Neuregulins are essential for spermatogonial proliferation and meiotic initiation in neonatal mouse testis

JiDong Zhang¹, Ko Eto¹, Asuka Honmyou¹, Kazuki Nakao², Hiroshi Kiyonari² and Shin-ichi Abé¹,*

SUMMARY
The transition from mitosis to meiosis is unique to germ cells. In murine embryonic ovaries and juvenile testes, retinoic acid (RA) induces meiosis via the stimulation by retinoic acid gene 8 (Stra8), but its molecular pathway requires elucidation. We present genetic evidence in vivo and in vitro that neuregulins (NRGs) are essential for the proliferation of spermatogonia and the initiation of meiosis. Tamoxifen (TAM) was injected into 14-day post-partum (dpp) Sertoli cell-specific conditional Nrg1Ser⁻ ⁻ mutant mice. TAM induced meiosis in neonatal testes, increased the number of STRA8-positive and TUNEL-positive cells, respectively. In testicular organ cultures from 5-6 dpp wild-type mice and cultures of their re-aggregated spermatogonia and Sertoli cells, FSH, RA [all-trans-retinoic acid (ATRA), AM580, 9-cis-RA] and NRG1 promoted spermatogonial proliferation and meiotic initiation. However, TAM treatment of testicular organ cultures from the Nrg1Ser⁻ ⁻ mutants suppressed spermatogonial proliferation and meiotic initiation that was promoted by FSH or AM580. In re-aggregated cultures of purified spermatogonia, NRG1, NRG3, ATRA and 9-cis-RA promoted their proliferation and meiotic initiation, but neither AM580 nor FSH did. In addition, FSH, RAs and NRG1 promoted Nrg1 and Nrg3 mRNA expression in Sertoli cells. These results indicate that in juvenile testes RA and FSH induced meiosis indirectly through Sertoli cells when NRG1 and NRG3 were upregulated, as NRG1 amplified itself and NRG3. The amplified NRG1 and NRG3 directly induced meiosis in spermatogonia. In addition, ATRA and 9-cis-RA activated spermatogonia directly and promoted their proliferation and eventually meiotic initiation.

KEY WORDS: Neuregulins, Retinoic acid, Meiotic initiation, Neonatal mouse testis

INTRODUCTION
Meiosis is a process that produces gametes with extensive genetic diversity. In testis and ovary spermatogonia and oogonia, respectively, undergo species-specific rounds of mitotic divisions followed by the initiation of meiosis. In the mammalian ovary, germ cells enter meiosis during embryogenesis, whereas those in the testis do not until puberty. Initiation of meiosis in developing ovaries, but not in testes, may involve environment cues (McLaren, 1984; McLaren et al., 1995).

Vitamin A is required for normal spermatogenesis in the form of retinoic acid (RA), which is an active metabolite of vitamin A and is essential for mammalian spermatogonial differentiation and meiotic initiation (Hogarth and Griswold, 2010). In vitamin A-deficient (VAD) animals, germ cells are depleted from the seminiferous tubules and only undifferentiated type A spermatogonia remain. Replacement of vitamin A reinitiates spermatogenesis (Van Pelt and de Rooij, 1990). RA regulates the initiation of meiosis in mouse embryonic ovaries and juvenile testes via the Stra8 gene (stimulated by retinoic acid gene 8) (Koubova et al., 2006; Bowles et al., 2006; Baltus et al., 2006; Anderson et al., 2008). CYP26B1, a P450 enzyme that catabolizes all-trans RA (ATRA) into inactive metabolites, is expressed more in male than in female gonads between 12.5 and 14.5 dpc, the crucial time when female, but not male, germ cells enter meiosis (Menke and Page, 2002). As germ cells enter meiosis precociously in the testes of Cyp26b1-knockout (KO) mouse embryos (Bowles et al., 2006), CYP26B1 might protect male gonads from the action of RA in embryos by reducing retinoid levels during fetal gonadal development (Bowles and Koopman, 2007). Fibroblast growth factor 9 (FGF9) produced in the fetal testis acts directly on germ cells and inhibits meiosis (Bowles et al., 2010). However, the molecular pathway in which RA induces meiosis remains to be clarified.

We have previously identified neuregulin (NRG) 1 in newt testes as a FSH-upregulated clone that is homologous to mouse NRG1 (Oral et al., 2008). Nrg1 is one of the Nrg genes (Nrg1-Nrg4) that belong to the epidermal growth factor (EGF) family, and play essential roles in the nervous system, heart and breast (Falls, 2003). The family of EGF receptors includes four closely related transmembrane tyrosine kinases: ERBB1 (EGFR – Mouse Genome Informatics), ERBB2, ERBB3 and ERBB4. NRG1 and NRG2 both bind ERBB3 and ERBB4, whereas NRG3 and NRG4 bind ERBB4, but not ERBB3 (Hynes et al., 2001). ERBB2 has no direct ligand, but works as a co-receptor. NRGs induce not only homodimers of ERBB3 and ERBB4, but also heterodimers of ERBB2/ERBB3 and ERBB2/ERBB4. As Nrg1 and Nrg3 mRNAs are expressed in mouse neonatal testes, in this report we analyzed the functions of NRG1 and NRG3 in the proliferation of spermatogonia and their meiotic initiation in relation to retinoic acid and FSH in vivo and in vitro. Because Nrg1 is expressed in Sertoli cells only, and pan-Nrg1 KO mice are known to be embryonic lethal (Falls, 2003), we generated conditional Sertoli cell-specific Nrg1Ser⁻ ⁻ mutant mice. RA acts through the nuclear
RA receptors (RARs) (α, β, γ) and retinoid X receptors (RXRs) (α, β, γ). RARs and RXRs may function as homodimers and heterodimers (RAR/RXR) (Mark and Chambon, 2003). As the expressed receptor isoforms of Sertoli cells and spermatogonia are different, we examined the effect of three kinds of retinoids on the proliferation of spermatogonia and their meiotic initiation in vitro: AM580, a stable retinoic acid receptor α (RARα)-specific agonist; ATRA, which can bind RARs; and 9-cis-RA, which can act on both RARs and RXRs (Mark and Chambon, 2003).

**MATERIALS AND METHODS**

**Mice**

C57BL/6j mice were purchased from JapanCLEA (Tokyo, Japan) and maintained on a 12-hour-light/12-hour-night schedule at constant temperature and humidity in the Center for Animal Resources and Development of Kumamoto University according to protocols for animal experiments approved by the Institutional Animal Care and Use Committee.

