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REVIEWS

Retinoic Acid, Local Cell–Cell Interactions, and Pattern Formation in Vertebrate Limbs

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Retinoic acid (RA), a derivative of vitamin A, has remarkable effects on developing and regenerating limbs. These effects include teratogenesis, arising from RA’s ability to inhibit growth and pattern formation. They also include pattern duplication, arising as a result of the stimulation of additional growth and pattern formation. In this review we present evidence that the diverse effects of RA are consistent with a singular, underlying explanation. We propose that in all cases exogenously applied RA causes the positional information of pattern formation–competent cells to be reset to a value that is posterior–ventral–proximal with respect to the limb. The diversity of outcomes can be seen as a product of the mode of application of exogenous RA (global versus local) coupled with the unifying concept that growth and pattern formation in both limb development and limb regeneration are controlled by local cell–cell interactions, as formulated in the polar coordinate model. We explore the possibility that the major role of endogenous RA in limb development is in the establishment of the limb field rather than as a diffusible morphogen that specifies graded positional information across the limb as previously proposed. Finally, we interpret the results of the recent finding that RA can turn tail regenerates into limbs, as evidence that intercalary interactions may also be involved in the formation of the primary body axis.

INTRODUCTION

Developing and regenerating limbs respond in remarkable and diverse ways to retinoic acid (RA). In different organisms and under different conditions, these responses lead to various types of pattern duplications as well as to teratogenesis. Over the last year or two, there has been a flurry of reviews that have focused either on limb development or limb regeneration (see for example: Brockes, 1990; Thaller and Eichele, 1990b; Duboule, 1991; Maini and Solursh, 1991; Stocum, 1991a; Tabin, 1991; Tickle, 1991). All have drawn attention to some of the known and interesting facts about the effects of RA on limbs. Our article differs from the previous reviews by providing a unified view of the mode of action of RA and in presenting evidence that the apparently varied RA responses can be comprehensively understood in terms of the local cell–cell interactions that drive limb outgrowth and pattern formation. Further, we propose that the principle function of endogenous RA in limb development is in the establishment of the limb field. We begin with the two premises concerning limb development and regeneration that underlie our interpretation.

A. PREMISES

1. Developing and Regenerating Limbs of All Vertebrates Use the Same, Basic Mechanisms for Limb Outgrowth and Pattern Formation

There are several lines of evidence that support this premise and none that argue against it. This issue is also addressed in Muneoka and Sassoon (1992). First is the fact that the basic features of limb outgrowth, such as proximal to distal elaboration of the pattern and the requirement for a permissive epidermis, are the same for the developing limbs of all the different classes of vertebrates and for regenerating amphibian limbs (see Bryant and Muneoka, 1986).

Next is the fact that all vertebrate limbs share a common repertoire of regulative responses to experimental interventions (Bryant and Muneoka, 1986) (Fig. 1). These include the development of supernumerary limbs in response to positional disparities, the ability to intercalate missing parts of the pattern along the proximal–distal axis, and the ability to regenerate amputated distal parts of the pattern. In urodele amphibians these responses can be evoked from limb cells at any time.
During the life of the animal. In all other tetrapods, the regulative responses are more restricted, both spatially and temporally.

In the developing chick limb, for example, only cells in the distal mesenchyme (progress zone) under the apical ectodermal ridge (AER) exhibit these behaviors (Summerbell et al., 1973; Summerbell, 1977; Kieny and Pautou, 1977). Once cells leave the progress zone during limb outgrowth, they no longer participate in the creation of new pattern. The regulative response of distal limb bud cells is further restricted by the limited spatial extent of the permissive AER. For example, supernumerary limbs develop in response to anterior–posterior confrontations because the site of the positional disparity is beneath the AER, which is present as a narrow distal band extending anterior to posterior (Tickle et al., 1975). In contrast, if dorsal and ventral cells are confronted by bud tip grafting, the positional disparity does not lay beneath the AER, and supernumerary limbs do not develop. Despite the absence of a full supernumerary response, supernumerary muscles are generated in response to dorsal–ventral confrontations (Javois and Iten, 1982). Similarly, the lack of regenerative ability in chick limb buds can be viewed as a consequence of a lack of a permissive epidermis. Numerous experiments have shown that if the AER is removed in chicks, the epidermis heals but an AER is not reformed. As a consequence, further pattern formation ceases (Saunders, 1948). We have recently confirmed earlier reports that if amputated chick limb buds are resupplied with an intact AER, they are capable of complete regeneration (Rubin and Saunders, 1972; Muneoka et al., 1992). Hence, cells that have recently exited the progress zone can be caused to reenter it. In mice, a distinct AER does not form until after outgrowth is well underway (see Wanek et al., 1989b), and it is possible that this lesser dependence upon a maximally developed AER could account for the ability of early stage rodent limb buds to “regenerate” in culture (Deuchar, 1976; Chan et al., 1991) and of older limb buds to regenerate their peripheral digits (Wanek et al., 1989a).

The limitations discussed above can account for all the deficits in regulative behavior displayed by higher vertebrate limbs when compared to the spectrum shown at its most complete by the limbs of urodele amphibians. In urodeles, outgrowth permitting epidermis covers the entire outgrowth, and it is readily reformed after removal (Stocum and Dearlove, 1972). Further, limb cells that leave the progress zone at the limb tip can reinitiate pattern formation in response to injury.

