to protect against the failure of one another? What ultimately happens when the spatial specificity of signaling is disturbed? These and other interesting questions about niche regulation await future study.

### **Acknowledgments**

Grant sponsor: National Institutes of Health, grant number: R01AG047318 (to MB), 1R01GM118308-01 (to YY)

Grant sponsor: Howard Hughes Medical Institute

We thank Yamashita and Buszczak lab members for discussion. The research in the Yamashita laboratory is supported by Howard Hughes Medical Institute, National Institutes of Health (1R01GM118308-01). YMY is supported by MacArthur Foundation. MB is supported by the National Institutes of Health (R01AG047318)

## **Abbreviations**



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#### **Figure 1. Short range niche signaling in** *C. elegans* **and** *Drosophila* **gonadal niches**

(**A**) In C. elegans, one distal tip cell (DTC) forms the niche for germline stem cells located at the distal end. DTC extends long projections that contact stem cells. (**B**) Asymmetric fate determination of Drosophila germline stem cells (GSCs) largely depends on the differential placement of two stem cell daughters to distinct locations: cells within the niche self-renew while cells outside the niche differentiate. The niche cell cluster (hub cells in males, terminal filament and cap cells in females) provides signals for stem cell self-renewal to the juxtaposed stem cells, but not other daughter cells displaced 1 cell diameter away from niche cells (Gonialblast in male, Cystoblast in female).



#### **Figure 2. MT-nanotube mediated niche-stem cell signaling**

(**A**) Model for MT-nanotube mediated signaling. Dpp induces MT-nanotube formation, and receptor–ligand interaction occurs at the surface of MT-nanotubes, leading to signaling activation in GSCs. (**B**) Dpp-mCherry (red) expressed in hub cells together with GFP-α-Tubulin (green, hub cell cortex) using hub specific unpaired (Upd) promoter. DppmCherry forms punctae along hub cell cortex (arrowheads). Scale bar: 10μm. Entire hub area is encircled by white broken line. GSCs are attached to hub from surrounding area (not visible here).