**Generation of conditional Nrg1 mutant mice**

Conditional Nrg1 mutant mice were generated as shown in Fig. S1 in the supplementary material.

**Injection of TAM into mutant and control mice**

TAM was injected daily into Nrg1<sup>−/−</sup> mutant and control mice (0.3 mg/10 g weight) for 5 days, and 1 month later the testes were fixed in Bouin’s solution.

**Organ cultures of testicular fragments from wild-type and mutant mice**

One testis from a 6 dpp wild-type mouse was cut into 16 fragments (about 1×1 mm). Eight fragments were placed on a float of nucleapore filter (Track-Etch Membrane, 25 mm, 0.2 µm pore size, Whatman) in a 35 mm plastic dish (Falcon; #1008) and cultured in Dulbecco’s Modified Eagle’s Medium (DMEM, 2 ml) containing 0.1% bovine serum albumin (BSA, Sigma) for 72 hours at 32°C with 5% CO2 aeration. Porcine FSH (National Hormone & Peptide Program [NHPP]), ATRA (1 µM; Sigma), AM580 (1 nM; Sigma), 9-cis-RA (1 µM; Biomol), or recombiant EGFR-domain of NRG1 or NRG3 was added to the culture dishes. Various inhibitors [AG879 (Calbiochem), an ERBB2 inhibitor; AG1478 (Wortmannin), an EGFR (ERBB1/ERBB4 inhibitor; and PD153035 (Biaffin), an ERBB1/ERBB4 inhibitor] were added to the medium 3 hours prior to the addition of FSH, NRGs or RA. Two to six mice were used for one experiment, depending on the number of different reagents added to the cultures.

Organ cultures of 5 dpp testes from Nrg1<sup>−/−</sup> mutant mice were produced as follows. One testis was cut into 12 fragments (about 1×1 mm) and only six dishes (four fragments/dish) could be cultured from one mutant mouse. TAM (100 ng/ml) was added to the medium on day 0, followed 1 day later by the addition of FSH, ATRA, AM580, 9-cis-RA, NRG1 or NRG3, and cultured for 6 days.

**Cell fractionation and re-aggregated cultures**

Re-aggregated cultures of spermatogonia and Sertoli cells were produced as follows. After the tunica albuginea was removed from 6 dpp testes, whole testes were treated with 0.1% collagenase (type N-2, Nitta Gelatin, Japan) for 30 minutes at 25°C. The supernatant was used as a Leydig cell-rich fraction. The tubules were then dissociated by 0.1% collagenase and 1.5 KU/ml DNase I (type IV, Sigma) for 1 hour at 25°C, followed by pipetting (~100 times). Then the tubules were dissociated completely by 0.05% trypsin containing 0.1 mM EDTA for 10 minutes at 37°C. The cell suspension was filtered through nylon gauze (50 µm) and washed in DMEM, and 5 ml of the cell suspension in DMEM was placed on a 15% Nycodentz (Sigma) solution in Krebs ringer (5 ml) and centrifuged at 110 g for 3 minutes. Dead cells sedimented at the bottom of the tube. The layer formed between DMEM and 15% Nycodentz was collected as live germ and Sertoli cells, and washed with DMEM three times. The dissociated cells were re-aggregated by rotation culture (70 rotations/minute, R-30, TAIJEC, Koshigaya, Saitama, Japan) for 1.5 hour at 25°C, followed by centrifugation in a 0.5 ml siliconized tube (Assist, Japan) (~10<sup>6</sup> cells/tube) at 230 g for 5 minutes. The cell pellet formed was detached from the tube by a needle in 50 µl medium, to which DMEM (200 µl) containing 0.24 mg/ml collagen (Cellmatrix type I-P, Nitta Gelatin) was added. The solution containing a re-aggregate was then placed on a nucleopore filter for 1 hour until the collagen hardened and the filter was floated on the medium as an organ culture. The medium was changed after 3 days of culture.

The fractionation of Sertoli cells and spermatogonia was performed as follows. After the testes were dissociated by collagenase and the cell suspension was filtered through nylon gauze, the supernatant (5 ml) was placed on 5% Nycodentz in Krebs ringer (5 ml) that was underlayered by 15% Nycodentz (5 ml) and centrifuged at 110 g for 3 minutes. The cells recovered from the layer formed between 5% Nycodentz and DMEM were used as a Sertoli cell-rich fraction, and the cells between 5% and 15% Nycodentz were used as a germ cell-rich fraction. Each fraction was further purified by seeding on dishes coated with collagen (type I; Nitta Gelatin) and incubated at 32°C. For the Sertoli cell-rich fraction, after the floating germ cells were removed daily for 3 days, the attached Sertoli cells were used directly as purified Sertoli cells (purity 94.3±2.6%). For the germ cell-rich fraction, the floating germ cells were collected after 6 hours incubation in collagen-coated dishes and, after pipetting, inoculated again into another collagen-coated dish. Then the supernatant spermatogonia were recovered after 6 hours incubation as purified spermatogonia (purity 90.3±2.7%). Re-aggregated cultures containing only spermatogonia were produced in the same way as that described above for spermatogonia and Sertoli cells.

**Histology, and analysis of spermatogonial proliferation and differentiation**

Cultured testicular fragments, re-aggregates and whole testes were fixed in Bouin’s solution and dehydrated in a graded ethanol series. All testicular fragments cultured in one dish were embedded together in paraffin wax. Sections were made serially at 5 µm thickness, treated with xylene, dehydrated in an ethanol series and stained with Delafield’s Hematoxylin followed by Eosin.

To assay for spermatogonial proliferation, 5-bromo-2-deoxy-uridine (BrDU) was injected intraperitoneally (0.3 mg/10 g weight) or incubated in the cultures, for 3 hours prior to fixation and processed for immunohistochemistry with a kit according to the manufacturer’s instructions (Amersham Pharmacia Biotech, Buckinghamshire, England). The number of positive cells was counted in three photographed areas (157,500 µm<sup>2</sup>/area), taken at random in a section (in vivo) containing three or four fragments (in organ culture) × three sections, each derived from five or six sections. Sertoli cells and germ cells were identified by cell size, nuclear morphology and localization within the seminiferous tubules (in the case of in vivo and organ culture), according to the descriptions by Huckins and Oakberg (Huckins and Oakberg, 1978) and Russell et al. (Russell et al., 1990).