A third line of evidence that outgrowth of developing and regenerating limbs involves the same mechanisms comes from a direct test that asks whether developing and regenerating limb cells can interact appropriately with one another to form a limb (Fig. 2). This test has been carried out in urodeles, the only group to both develop and regenerate their limbs. Grafts to confront anterior cells of a developing limb with posterior cells of a regenerating limb and vice versa lead to the formation of supernumerary limbs of normal structure (Muneoka and Bryant, 1982). Further, these limbs are derived in equal parts from cells that originated on the developing
limb buds anti regenerating hlastemas of urodeles, developing and regenerating walls on either side of the confrontation interact to form from the result of grafting within developing or regenerating limbs. The contribution pattern is indistinguishable from supernumerary limbs. The development of uncovering the genes involved in limb pattern for-...on the regenerating limb side (Muneoka and Bryant, 1984a). A final line of evidence concerning the universality of limb outgrowth and pattern formation mechanisms is that genes considered likely to be involved in pattern formation are expressed in both developing and regenerating limbs. Although we are only now at the threshold of uncovering the genes involved in limb pattern formation, sufficient evidence is already available to support this conclusion. Candidate pattern formation genes that have been reported to date include the various nuclear and cytoplasmic receptors for RA (see Mendelsohn et al., 1992), as well as various homeobox genes (see Izpisúa-Belmonte and Duboule, 1992; Muneoka and Sassoon, 1992). Although data for only two homeobox genes from regeneration blastemas have been published (Hox 3.3: Savard et al., 1988; Hox 4.6: Brown and Brockes, 1991), it is clear that several more are expressed. In our own efforts to isolate and characterize homeobox genes expressed during regeneration, we have thus far identified the axolotl homologues of more than a dozen additional homeobox genes (Gardiner and Blumberg, unpublished data).

The differences that do exist between limb development and limb regeneration all seem to involve aspects of cell biology that are not directly related to the control of growth and pattern formation. Most of these differences are associated with events involved in dedifferen-
In summary, in all essential features concerning growth and pattern formation, limb development, and limb regeneration are alike.

2. Local Positional Differences Drive Growth and Pattern Formation: in the Absence of Positional Differences, Growth and Pattern Formation Cease

The clearest demonstrations that positional differences are essential for growth and pattern formation come from grafting experiments to generate positional disparities (French et al., 1976; Bryant et al., 1981). For example, when posterior cells are placed next to anterior cells, growth and pattern formation are stimulated and a new limb forms between the confronted cells. In higher vertebrates, the pattern of cellular contribution has become very one-sided, with anterior cells contributing the majority of the cells for the new outgrowth (Honig, 1983; Javois and Iten, 1986). Hence, when a small group of posterior chick wing bud cells is grafted into an anterior (responding) location, a full supernumerary set of digits is obtained (Tickle et al., 1975). In contrast, when small grafts of anterior (responding) cells are made into a posterior site, the supernumerary response is more limited, undoubtedly due to the small pool of cells available to respond (Iten and Murphy, 1980; Honig, 1983). Nevertheless, when the whole tip is rotated, thereby bringing anterior and posterior regions of the bud into contact on both edges of the limb bud, two full supernumerary limbs are produced as expected (Saunders et al., 1958; Javois and Iten, 1986).

In amphibians, which we assume represent the more generalized condition, it has been clearly demonstrated that the cells that form supernumerary limbs are contributed equally from anterior and posterior parts of the confrontation (Muneoka and Bryant, 1984a,b; Muneoka and Murad, 1987). Even when small clumps of cells are transplanted from posterior to anterior and vice versa, the frequency and completeness of supernumerary outgrowths are the same (Rollman-Dinsmore and Bryant, 1982; Groell and Gardiner, unpublished data). These results require that the pattern formation mechanism involves both anterior cells and posterior cells in both signaling and responding. In the simplest case, cells will respond by growth and intercalation when they are confronted with neighbors from a distant part of the pattern. If some condition has occurred to prevent one of the partners in the interaction from contributing, as appears to be the case in chick limbs and probably also in mice (see Wanek et al., 1989a), the pattern is generated by the remaining partner. Equivalent one-sided contribution can be demonstrated experimentally in amphibians, where the pattern can be generated by either anterior or posterior cells if the other partner has been prevented from participating by X-irradiation (Holder et al., 1979).

An additional feature of the growth response to positional confrontations that has been demonstrated in amphibians is that the response has an inherent directionality or polarity, with anterior cells contributing more to the ventral and posterior cells contributing more to the dorsal parts of the new pattern (Muneoka and Bryant, 1984a; Muneoka et al., 1986b). The basis for this polarity in the patterning process is presently unknown, but it could be linked to an obligatory polarity in the spatial and temporal pattern of relevant gene expression, similar to that described for the Hox-4 complex of homeobox genes in developing limbs (Dolé et al., 1989a; Izpisúa-Belmonte et al., 1991; Nohno et al., 1991; Yokouchi et al., 1991). Regardless of its basis, the existence of directional intercalation provides a possible mechanism whereby the limb field could be generated at the interface between two oppositely specified cell populations: posterior-ventral and anterior-dorsal. This idea is explored further in Section D2 (see also Fig. 12).

In addition to evidence indicating that whenever positional disparities are created, growth and pattern formation occur, there is reciprocal evidence that when positional disparities are reduced or eliminated, growth and pattern formation are concomitantly curtailed. Evidence of this type came from surgically created limb stumps in newts that were symmetrical, double half limbs (Bryant, 1976). These limb stumps were lacking half of the normal circumference of positional values, and the other half was present twice, arranged in mirror symmetry. After allowing sufficient time for these limbs to be fully healed (reintegrated, revascularized, and reinnervated), they were amputated to study their regenerative ability. As predicted, due to the lack of positional disparities, regeneration was either very reduced (and symmetrical) or it did not occur at all. Subsequent experiments have confirmed and extended these results (reviewed in Bryant et al., 1982).

Perhaps the most graphic demonstration of the relationship between positional disparities, growth, and pattern formation comes from experiments by Emile Lheureux on X-irradiated salamander limbs (Lheureux, 1975) (Fig. 3). In these experiments, limbs were X-irradiated to inhibit regeneration and then supplied with various types of skin grafts to determine the minimal number of different circumferential qualities of positional information (i.e., anterior, posterior, dorsal, ventral) that are required for outgrowth. As illustrated in Fig. 3, a full cuff of skin containing information from the entire limb circumference is able to support the regeneration of a normally patterned limb. Parenthetically, this experiment shows that dermal fibroblasts alone are capable of generating the entire limb pattern.
The telling experiment is the one in which a cuff of skin derived from only one circumferential position is wrapped around the irradiated limb (Lheureux, 1975). Despite the presence of unirradiated fibroblasts, regeneration does not occur. Only when at least two different qualities of positional information from the limb circumference are present (e.g., dorsal and ventral or anterior and posterior) does regeneration take place.

In summary, positional disparities are essential for growth and pattern formation; without them, growth and pattern formation do not occur. In normal limb outgrowth, in both development and regeneration, the circumferential positional disparities essential for pattern formation (French et al., 1976; Bryant et al., 1981) exist within the progress zone. As cells divide in response to these positional disparities, they adopt a more distal positional identity (Bryant et al., 1981), thereby accounting for the properties that characterize the progress zone (Summerbell et al., 1973).

The above two premises form the basis of our interpretation of the responses of limb cells to RA.

B. INTERPRETATION OF RESPONSES OF LIMBS TO EXOGENOUS RA

In this section we will focus on the effects of exogenously applied RA on limbs and defer until later a discussion concerning the role that endogenous retinoids might play in normal development.

1. Exogenously Applied RA Changes the Positional Values of Limb Cells to a Value That Is Posterior–Ventral–Proximal with Respect to the Limb

This conclusion is based on experiments in both developing and regenerating limbs (anterior–posterior axis: Tickle et al., 1982; Kim and Stocum, 1986; dorsal–ventral axis: Ludolph et al., 1990; proximal–distal axis: Nuzzi and Saxena, 1978; Maden, 1982). At the present time, very little is known concerning how this change is effected, and even less is known regarding the molecular nature of the positional values that are changed by RA.

As is apparent from other reviews in this issue (e.g., Mendelsohn et al., 1992), molecules likely to be involved in mediating the effect of RA in limbs include cellular retinoic acid binding proteins (CRABPs), as well as DNA-binding retinoic acid receptors (RARs; RXRs). Homeobox genes presently are the best candidates for limb pattern formation genes (Duboule, 1991), and thus would be likely targets of RA, a subject also reviewed in this issue (see Izsiké-Belmonte and Duboule, 1992; Muneoka and Sassoon, 1992). Representatives of all these classes of molecules have been shown to be present in developing and regenerating limbs, and hence could be involved in the change in positional value effected by RA.

In several instances an apparent dose response to RA has been observed, with higher doses for longer periods leading to more extreme results than lower doses for shorter periods of time (Tickle et al., 1985). At present, it is not possible to distinguish between two possibilities for the dose effect. The first possibility is that cells are converted from one extreme toward the other (e.g., from

Fig. 3. Relationship between positional disparities, growth, and patterning. In this experiment limb stumps are shown in cross section on the left, and the outcome after regeneration on the right. The outer ring represents skin; the inner circle represents the core of the limb. Hatched tissues have been X-irradiated to prevent participation in regeneration. In A, when both the skin and the core of the limb are X-irradiated, limbs do not regenerate. In B, limbs with an X-irradiated core can regenerate when provided with a normal circumference of unirradiated skin. In C, if the unirradiated skin graft is oriented such that only one quality of circumferential information is present at the amputation plane, limbs with an X-irradiated core do not regenerate. In D, when a minimum of two qualities of circumferential information are present at the amputation plane, limbs with an X-irradiated core can regenerate. Abbreviations used are as in the legend for Fig. 1. After Lheureux (1975).
anterior toward posterior) along a continuum, the higher the dose or the longer the exposure, the further the change along this continuum. For example, in the chick limb it has been suggested that a RA concentration of 0.9 nM specifies digit 2, a concentration of 2.5 nM RA specifies digits 3 and 2, and 25 nM RA specifies digits 4, 3, and 2 (Tickle et al., 1985). It has also been hypothesized that digit 2 must be specified before digit 3 and similarly, specification of digits 2 and 3 must precede that of digit 4. According to this view, it would obviously take a longer period of exposure, and a higher dose of RA, to specify a digit 4 than a digit 2. Evidence for a sequential activation of expression of homeobox genes in EC cell lines in response to different RA concentrations (Boncinelli et al., 1991) has been cited as indirect evidence for the likelihood of a graded response to RA in limbs (Stocum, 1991b; Tabin, 1991). However, the RA concentrations (10^{-5} M) and duration of exposure (several days) necessary to obtain sequential expression of Hox genes in vitro (Boncinelli et al., 1991) are not consistent with the kinetics of the RA effects on limb pattern formation (less than 5 \times 10^{-6} M for less than 20 hr; Eichele et al., 1985). In addition, a recent report indicates that the 5' members of the Hox-4 complex that are sequentially expressed during limb development (and are considered to be involved in anterior–posterior specification), are not activated but inhibited by RA in the in vitro model system (Simeone et al., 1991).