The proliferative activity and extent of differentiation were expressed as the number (mean±s.e.m.) of BrDU-positive cells and SYCP3 (synaptosomal complex protein 3)-positive cells/tube/section for independent experiments. For re-aggregated cultures the number (mean±s.e.m.) of BrDU-positive cells and SYCP3-positive cells/100,000 µm<sup>2</sup>/section × three sections from three independent experiments indicated their proliferative activity and extent of differentiation.

**Immunofluorescence**

Testes, cultured testicular fragments or re-aggregates were fixed in Bouin’s and embedded in paraffin wax. Sections (5 µm) were boiled in 0.01 M citrate (pH 6.0), washed three times in 0.1% Tween20/PBS, transferred to blocking solution containing 3% BSA and 10% goat serum (without serum in case of goat anti-GATA4) in 0.1% Triton-PBS for 1 hour, and incubated with primary antibody at 4°C overnight. After washing, the secondary antibody was added and the sections were incubated for 2 hours at room temperature. Combinations of the first antibody and the second antibody were as follows: rabbit anti-SYCP3 polyclonal antibody (Abcam, 1:200) or rabbit anti-STR A8 (Abcam, 1:200) and Alexa Fluor 488 goat anti-rabbit.
Expression and purification of the EGF-like domain of recombinant NRGs

Expression and purification of the EGF-like domain of recombinant NRG1 and NRG3 was performed as previously reported (Eto et al., 2010). The sense and antisense primers were 5′-CAT ATG ACC AGC CAT CTC ATA AAG TGT GC-3′ (NRG1) and 5′-CAT ATG TCT GAG CAC TTC AAA CCC TGT C-3′ (NRG3), and 5′-CTC GAG TTA CTC CGC TTC CAT AAA TTC-3′ (NRG1) and 5′-ATT CCT CGA GCT AGT GGT CTG TTG GA-3′ (NRG3).

Statistics

Cell proliferation and differentiation activities were analyzed using a Student’s t-test. P values less than 0.05 were considered to be statistically significant.

RESULTS

Expression of mRNA for Nrg and Erbb genes in neonatal testes

mRNAs of Nrg1, Nrg3, Erbb1, Erbb2, Erbb2/2 and Erbb4 were expressed in the neonatal testes during the first 8 weeks (see Fig. S2A in the supplementary material), but mRNAs for Nrg2, Nrg4 and Erbb3 were barely detected. RT-PCR analysis of mRNAs from fractionated Sertoli cells, Leydig cells and spermatogonia from 6 dp testes, where anti-Mullerian hormone (Amh), luteinizing hormone receptor (Lhr; Lhcgr – Mouse Genome Informatics) and A-kinase anchor protein 12 (Akap 12) were used as markers for the respective cell types, showed that Nrg1 and Erbb2 were expressed only in Sertoli cells, while Nrg3, Erbb1, Erbb2/2 and Erbb4 were expressed in both somatic cells (Sertoli cells and Leydig cells) and spermatogonia (see Fig. S2B in the supplementary material).

Tamoxifen injected into Sertoli cell-specific Nrg1Ser−/− mutant mice induces testis degeneration

To examine the functional role of NRG1 in spermatogenesis, we generated Sertoli cell-specific Nrg1Ser−/− mutants (Accession Number CDCB0743K: http://www.cdb.riken.jp/aug/mutant%20mice%20list.html), in which both alleles of Nrg1 exon 6 were excised.

Table 1. Sequences of primers, number of cycles and temperature by which PCR was performed for each gene