The other interpretation is that cells are switched from anterior to posterior at some threshold level of RA, and that higher doses and longer exposures generate more switched cells (Wanek et al., 1991; also see Tickle et al., 1985). Evidence that RA converts anterior cells to posterior edge cells but not to cells with intermediate positional values is presented and discussed below (Section B2). Cheryl Tickle has presented direct evidence that a graded patterning response can be stimulated by varying the numbers of posterior (ZPA) cells (Tickle et al., 1985). According to this view, it would appear as though the cells had been only partially posteriorized. As discussed below (Section B4), this view can also account for the equivalent, dose–response phenomenon in RA-treated regenerating amphibian limbs.

Recent investigations of the molecular basis of graded, inducible responses at the level of gene transcription provide evidence consistent with this second interpretation (Ko et al., 1990). These studies have been carried out in a model system for the dose-dependent transcription of a reporter gene with a glucocorticoid response element in its enhancer/promoter region. The glucocorticoid nuclear receptor, along with those for steroid and thyroid hormones and retinoids, belongs to the steroid/thyroid hormone receptor superfamily of ligand-dependent transcription factors (see Evans, 1988; Green and Chambon, 1988). In these studies it was found that individual cells do not respond in a graded way to increasing doses of dexamethasone; rather the dose dependence is a cell population phenomenon, in which increasing numbers of cells initiate transcription in response to increased concentrations of inducer.

Experiments designed to distinguish between these or other possibilities would provide an important, missing piece to the puzzle of how RA changes a cell's positional value.

2. Locally Applied RA Causes Local Changes in Positional Values and Is Equivalent to Grafting

Although the pattern-duplicating effects of RA were first described for the proximal–distal axis of regenerating amphibian limbs (Niazi and Saxena, 1978; Maden, 1982), it has been the effect of RA on the anterior–posterior pattern of chick wing buds that has received the most attention. The experimental paradigm is to load RA onto ion-exchange resin beads and then to implant a single bead under the anterior edge of the AER (Tickle et al., 1985). The bead acts as a slow release source of RA. When the limb is examined several days later, a supernumerary set of digits is found with its posterior edge adjacent to the site of the bead. Hence, the RA–bead appears to mimic a graft of posterior cells into an anterior location. We have confirmed the generality of this phenomenon by eliciting similar responses in axolotl limbs (Sessions et al., 1989, and manuscript in preparation). Hence, when RA–beads are implanted into the anterior of axolotl limbs, either developing buds or regenerates, they cause pattern duplications similar to those described for the chick.

There are at least two explanations for pattern dupli-
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A

DIGITS

B

FIG. 4. RA-induced pattern duplication in chick limb buds. (A) An RA-bead in the anterior progress zone of a chick wing bud, 15–24 hr after grafting. The bud is shown end on; numbers represent positional values in the periphery of the limb mesenchyme. Cells adjacent to the bead have been converted to posterior–ventral–proximal positional value by exposure to RA (see text). Asterisks indicate where intercalation will be stimulated by the proximity of posterior cells with anterior cells. The bar below indicates the digits that are represented by the positional information shown. (B) After intercalation, a fully duplicated limb pattern is formed.

A second explanation is that RA first converts cells adjacent to the bead into posterior edge (ZPA) cells (Wanek et al., 1991; see also Tickle et al., 1985). Since these converted posterior cells are surrounded by anterior cells, the positional disparity thereby created results in the stimulation of growth and the intercalation of new pattern in the same way as proposed for posterior to anterior tissue grafts (Fig. 4). As a test of whether cells next to an RA-bead become functional ZPA (posterior edge) cells, we removed RA-beads from chick limb buds at various times after implantation and assayed the wedge of tissue next to the bead for ZPA properties by grafting into the anterior of a host wing bud (Wanek et al., 1991). We found that the cells next to the bead become functional ZPA cells beginning at about 15 hr after bead implantation. Further, by using the chick/quail marker, we found that all of the cells in the 200- to 250-μm wedge adjacent to the bead become ZPA cells; i.e., the grafted wedge of RA-treated anterior cells makes the same small contribution to the posterior of digit 4 as does a ZPA graft. In other words, the region of the limb bud in which the rudiments of the specified digits 4, 3, and 2 should be found according to the RA-as-morphogen view, consists only of cells that behave like ZPA cells.

Based on the results from this and other (Summerbell and Harvey, 1983; Noji et al., 1991; Tickle, 1991) experiments, we interpret the sequence of events that lead to RA-bead-induced pattern duplication in chick limbs as follows. Cells next to the bead are exposed to RA which initiates changes that lead eventually to changes in positional value. We propose that the cells acquire a posterior–ventral–proximal positional identity. In chicks because pattern regulation only occurs along the anterior–posterior axis (see Section A1), the experiments only reveal the posteriorization of the cells. In order to see unambiguous proximal–distal effects, limbs that had already formed distal parts of the pattern would need to be treated. As we discuss later, the time required for the response to RA to be completed by the pattern formation-competent cells precludes the possibility of this test. Nevertheless some evidence of a proximal–distal effect has been noted by Oliver et al. (1990). In addition, as discussed above (Section A1), dorsal–ventral effects are expected to be masked if pattern regulation is only possible beneath the AER. After the start of RA exposure, no stable change in positional value is achieved before 15 hr of exposure. Removal of the bead at this time leads to some partially duplicated limbs (Eichele et al., 1985). At this time point, grafts of cells that were adjacent to the bead into an anterior site in a host wing bud show ZPA activity, although the frequency of full duplications increases the longer the cells are exposed to RA prior to testing. By 24 hr after RA exposure, cells next to the bead have developed near-normal ZPA activity. We have proposed that, left in situ, such cells will interact with adjacent anterior cells to generate the parts of the pattern that normally lie between the newly generated extreme posterior edge and digit 2 (Wanek et al., 1991) (Fig. 4).

The time of conversion of cells adjacent to the bead into posterior edge cells proceeds the onset of ectopic expression of the Hox genes that are normally expressed in a restricted posterior domain. Posterior edge cells are present after 15–24 hr of exposure to RA. Cells next to the bead initially express Hox-4.6 at about 20–30 hr, and later express Hox-4.8 at about 24–48 hr (Izpisúa-Belmonte et al., 1991, Nohno et al., 1991). The timing of the Hox gene expression relative to the timing of con-
version to posterior suggests that the initiation of Hox-4 gene expression is an indirect result of RA exposure, with RA initially converting cells to posterior identity. As new cells are intercalated, they begin to express the posterior Hox-4 genes.

One aspect of the comparison between ZPA- and RA-bead-induced duplications that has generated confusion is that, in each case, the duplication-inducing stimulus must be present for about 15 hr in order for its effect on limb pattern to become irreversible (Tickle and Brickell, 1991). This similarity in timing has contributed to the conclusion that the ZPA is in fact a source of endogenous RA, and that RA acts as a natural morphogen to specify graded positional values across the normal anterior–posterior axis of the chick limb. It is also possible to view the similarity in time to independence from the stimulus in a different way (Fig. 5). In both RA-bead- and ZPA-induced duplications, we define the time to independence as the time after which positionally altered cells have been generated next to the bead or graft. The presence of these cells allows for the generation of the remaining intermediate positional values after the removal of the stimulus.

Let us first consider what happens after a ZPA graft (Fig. 5). Following an interval for graft healing (most likely less than an hour), local interactions between graft and host cells stimulate cells to enter S-phase of the cell cycle. An increase in S-phase cells next to a ZPA graft has been documented as early as 4–5 hr after grafting (Cooke and Summerbell, 1980), which would correspond to 3–4 hr after healing-in of the graft. Twelve hours later, i.e., 16 hr after grafting, the increase in S-phase cells results in an increase in mitosis (Cooke and Summerbell, 1980) and the start of the widening of the limb bud (Smith and Wolpert, 1981). We view this as the time when the intercalated pattern begins to be generated. The limb becomes independent of the graft after 15–17 hr (Smith, 1980), presumably because newly intercalated cells with positional values intermediate between the graft and the host have been generated in the host tissue. The remaining parts of the duplicated pattern can be generated by intercalation after the limb bud has become independent from the graft stimulus.

In the case of RA-beads, the time to independence also suggests an involvement of the cell cycle. We propose that cells only become reprogrammed by RA as a result of going through some critical part of the cell cycle in the presence of RA. Once enough of the cells are reprogrammed, beginning at about 15 hr, they are independent of the stimulus (Eichele et al., 1985; Wanek et al., 1991), and can therefore go on to interact with adjacent host cells to generate the new pattern through intercalary growth. As the population of cells in the vicinity of the bead is exposed to RA for longer periods, increasing numbers of cells pass through the critical phase of the cell cycle in the presence of RA, and thus the population of reprogrammed cells can continue to expand with time. As discussed above (Section B1) we predict that the increase in the number of reprogrammed cells will eventually result in a more complete supernumerary response. This view predicts that the growth necessary for the expanded pattern in RA-induced duplications will take place after and as a consequence of the initial re specification interval (Fig. 5). In other words, the similarities in time to independence reflect different processes, both of which involve the cell cycle and both of which result in the generation of some threshold number of respecified cells that are intermediate between anterior and posterior extremes. Hence, the increase in S-phase observed after ZPA grafts at 4–5 hr is expected to occur at about 18–20 hr after bead im-

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**Fig. 5.** Proposed sequence of events leading to pattern duplication after a ZPA graft and an RA-bead. Additional detail is provided in the text.
plantation (a 15-hr respecification interval plus 3–4 hr to recruit cells into S-phase). By the same reasoning, an increased mitotic rate and the onset of bud widening is expected to occur another 10–12 hr after that, or at about 28–30 hr after initial bead implantation (Fig. 5).