<table>
<thead>
<tr>
<th>Gene</th>
<th>Forward primer</th>
<th>Reverse primer</th>
<th>Cycles</th>
<th>Temperature (°C)</th>
<th>Reference or Accession Number</th>
</tr>
</thead>
<tbody>
<tr>
<td>Nrg1</td>
<td>GCAGCTGGAGGGCAAAGAGAGG</td>
<td>CGTACCTTGACAAAGATTC</td>
<td>40</td>
<td>54</td>
<td>NM_178591.2</td>
</tr>
<tr>
<td>Nrg3</td>
<td>GGCTAGCGCGGATGGAAC</td>
<td>GTCCTCATCGGGCTACCC</td>
<td>40</td>
<td>58</td>
<td>NM_008734.2</td>
</tr>
<tr>
<td>Erbb1</td>
<td>GTGACATGCTGGTCCAGTCAGT</td>
<td>CTAGTGCATTTGTTCTGCTGC</td>
<td>40</td>
<td>58</td>
<td>NM_007912.4</td>
</tr>
<tr>
<td>Erbb2</td>
<td>GACTACGTGTCAGTCCAGC</td>
<td>TGGTTGTTCTTCTAGTGA</td>
<td>40</td>
<td>54</td>
<td>NW_00130424.1</td>
</tr>
<tr>
<td>Erbb2/2</td>
<td>CAATCGTTGGGCGCTGTCGC</td>
<td>GAGTGACAGGACTCACCCTC</td>
<td>40</td>
<td>54</td>
<td>NW_001003817.1</td>
</tr>
<tr>
<td>Erbb3</td>
<td>GTGACTTATACCTCAGGAAC</td>
<td>GTTACACAGTCTAGACCC</td>
<td>40</td>
<td>54</td>
<td>NM_010153.1</td>
</tr>
<tr>
<td>Erbb4</td>
<td>GACCATGTGCTGGTGTCCG</td>
<td>CATGATGACAGGATGAAG</td>
<td>40</td>
<td>54</td>
<td>NM_010154.1</td>
</tr>
<tr>
<td>β-Tubulin</td>
<td>TTTCTGGGAGGCTCATAAGGATG</td>
<td>GGGAAAGGGGGGAGGTGTG</td>
<td>25</td>
<td>58</td>
<td>NM_011655.4</td>
</tr>
<tr>
<td>Akap 12</td>
<td>GCTAAGGGCTTGGTACGACA</td>
<td>GTGACTGCTGCACCTTCTAG</td>
<td>40</td>
<td>58</td>
<td>NM_0013185.3</td>
</tr>
<tr>
<td>Amh</td>
<td>TGACAGTGGAGGAGGAGG</td>
<td>TATCCACTCCACGCAGATGGA</td>
<td>40</td>
<td>56</td>
<td>Guyot et al., 2004</td>
</tr>
<tr>
<td>Fshr</td>
<td>CGGCAAACTCTGAACCTACAATACCT</td>
<td>GTGTTAGTGGCTGTGAGCAGA</td>
<td>35</td>
<td>56</td>
<td>NM_013523.2</td>
</tr>
<tr>
<td>Rara</td>
<td>AGAAGCCTCATCAACAGAAGGA</td>
<td>TCCCGTGTGGCTGCTGTTGTCG</td>
<td>28</td>
<td>55</td>
<td>Volle et al., 2007</td>
</tr>
<tr>
<td>RarB</td>
<td>TGCCAGGAGGGCAGGGAGG</td>
<td>GCTGAGGTGCTGGGAGCAGC</td>
<td>28</td>
<td>55</td>
<td></td>
</tr>
<tr>
<td>RarC</td>
<td>GCAGCTGGGAGGGAAGAGAGG</td>
<td>CATGATGGTGGTACGACCC</td>
<td>40</td>
<td>60</td>
<td>Bek et al., 2003</td>
</tr>
<tr>
<td>RarX</td>
<td>GTTTCTGGGAGGGAAGAGAGG</td>
<td>CATGATGGTGGTACGACCC</td>
<td>28</td>
<td>55</td>
<td></td>
</tr>
<tr>
<td>RarX</td>
<td>CTAGCTGGAGGGAAGAGAGG</td>
<td>CATGATGGTGGTACGACCC</td>
<td>28</td>
<td>55</td>
<td></td>
</tr>
<tr>
<td>Sard</td>
<td>AATAGGGCTGGAAAGAGAGG</td>
<td>CATGATGGTGGTACGACCC</td>
<td>40</td>
<td>60</td>
<td></td>
</tr>
<tr>
<td>Str8</td>
<td>AGCTGGGAGGGGAGGAGG</td>
<td>CATGATGGTGGTACGACCC</td>
<td>40</td>
<td>60</td>
<td></td>
</tr>
<tr>
<td>Cyclophilin</td>
<td>GGCTCCTGACATCATCAACAGA</td>
<td>TCCCGTGTGGCTGCTGTTGTCG</td>
<td>28</td>
<td>55</td>
<td>Volle et al., 2007</td>
</tr>
<tr>
<td>Nrg1-lop</td>
<td>GCTGCGCCTGCTAGGCTGAGT</td>
<td>GCTGAGGTGCTGGGAGCAGC</td>
<td>28</td>
<td>55</td>
<td></td>
</tr>
<tr>
<td>Mis cre ER</td>
<td>AGCTGGGAGGGGAGGAGG</td>
<td>CATGATGGTGGTACGACCC</td>
<td>40</td>
<td>60</td>
<td>Hofmann et al., 2003</td>
</tr>
</tbody>
</table>
in Sertoli cells when tamoxifen (TAM) was added, by crossing mice carrying loxP-flanked alleles of Nrg1 exon 6 with mice bearing the MIS-CreER transgene (see Fig. S1 in the supplementary material). Mutant mice (14 dpp) containing primary spermatocytes were injected with TAM daily for 5 days and 1 month later the testes were fixed. The testicular size and weight in mutant mice were remarkably reduced compared with that in heterozygous mice (Fig. 1A,B). Histological sections showed that vacant areas appeared in the central or peripheral regions of seminiferous tubules (44.3±3.0%) of mutant testes (Fig. 1D, D'), whereas no such areas were observed in those of the heterozygous mice (Fig. 1C). In some cases, only Sertoli cells and undifferentiated spermatogonia remained (Fig. 1D), whereas in other cases Sertoli cells, primary spermatocytes and spermatids were present, but not spermatogonia (Fig. 1D'). Some moribund and dead cells were observed; they were probably spermatogonia (Fig. 1F,H) or primary spermatocytes (Fig. 1J,L), because such live cell types were nearby and also those were the cell types in seminiferous tubules of heterozygous mice (Fig. 1E,G,I,K). This interpretation is consistent with the fact that most of the TUNEL-positive cells were located in the central and peripheral regions of the tubules (Fig. 1N,O). The numbers of TUNEL-positive cells/tubule and TUNEL-positive tubules/section in mutant testes increased to more than twice that in heterozygous mouse testes (Fig. 1P,Q), indicating that NRG1 is required for the survival of spermatogonia and/or primary spermatocytes. The numbers of BrdU-labeled cells/tubule and of BrdU-positive tubules/section in mutant testes were less than half of those in heterozygous mouse testes (Fig. 1R-U), indicating that NRG1 is indispensable for the proliferative activity of spermatogonia and/or DNA synthesis of pre-leptotene spermatocytes. As the majority of STRA8-immunopositive cells in adult testes are pre-leptotene/early leptotene spermatocytes (Zhou et al., 2008b), we compared the number of STRA8-positive cells in the mutant testes with that in heterozygous testes. The numbers of STRA8-positive cells/tubule and of STRA8-positive tubules/section in mutant testes were less than half of those in heterozygous testes (Fig. 1V-Y). In addition, the percentage of BrdU-positive cells in the STRA8-positive cells in mutant testes was significantly lower than that in heterozygous mouse testes (Fig. 1Z). These results indicate that NRG1 plays an important role in the differentiation of spermatogonia into pre-leptotene spermatocytes and their DNA synthesis.

Finally, examination of Hematoxylin and Eosin-stained sections revealed several differences in cell types between mutant and wild-type mice (Fig. 1M): (1) the numbers of Sertoli cells per seminiferous tubule in mutant and heterozygous testes were almost
the same; and (2) the number of undifferentiated spermatogonia per 1000 Sertoli cells was almost the same in mutant and heterozygous testes; but (3) the numbers of differentiating type A and type B spermatogonia and pre-leptotene and leptotene-zygotene primary spermatocytes per 1000 Sertoli cells in mutant testes were remarkably lower than those in heterozygous testes; (4) the ratio of differentiating type A spermatogonia to undifferentiated type A spermatogonia in mutant testes (1.15) was remarkably reduced compared with that in heterozygous testes (1.75); (5) the ratio of pre-leptotene primary spermatocytes to type B spermatogonia in mutant testes (1.34) was remarkably lower compared with that in heterozygous testes (1.78); however, (6) the ratio of type B spermatogonia to differentiating type A spermatogonia and that of leptotene-zygotene primary spermatocytes to pre-leptotene primary spermatocytes were similar in mutant testes (1.31 and 2.78, respectively) and heterozygous testes (1.25 and 2.35, respectively). These results indicate that development from the undifferentiated type A spermatogonia to the differentiating type A spermatogonia was blocked and/or that the proliferative activity of the differentiating type A spermatogonia was inhibited in the mutant testes. In addition, we conclude that the process from type B spermatogonia to pre-leptotene primary spermatocytes was largely suppressed in the mutant testes.

**FSH, RA and NRG1 promote spermatogonial proliferation and meiotic initiation in organ culture of wild-type mouse testes**

As the above results indicate that NRG1 is indispensable for the proliferative activity of spermatogonia and/or DNA synthesis of pre-leptotene spermatocytes and/or survival of germ cells, we examined the roles of RA and NRG1 in the proliferation of spermatogonia and their entrance into meiosis in 3-day organ cultures from 6 dpp testes when differentiating A-type spermatogonia were the most advanced stage. Recombinant EGF-like domains of NRG1 and NRG3, sufficient for binding and activating receptors to induce cellular responses (Holmes et al., 1992; Wen et al., 1994), promoted spermatogonial proliferation in a dose-dependent manner (Fig. 2A). Both AM580 and 9-cis-RA...
stimulated the proliferation of spermatogonia to an extent similar to NRG1, whereas FSH strongly stimulated their proliferation to more than double that of the control number (Fig. 2B). Only FSH, but neither NRGs nor RAs stimulated the proliferation of Sertoli cells (Fig. 1C).

To determine whether NRG1 and RA promote the differentiation of spermatogonia into primary spermatocytes, we examined in organ cultures the nuclear morphology and the expressions of Spool mRNA, which is specifically expressed in meiotic cells (Keeney et al., 1999) and the protein that it encodes is required for double-strand break formation and synopsis (Romanienko and Camerini-Otero, 2000), by RT-PCR. We also examined SYCP3 (Heyting, 1996) and γ-H2AX (phosphorylated histone H2AX) (Richardson et al., 2004), both of which appear during early meiotic prophase, by immunofluorescence. In culture media, containing FSH, AM580 or NRG1 (Fig. 2F-H), some nuclei showed chromosomal condensation that was characteristic of the zygotene or pachytene stages, whereas no such nuclei were observed in the initial (Fig. 2D) or control (Fig. 2E) cultures. Spool mRNA was detected in the presence of FSH, AM580 or NRG1, whereas it was barely detectable in the initial or control cultures (Fig. 2I). In addition, SYCP3 (Fig. 2L-N) and γ-H2AX (Fig. 2P-R) proteins were expressed when FSH, AM580 or NRG1 was added, but not in the initial or control cultures (Fig. 2J,K). Extended chromosomes revealed that the SYCP3 protein was distributed throughout the chromosomes (Fig. 2O). The number of TUNEL-positive cells was few in the control (1.20±0.20 cells/tubule) and in the cultures to which AM580 (1.35±0.24 cells/tubule) and NRG1 (1.28±0.21 cells/tubule) (Fig. 2T-V) were added. Thus, FSH, AM580 and NRG1 stimulated differentiation of spermatogonia into primary spermatocytes in culture (Fig. 2S), indicating that both RA and NRG1 play a pivotal role in the initiation of meiosis.

Relationship between the NRGs/ERBB and the RA and FSH pathways

To determine the relationship between the NRGs/ERBB and the RA and FSH pathways, we first examined whether Nrg mRNA expression is activated by AM580 and FSH. Nrg1 and Nrg3 mRNA expression was stimulated by AM580 and FSH (Fig. 2W), indicating that NRG1 and NRG3 act downstream of RA and FSH. Next, we examined whether ERBB inhibitors suppress the spermatogonial proliferation that was stimulated by AM580, because, if RA operates upstream of the NRG pathway, ERBB inhibitors should suppress the effect of AM580. Although NRG1 can potentially bind to ERBB3 and ERBB4 (Holmes et al., 1992), no Erbb3 mRNA expression was detected in mouse testes (1-8 weeks) (see Fig. S2A in the supplementary material). Therefore, NRG1 may signal through homodimers of ERBB4 and/or heterodimers of ERBB2/ERBB4 and/or ERBB1/ERBB4. The effects of various inhibitors for ERBB1, ERBB2 and ERBB1/ERBB4 on spermatogonial proliferation in the presence of NRG1 revealed that both ERBB2 and ERBB1/ERBB4 inhibitors suppressed spermatogonial proliferation, whereas there was no significant inhibition without NRG1 compared with the control (Fig. 2X). These results indicate that NRG1 binds to ERBB4 and signals via ERBB2/ERBB4 and/or ERBB4/ERBB4. We then examined the effects of the inhibitors on spermatogonial proliferation in the presence of AM580 (Fig. 2X). Both ERBB2 and ERBB1/ERBB4 inhibitors significantly suppressed the proliferative activity that was promoted by AM580. These results indicate that NRG1 operates downstream of the RA signaling system.

Neither FSH nor AM580 promote meiotic initiation in testes from Sertoli cell-specific Nrg1Ser−/− mutants in the presence of TAM in vitro

To examine the functional role of NRG1 in spermatogonial proliferation and meiotic initiation, TAM or its vehicle was added to organ cultures of 5 dpp mutant testes. Then, 1 day later AM580, ATRA, 9-cis-RA, FSH, NRG1 or NRG3 was added and the testes were cultured for 6 days. Analysis of histological sections revealed only a few areas with pyknotic cells (data not shown). TUNEL staining also showed very few dead cells in the presence of the vehicle ±TAM (1.32±0.18 cells/tubule, 1.35±0.24 cells/tubule, respectively) and NRG1 ±TAM (1.38±0.22 cells/tubule, 1.42±0.20 cells/tubule, respectively). Thus, cell viability was very good. The proliferative activity of spermatogonia and their differentiation into primary spermatocytes were promoted by FSH, ATRA, AM580, 9-cis-RA, NRG1 and NRG3 in the absence of TAM (Fig. 3A-F). However, the proliferative activity of spermatogonia and their differentiation stimulated by FSH and AM580 were significantly suppressed by TAM (Fig. 3A,B,D,E). However, the addition of NRG1 or NRG3 promoted spermatogonial proliferation and their differentiation to primary spermatocytes in organ cultures of 5 dpp mutant testes. Then, 1 day later AM580, ATRA, 9-cis-RA, FSH, NRG1 or NRG3 was added and the testes were cultured for 6 days. Analysis of histological sections revealed only a few areas with pyknotic cells (data not shown). TUNEL staining also showed very few dead cells in the presence of the vehicle ±TAM (1.32±0.18 cells/tubule, 1.35±0.24 cells/tubule, respectively) and NRG1 ±TAM (1.38±0.22 cells/tubule, 1.42±0.20 cells/tubule, respectively). Thus, cell viability was very good. The proliferative activity of spermatogonia and their differentiation into primary spermatocytes were promoted by FSH, ATRA, AM580, 9-cis-RA, NRG1 and NRG3 in the absence of TAM (Fig. 3A-F). However, the proliferative activity of spermatogonia and their differentiation stimulated by FSH and AM580 were significantly suppressed by TAM (Fig. 3A,B,D,E). However, the addition of NRG1 or NRG3 promoted spermatogonial proliferation and their differentiation to primary spermatocytes in organ cultures of 5 dpp mutant testes. Then, 1 day later AM580, ATRA, 9-cis-RA, FSH, NRG1 or NRG3 was added and the testes were cultured for 6 days. Analysis of histological sections revealed only a few areas with pyknotic cells (data not shown). TUNEL staining also showed very few dead cells in the presence of the vehicle ±TAM (1.32±0.18 cells/tubule, 1.35±0.24 cells/tubule, respectively) and NRG1 ±TAM (1.38±0.22 cells/tubule, 1.42±0.20 cells/tubule, respectively). Thus, cell viability was very good. The proliferative activity of spermatogonia and their differentiation into primary spermatocytes were promoted by FSH, ATRA, AM580, 9-cis-RA, NRG1 and NRG3 in the absence of TAM (Fig. 3A-F). However, the proliferative activity of spermatogonia and their differentiation stimulated by FSH and AM580 were significantly suppressed by TAM (Fig. 3A,B,D,E). However, the addition of NRG1 or NRG3 promoted spermatogonial proliferation and their differentiation to primary spermatocytes in organ cultures of 5 dpp mutant testes. Then, 1 day later AM580, ATRA, 9-cis-RA, FSH, NRG1 or NRG3 was added and the testes were cultured for 6 days. Analysis of histological sections revealed only a few areas with pyknotic cells (data not shown). TUNEL staining also showed very few dead cells in the presence of the vehicle ±TAM (1.32±0.18 cells/tubule, 1.35±0.24 cells/tubule, respectively) and NRG1 ±TAM (1.38±0.22 cells/tubule, 1.42±0.20 cells/tubule, respectively). Thus, cell viability was very good.