A puzzling difference between ZPA graft- and RA-bead-induced duplications becomes interpretable in the context of this view. Although limbs up to stage 24 can respond to ZPA grafts by generating extra digits (Iten et al., 1983; Summerbell, 1974), their response to RA–beads is much reduced at stages 21–22, and absent at stage 23 (Eichele et al., 1985; Tickle and Crawley, 1988). We suggest that this difference is related to the prediction that RA–bead-induced duplications require a longer time, on the order of 12–24 hr, for completion of the intercalary growth necessary for the generation of the new pattern. After stage 20/21, there is insufficient time remaining in the limb pattern formation period for cells next to a bead to first become reprogrammed by RA and then to complete intercalation before cells lose their responsiveness to positional signals. We thus far have been unsuccessful in attempts to generate pattern duplications in developing mouse limbs by implanting RA–beads, despite being able to obtain a limited regulative response from grafts to confront anterior and posterior cells (Wanek et al., 1989a). It has not been possible to extend these studies to earlier time points due to the technical limitations of ex utero surgery (Muneoka et al., 1986c). It is likely that the amount of time available in the mouse studies for pattern duplication by locally applied RA (in contrast to the time needed for duplication as a result of direct anterior–posterior confrontations created by grafting) is insufficient for the reasons outlined above.

In summary, we propose that locally applied RA in developing chick limbs (as well as in the limbs of other vertebrates) causes pattern duplication via a two-step process. In the first step, cells adjacent to the RA–bead are reprogrammed to posterior–ventral–proximal with respect to the limb. The time required for this step suggests an involvement of the cell cycle. Once reprogrammed cells have been generated, they interact with adjacent anterior cells stimulating growth and the intercalation of new pattern by the same means as a graft of posterior cells.

3. Globally Applied RA Leads to Uniformity of Positional Values and Hence to Hypomorphic Outgrowth in Limb Development

Before RA was found to cause interesting pattern duplications in amphibians and chicks, retinoids were known to be teratogens in mammals (see Lammer et al., 1985). Indeed, at the present time, a retinoid used in acne medication is considered one of the most potent of human teratogens. When retinoids are supplied to the fetus via the placenta, a number of different organ systems, among them limbs, develop abnormally (Kochhar, 1977; Kochhar et al., 1984). The types of limb defects encountered are varied, and are sensitive to dose, to the particular form of retinoid administered and to the time of exposure relative to stage of limb development (Howard and Willhite, 1986; Kochhar, 1977). However, in almost all instances, the affected limbs are hypomorphic, ranging from complete truncations to missing or reduced skeletal elements at different proximal-distal levels. The effects of systemic administration in mammals can be replicated in chicks by adding retinoids to the amniotic cavity (Larsen and Jannasr, 1987) or by implanting very high dose RA–beads into limb buds (Tickle and Crawley, 1988). Similar results can also be generated in developing (but not regenerating, see Section 4 below) axolotl and Xenopus limbs by exposure to retinoids in the aquarium water (Scadding and Maden, 1986a,b).

Various explanations for the teratogenic effect of retinoids have been proposed, among them cell death in the mesenchyme and destruction of the AER. However, recent studies have shown that teratogenic doses of RA do not increase the amount of cell death in mouse limb buds above normal (Abbott et al., 1990). In chicks treated to give truncated limbs, the AER is affected and becomes flat. However, this appears to be a secondary consequence of RA-induced changes in the mesenchyme, since transplants of AERs that have been flattened by RA exposure, to untreated mesenchymes, leads to the restoration of the AER and the outgrowth of a normal limb (Tickle et al., 1989).

The teratogenic effects of RA can be interpreted as a consequence of the ability of RA to change limb cell positional information to posterior–ventral–proximal. As described in the preceding section, local reprogramming generates positional disparities that in turn lead to extra pattern. The converse is true for global application of RA. If a sufficient dose of RA is administered, then all or most cells in the progress zone will be reprogrammed. Rather than generating positional diversity, globally applied RA will reduce positional diversity. Since positional differences are required to sustain growth and pattern formation (see Section 2A), reduction of positional diversity will lead to reduced or truncated structures (Fig. 6). Studies in mouse limb buds have documented an inhibition of growth in response to RA exposure in vivo (Abbott et al., 1990), and we have observed a growth reduction in cultures of mouse limb bud cells treated with RA (Gardiner et al., 1992). The view we have presented here predicts that teratogenic effects in limbs would be expected at similar stages to
those at which pattern duplication can be elicited, as well as at later stages, since no enhanced growth response to RA-induced changes in positional information is required in order to observe the effect. Studies in chick limbs show that, whereas RA-induced duplications do not occur beyond stage 21/22, RA-induced truncations and reductions occur from prelimb bud stages (Larsen and Janners, 1987) at least through stage 24, the latest stage tested (Tickle and Crawley, 1988). Another prediction of this view is that it should be possible to rescue limbs whose cells have been reprogrammed to positional uniformity by removing any further influence of RA and providing anterior cells as a source of positional diversity.

An outstanding exception to the discussion above concerning the effects of global application of RA can be found in regenerating amphibian limbs. This exception is discussed below, where we present arguments that this is the exception that proves the rule.

In summary, we propose that global application of RA in developing vertebrate limb buds causes all cells to be reprogrammed toward uniform positional values. Lack of positional diversity in the progress zone leads to the failure of growth and pattern formation, and to the formation of reduced or truncated limbs.