Fig. 3. Proliferative activity of spermatogonia and their differentiation to primary spermatocytes in organ cultures of mutant and heterozygous testes. In organ culture of 5 dpp testes from Nrg1Ser−/− mutants (A-F) and heterozygous mice (G-L), TAM or vehicle was added to culture medium, followed 1 day later by addition of various factors, and cultured for further 6 days. Data are mean±s.e.m.; *P<0.01.
differentiation even in the presence of TAM to almost the same level as when TAM was absent (Fig. 3C,F), indicating that NRG1 and NRG3 rescued the mutant phenotype. On the other hand, ATRA and 9-cis-RA also promoted the proliferative activity and the differentiation even in the presence of TAM (Fig. 3A,B,D,E). TAM had no effect on heterozygous mouse testes in littermates (Fig. 3G-L). These results indicate that NRG1 acts as a downstream factor of FSH and AM580 in promoting spermatogonial proliferation and the initiation of meiosis.

NRGs promote the initiation of meiosis in re-aggregated cultures of purified spermatogonia

To determine whether NRGs and/or RA stimulate spermatogonial proliferation and their differentiation into primary spermatocytes, we established cultures (1) consisting of spermatogonia and Sertoli cells that were re-aggregated, and cultures (2) consisting of only re-aggregated spermatogonia, within a collagen matrix. In the re-aggregated cultures of spermatogonia and Sertoli cells, AM580 and NRG1, as well as FSH, promoted spermatogonial proliferation (see Fig. S3A in the supplementary material), whereas only FSH stimulated the proliferation of Sertoli cells (see Fig. S3B in the supplementary material). In 7-day cultures, primary spermatocytes in the zygote-pachytene stage appeared in the media supplemented either with FSH, AM580 or NRG1 (see Fig. S3D-F in the supplementary material), but not in the control medium (see Fig. S3C in the supplementary material). Differentiation into primary spermatocytes was confirmed by the expression of SYCP3 (see Fig. S3G-J,O in the supplementary material) and γ-H2AX (see Fig. S3K-N in the supplementary material).

To determine whether NRGs and RA act on spermatogonia directly, NRG1, NRG3, AM580, ATRA and 9-cis-RA were added separately in re-aggregated cultures of spermatogonia and cultured for 1 week. NRG1, NRG3, ATRA and 9-cis-RA promoted spermatogonial proliferation, whereas AM580 did not (Fig. 4A). Zygotene-pachytene spermatocytes were identified by Hematoxylin and Eosin staining (Fig. 4B-G), and by immunofluorescence for SYCP3 (Fig. 4H-M) and γ-H2AX (Fig. 4N-P) in the presence of NRG1, NRG3 and ATRA, but were not seen in the presence of AM580 or FSH. NRG1 and NRG3, as well as ATRA, promoted the expression of mRNA for Spo11 and Stra8, while NRG1 and ATRA also upregulated Nrg3 mRNA, but NRG3 did not. Conversely, AM580 did not activate the three kinds of mRNA (Fig. 4R). Analysis of histological sections and TUNEL staining showed a few dead cells in the control and in the cultures in which NRG1 was added (see Fig. S4A-D in the supplementary material). That FSH did not promote spermatogonial proliferation (Fig. 4A) was consistent with the fact that the percentage of contaminated Sertoli cells was quite low (see Fig. S4E-G in the supplementary material). These results strongly suggest that NRG1, NRG3, ATRA and 9-cis-RA directly stimulated spermatogonial proliferation and the initiation of meiosis.

AM580, ATRA and FSH promote the expression of Nrg1 and Nrg3 mRNAs, whereas NRG1 amplifies itself and Nrg3 mRNAs in Sertoli cells

To determine the relationship between RA signaling and the NRGs/ERBB pathway in Sertoli cells, we examined the effects of FSH, AM580, ATRA, NRG1 and NRG3 on mRNA expression of Nrg1 and Nrg3 in Sertoli cells cultured for 3 days (Fig. 4S). FSH, AM580 and ATRA promoted the expression of Nrg1 and Nrg3 mRNAs. Interestingly, NRG1 also stimulated the expression of Nrg1 and Nrg3 mRNA, but NRG3 did not. These results indicate that the NRGs/ERBB system operates downstream of RA and FSH, and that NRG1 exerts an autocrine effect on the production of NRG1 and NRG3 in Sertoli cells (Fig. 5).