4. Globally Applied RA in Regenerating Amphibian Limbs Leads to Uniformity of Positional Values in the Blastema; Outgrowth Is Rescued by Dedifferentiating Cells of the Stump

All the arguments presented above for the teratogenic effects of systemically applied retinoids in developing limbs might also be expected to apply to the regenerating limbs of amphibians. Indeed, a significant growth inhibition in the blastema during the period of exposure to retinoids has been well documented (Maden, 1983b; Pietsch, 1987). However, when urodele limbs are amputated through the lower arm and the animals are swum in a retinoid solution, or when they are injected with a retinoid in the first few days after amputation, the striking result is the development of limbs with serially duplicated elements along the proximal-distal axis (Maden, 1982; Thoms and Stocum, 1984). The result is similar to what would be expected if a proximal blastema were developing on a distal stump.

In contrast to the conclusions of others (Scadding and Maden, 1986a,b; Stocum, 1991a), we view the difference in the eventual outcome of exposure to RA in developing and regenerating limbs as attributable to the special circumstances of regeneration, not to any difference in the fundamental mechanisms of limb outgrowth and pattern formation in the two systems. The special circumstances we are referring to are associated with the process of dedifferentiation of mature limb tissues, leading to the creation of a group of undifferentiated cells from which outgrowth and pattern formation can proceed. Of particular significance is that dedifferentiation continues to progress back from the amputation plane during the early phases of blastema outgrowth, causing the addition of new pattern formation-competent cells to the base of the growing blastema (Tank, 1977).

The observation that the period of maximal sensitivity to retinoids coincides with the period during which dedifferentiation is occurring (Maden, 1983b; Maden et al., 1985; Thoms and Stocum, 1984) indicates to us that retinoids are only affecting the undifferentiated cells of regenerating limbs. This parallels the finding that it is cells in the progress zone of developing limbs that are sensitive to RA (see Tickle and Brickell, 1991). In regenerating limbs, administration of RA either too soon, before cells have dedifferentiated, or too late, after cells have redifferentiated, results in little or no pattern duplication (Maden, 1983b; Thoms and Stocum, 1984; Maden et al., 1985; Niazi et al., 1985). We propose that the effect of retinoid treatment in regenerating limbs is the same as that in developing limbs; namely, blastema cells are converted to a posterior-ventral-proximal positional value (Fig. 7). We propose that in the presence of retinoids, growth and pattern formation are arrested due to the uniformity of positional values present in the blastema. During the exposure interval, any newly dedifferentiating blastema cells that enter the blastema from the stump will be reprogrammed along with the original blastema cells. However, after retinoids are withdrawn, newly dedifferentiating cells entering the base of the blastema will be able to maintain their original positional values. These newcomers are therefore able to provide the necessary circumferential positional diversity (anterior and dorsal) to stimulate outgrowth.

Globally Applied RA to Developing Vertebrate Limbs: RA affects all cells in the pattern-forming region of the limb bud similarly by converting them to cells with posterior-ventral-proximal positional information. Without positional disparities, limb outgrowth ceases, leading to limb defects. Details of the diagrams are the same as described in the legend for Fig. 4.
Regenerating limb

Fig. 7. Rescue of outgrowth by dedifferentiating cells in regenerating limbs exposed systemically to RA. The details of these diagrams are similar to those described in the legends for Figs. 4 and 6, except that in this figure cells in the blastema are represented by the inner circle and stump cells that dedifferentiate and enter the base of the blastema are represented by the outer ring. The bar below represents digits as depicted in Figs. 4 and 6. RA is shown converting blastema cells uniformly to posterior-ventral-proximal positional values. After withdrawal of RA, newly dedifferentiating cells enter the base of the blastema and create the positional disparities (**) necessary for outgrowth (see text).

from the RA-treated posterior-ventral-proximalized cells of the blastema (Fig. 7). The positional confrontations necessary to promote outgrowth of an RA-treated blastema will be expected to occur on the anterior edge of the blastema, and Kim and Stocum (1986a) have reported that RA-treated blastemas appear to originate from the anterior part of the limb stump. The view that we present makes it unnecessary to propose the existence of a unique population of cells in the peripheral anterior-dorsal part of the blastema, whose properties render them refractory to the effects of RA (Stocum, 1991a,b).

At later stages of regeneration, there is still a distal region of undifferentiated cells; dedifferentiation in the stump has ceased; and the proximal regions of the blastema have left the progress zone and are beginning (or will shortly begin) to redifferentiate. Administration of RA at this stage leads to the formation of limbs that are truncated distally but normal proximally (Niazi et al., 1985). We interpret this result as follows: the proximal part of the blastema is unaffected by RA because it has left the progress zone at the time of exposure; the distal cells are reprogrammed as expected, but in the absence of newly dedifferentiated adjacent cells, they are not able to proceed with outgrowth and thus the distal part becomes hypomorphic or truncated.

According to the hypothesis presented above, the base of an RA-treated regeneration blastema is expected to consist of a mixture of proximal and distal cells. This leads to the prediction that intercalation will generate a small, reversed-polarity segment between the distal limb stump and the proximal boundary of the reprogrammed blastema cells (Fig. 8). The occurrence of such reversed-polarity segments has been noted by Kim and Stocum (1986a) and is evident in illustrations in other papers (e.g., Maden, 1983). We have found that the vast majority (83%, n = 29) of RA-duplicated regenerates have reversed-polarity segments (Fig. 9).