Fig. 4. Proliferative activity of spermatogonia and meiotic initiation in re-aggregated cultures of purified spermatogonia after 7 days of culture. (A) Proliferative activity of spermatogonia by various factors. Data are mean±s.e.m.; *P<0.01. (B-G) Hematoxylin and Eosin staining of the sections show meiotic cells (arrows). (H-P) SYCP3 (H-M) and γ-H2AX (N-P) expressed in the presence of NRG1 (K,N), NRG3 (L,O) and ATRA (M,P). (Q) Number of SYCP3-positive cells/105 μm² that differentiated in the presence of various factors. Data are mean±s.e.m. (R) RT-PCR shows expressions of mRNAs for Spo11, Stra8 and Nrg3 by various factors in purified spermatogonia. Cyclophilin was used as an internal control. (S) RT-PCR shows expressions of mRNAs for Nrg1 and Nrg3 by various factors in purified Sertoli cells cultured for 3 days.
aggregated cultures containing only spermatogonia (Fig. 4) indicate ERBB4 expressed on spermatogonia, in promoting spermatogonial only. Thus, NRG1 and NRG3 play a pivotal role, probably through consisting of spermatogonia and Sertoli cells, and of spermatogonia cell stage (Hamra et al., 2007). Our analysis of stimulated the formation of aligned rat spermatogonia up to the 32-recombinant NRG1 in combination with GDNF effectively spermatogenesis except for a few reports. For example, differentiations of RA in this process. It is also likely that NRG1 stimulates the proliferative activity of differentiating type A spermatogonia. It has been suggested that the SCF/c-kit interaction is required for the proliferation and/or survival of type A spermatogonia (Sette et al., 2000). As NRG1 stimulates the expression of SCF mRNA and vice versa in organ culture of testes (our unpublished results), it is possible that both SCF and NRG1 interact with each other and promote the proliferation of type A spermatogonia. Future experiments should examine the relationship between the SCF/c-kit and NRG/ERBB systems. The second process, the transition from type B spermatogonia to pre-leptotene primary spermatocytes, also seems to be largely dependent on the NRG1/ERBB system. RA was shown to regulate the initiation of meiosis in mouse embryonic ovaries and juvenile testes via Stra8 (Koubouva et al., 2006; Bowles et al., 2006; Baltus et al., 2006; Anderson et al., 2008). As the number of STRA8-positive cells in mutant testes was remarkably reduced compared with those in heterozygous testes (Fig. 1V-Z), we suggest that NRG1 plays a crucial role in regulating the initiation of meiosis. We further support this suggestion with our in vitro results to be discussed below.

**DISCUSSION**

NRGs are essential in spermatogonial proliferation and meiotic initiation

In the above study, we presented novel findings that FSH and RA activate Sertoli cells to produce NRG1 and NRG3, which act directly on spermatogonia, promoting their proliferation and initiation of meiosis (Fig. 5). Our findings are based on the analyses of Nrg1Ser–/– mutant mice and of re-aggregated cultures consisting of spermatogonia and Sertoli cells, and of spermatogonia only. Thus, NRG1 and NRG3 play a pivotal role, probably through ERBB4 expressed on spermatogonia, in promoting spermatogonial proliferation and their entrance into meiosis. Our results in re-aggregated cultures containing only spermatogonia (Fig. 4) indicate that ATRA and 9-cis-RA also act directly on spermatogonia.

Nrg1 is known to play essential roles in the nervous system, heart and breast (Falls, 2003), but little is known about its role in spermatogenesis except for a few reports. For example, recombinant NRG1 in combination with GDNF effectively stimulated the formation of aligned rat spermatogonia up to the 32-cell stage (Hamra et al., 2007). Our analysis of Nrg1Ser–/– mutants showed that ablation of NRG1 in Sertoli cells caused cell death of spermatogonia and/or primary spermatocytes, and also the reduction of spermatogonial proliferation and the number of STRA8-positive cells (Fig. 1). These results indicate that two processes were mainly affected in mutant testes: (1) the transition from undifferentiated type A spermatogonia to differentiating type A spermatogonia and (2) that from type B spermatogonia to pre-leptotene primary spermatocytes. The first process is known to involve the action of retinoic acid and stem cell factor (SCF)/c-kit system (de Rooij, 2001). RA was shown to induce directly the transition of undifferentiated spermatogonia to differentiating spermatogonia by stimulating Stra8 and Kit gene expression (Zhou et al., 2008a). It is possible that NRG works as a downstream factor of RA in this process. It is also likely that NRG1 stimulates the proliferative activity of differentiating type A spermatogonia. It has been suggested that the SCF/c-kit interaction is required for the proliferation and/or survival of type A spermatogonia (Sette et al., 2000). As NRG1 stimulates the expression of SCF mRNA and vice versa in organ culture of testes (our unpublished results), it is possible that both SCF and NRG1 interact with each other and promote the proliferation of type A spermatogonia. Future experiments should examine the relationship between the SCF/c-kit and NRG/ERBB systems. The second process, the transition from type B spermatogonia to pre-leptotene primary spermatocytes, also seems to be largely dependent on the NRG1/ERBB system. RA was shown to regulate the initiation of meiosis in mouse embryonic ovaries and juvenile testes via Stra8 (Koubouva et al., 2006; Bowles et al., 2006; Baltus et al., 2006; Anderson et al., 2008). As the number of STRA8-positive cells in mutant testes was remarkably reduced compared with those in heterozygous testes (Fig. 1V-Z), we suggest that NRG1 plays a crucial role in regulating the initiation of meiosis. We further support this suggestion with our in vitro results to be discussed below.

**RA and FSH promote spermatogonial proliferation and their meiotic initiation by generating NRG1 in Sertoli cells**

In 5-6 dpp testes from wild-type mice, we showed that RA (AM580, ATRA, 9-cis-RA) and FSH promoted spermatogonial proliferation and their meiotic initiation in organ culture (Fig. 2) and also in re-aggregated cultures of spermatogonia and Sertoli cells (see Fig. S3 in the supplementary material). All promoted the expression of NRG1 and NRG3 in Sertoli cells (Fig. 4S), indicating that they act on spermatogonia indirectly through Sertoli cells. FSH acts on the specific seven-transmembrane receptor of Sertoli cells (Sprengel et al., 1990). AM580, as well as FSH, act only indirectly through Sertoli cells, because in organ culture of Nrg1Ser–/– mutants, the stimulating effect of AM580 and FSH on spermatogonial proliferation and their meiotic initiation was abrogated by TAM (Fig. 3). ATRA and 9-cis-RA, however, can act on spermatogonia directly, as well as indirectly, because they promoted spermatogonial proliferation and meiotic initiation in cultures containing only spermatogonia (Fig. 4). The differential effects between AM580 and ATRA (or 9-cis-RA) may be due to different expressions of the isoforms of RAR and RXR in spermatogonia and Sertoli cells. RARα is expressed only in Sertoli cells, whereas RARγ is detected only in spermatogonia from 6 dpp testes (see Fig. S5 in the supplementary material) (Vernet et al., 2006b). Vernet et al. (Vernet et al., 2006a) showed that selective ablation of the RARα gene in mouse Sertoli cells (Rara+/+Ser–/– mutation) or in combination with RARβ and RARγ genes (RARα/β+/+/γ+/+Ser–/– mutation), delays spermatogenesis in pre-pubertal mice. That finding indicates that a cell-autonomous effect of RA-liganded RARα in immature Sertoli cells is required to promote spermatogonia differentiation during the prepubertal spermatogenic wave. Our current results, together with those of Vernet et al. (Vernet et al., 2006a), suggest that AM580, ATRA and 9-cis-RA act on Sertoli cells by binding to RARα. How RA promotes the expression of NRG1 and NRG3 in Sertoli cells through RARα remains to be elucidated.

**NRG1 activates Sertoli cells and amplifies NRG1 and NRG3**

FSH and RA stimulated the expression of Nrg1 and Nrg3 mRNA, and NRG1 upregulated the expression of Nrg1 and Nrg3 mRNA in Sertoli cells (Fig. 4S). This indicates that as long as FSH or RA is present, NRG1 and, accordingly NRG3, are amplified consecutively in Sertoli cells by NRG1 (Fig. 5). A potential mechanism for a positive-feedback loop of NRG1 is implicated in neuroblastoma in which Nrg1 gene transcription is activated by NF-κB, the transcriptional activity of which is increased by NRG1 (Frensing et al., 2008). It may be intriguing to know whether NF-κB is involved in Nrg1 gene transcription in Sertoli cells. It is also
Neuregulins are essential for spermatogenesis

RESEARCH ARTICLE 3167

interesting that although NRG1 promoted the expression of itself and NRG3, NRG3 did not (Fig. 4S). This may be due to differences of receptor affinity between NRG1 and NRG3; however, the receptors for NRG1 and NRG3 and their downstream signaling pathways in Sertoli cells remain to be identified.

NRG1 and NRG3 activate spermatogonia directly to promote their proliferation and meiotic initiation

NRG1 and NRG3 directly stimulated the proliferation of spermatogonia and their entrance into meiosis in re-aggregated cultures of purified spermatogonia (Fig. 4). These results are based on our successful cultivation of viable (~90%) spermatogonia for as long as 1 week. Although it has been reported that germ cells survived only for 48 to 72 hours under feeder cell-free and serum-free conditions (Zhou et al., 2008a), the reason for our successful culture may be due to the cell adhesion retained in the re-aggregates, and/or the three-dimensional structure of the cells maintained within the collagen matrix, and/or collagen as a substrate on which the cells attached. The reason for our success is now being investigated.

NRG1 is considered to signal through ERBB receptors in spermatogonia. ERBB receptors are indispensable not only because they have essential roles in normal physiological processes occurring during development, but also because of their involvement in various types of human tumors (Holbro et al., 2003). Ligand binding to ERBB receptors induces the formation of homo- and heterodimers leading to the activation of the intrinsic kinase domain, which in turn leads to the activation of intracellular pathways such as the mitogen-activated protein kinase (MAPK) and phosphatidylinositol-3 kinase (PI-3K) pathways (Olayioye et al., 2000; Holbro et al., 2003). Amplification of ERBB1 and ERBB2 contributes to processes linked to malignant development and ERBB2 bound with a partner in a heterodimer is responsible for the strong and prolonged activation of downstream signaling pathways leading to cell proliferation (Olayioye et al., 2000; Holbro et al., 2003). However, ERBB4 correlates in general with increased differentiation, such as that produced by NRG3 and ERBB4 signaling when regulating mammary bud specification (Muraoka-Cook et al., 2008). Therefore, distinct receptor combinations formed in response to NRG1 and NRG3 may determine the spermatogonial fates of proliferation or differentiation. The mechanism controlling the usage of the receptor combinations remains to be elucidated.

ATRA and 9-cis-RA act on spermatogonia directly to stimulate their proliferation and meiotic initiation

Our current study showed that ATRA and 9-cis-RA promoted spermatogonial proliferation and the initiation of meiosis in re-aggregated cultures containing only spermatogonia (Fig. 4), indicating that they act directly on spermatogonia. Our results are consistent with those of Zhou et al. (Zhou et al., 2008a) who showed, in cultures of THY1+ spermatogonia, that ATRA directly induced the meiotic entry of murine spermatogonia in cultures containing only spermatogonia. How, then, does ATRA stimulate spermatogonial proliferation and the initiation of meiosis in spermatogonia? As AMS80, a specific agonist of RARz, did not act on spermatogonia directly (Fig. 4), ATRA appears to act by binding RARy to form a heterodimer with RXR. In addition, our results indicate that ATRA acts on spermatogonia by promoting NRG3 expression within them (Fig. 4R). Thus, NRG3 is considered to act on spermatogonia as an autocrine growth factor. A similar example was reported by Xiao et al. (Xiao et al., 1999) using dominant-negative Rara mutants: they suggested that retinoid receptor heterodimers located in differentiated suprabasal cells mediated retinoid induction of heparin-binding EGF-like growth factor, which in turn stimulated basal cell growth via intercellular signaling. In summary, ATRA activates spermatogonia directly to promote proliferation and meiotic initiation, and also indirectly by activating NRG1 and NRG3 expression in Sertoli cells. Though it is currently unknown which pathway is dominant or equally effective, these two ways of retinoid action may provide a ‘fail-safe’ mechanism for RA to promote meiosis. Future study is required to clarify how ATRA activates NRG3 expression and whether the NRG3/ERBB signaling is essential for ATRA-mediated stimulation of spermatogonial proliferation and meiotic initiation.

Acknowledgements

We appreciate Marie A. DiBerardino for editorial review of the manuscript, Y. Nishimune for anti-TRA98 antibody, S. Chuma for anti-SYCP3 antibody, and the NHPP and A. F. Parlow for purified porcine FSH. This work was funded by Grants-in-Aid for Scientific Research (No. 22247008) from the Ministry of Education, Science, Sports and Culture of Japan (to S.-I.A.).

Competing interests statement

The authors declare no competing financial interests.

Supplementary material

Supplementary material for this article is available at http://dev.biologists.org/lookup/suppl/doi:10.1242/dev.062380/-/DC1

References