Demonstration that RA posteriorizes and ventralizes regenerating limb cells comes from studies of double half limbs in urodeles. We illustrate these results by reference to double anterior and double posterior limbs (Fig. 10), but similar conclusions can be drawn from the results of double dorsal and double ventral limbs. It has
been argued that when double anterior or double posterior limbs are amputated through the lower arm in the absence of RA, the reduced positional diversity at the amputation plane leads to regeneration of tapering symmetrical limbs with zero to three digits (Stocum, 1978; Krasner and Bryant, 1980; Kim and Stocum, 1986b). After RA treatment, double posterior limbs no longer regenerate at all, and double anterior limbs regenerate much more than before, forming double, mirror-imaged outgrowths (Stocum and Thorns, 1984; Kim and Stocum, 1986b). According to the view presented here, double posterior limbs cannot be rescued by newly arriving stump cells because these are posterior, like the reprogrammed blastema cells. Double anterior limbs regenerate two outgrowths because newly dedifferentiated anterior cells rescue the posterior-ventral-proximalized blastema on either side (see Fig. 10). Analogous experiments with double dorsal and double ventral limbs give equivalent results, whereby double ventral limbs regenerate less after RA treatment and double dorsal limbs regenerate more, leading to the conclusion that RA not only posteriorizes and proximalizes limb cells, but it also ventralizes them (Ludolph et al., 1990).

In another series of experiments, Stocum and colleagues have investigated the effects of RA on half limbs (Kim and Stocum, 1986b; Ludolph et al., 1990). These studies have provided results that can also be accounted for using the principles described here. In the case of half limb stumps grafted to the orbit to isolate them from all influences from the other half of the limb stump, anterior but not posterior halves can regenerate after retinoid treatment. Hence, outgrowth from the posterior-ventral-proximal blastema cells on an anterior half limb stump can be rescued as predicted by newly dedifferentiating anterior stump cells. Half posterior limbs cannot be rescued by newly arriving cells because these do not add any positional diversity to the blastema. Similar results have been obtained from RA-treated half ventral (no regeneration) and half dorsal (regeneration) limbs in situ, under conditions in which any cellular contribution from the other limb half was blocked by a head skin graft (Ludolph et al., 1990).

The effects of retinoids have also been studied in the regenerating limbs of anurans prior to the ontogenetic loss of regenerative ability. Here too, duplication of proximal limb elements is seen following systemic application and amputation at a distal level (Niazi and Saxena, 1978; Maden, 1983a; Scadding and Maden, 1986b) and a similar interpretation applies. At a lower
frequency, retinoids also induce duplication in the transverse axes in frogs, leading to the formation of mirror-imaged hands (Niazi and Saxena, 1978; Maden, 1983a; Scadding and Maden, 1986b). Similar results are also obtained in urodeles (Lheureux et al., 1986) but only when limb buds rather than mature limbs are amputated. From the available descriptions, it appears that such duplicated limbs are arranged in mirror symmetry, joined by their anterior or anterior-ventral edges (Maden, 1983a; Lheureux et al., 1986).

At present, it is not clear why the formation of mirror-imaged limbs is a response of amputated limb buds, but not of mature limbs, to RA treatment. However, in a recent study to map the organization of positional information in the interior of the axolotl limb, we were surprised to find that cells with anterior and ventral positional values predominate in the center of the limb (Gardiner and Bryant, 1989). Assuming that in limb buds, cells from central as well as peripheral regions of an amputated bud contribute equally to the regenerate, we suggest that anterior-ventral cells will be released into the center of an RA-treated blastema that consists of posterior-ventral-proximal cells. This would lead to the regeneration of mirror-imaged limbs. In contrast, it is known that the cellular contribution to the blastemas of mature limbs is dominated by peripheral cells (Muneoka et al., 1986a) and that the organization of positional information in the periphery dictates the pattern of the regenerate regardless of the organization of the central tissues (Tank, 1979; Slack, 1980). Hence, cells in the center of mature limbs and of limb buds may differ in the degree to which they are able to influence pattern formation, leading to the absence of mirror-imaged regenerates from RA-treated mature limbs.

As in the case of duplication in the anterior-posterior axis in chicks following local application of RA, the effect on the proximal-distal axis in regeneration is dose dependent, with higher doses and longer durations leading to more proximal duplications than lower doses for shorter times (Maden, 1983b; Kim and Stocum, 1986a). As with the chick results, it is not possible at present to decide whether with a lower dose fewer cells are converted to extreme proximal values or whether more cells are converted part of the way toward extreme proximal. As recognized in the “rule of distal transformation,” distal parts of the pattern are generated from more proximal parts, but not the reverse (Pescitelli and Stocum, 1980). During intercalation between cells with different proximal-distal positional values, we assume that progeny of the proximal cells take up a positional value that is intermediate between those of the neighboring cells. When many proximal cells are present, those that are at a distance from the site of interaction will maintain their most proximal positional value, and hence will stabilize the most proximal boundary of the regenerate. When few proximal cells are present, as they divide and take up intermediate positional values, the most extreme proximal part of the pattern will be lost. Hence, it will appear as though the cells had been partially proximalized. As discussed above (Section B1), recent evidence regarding the molecular mode of action of glucocorticoid hormones (Ko et al., 1990) is consistent with the interpretation that the dose dependence is a consequence of increasing numbers of cells being converted at higher doses. As in the case of the effect on chick limbs, it is important to distinguish between these possibilities if we are to understand the way in which retinoids affect the expression of positional information.

In summary, we have shown that the wide range of results that have been obtained in amphibian limbs using retinoids can be accounted for by virtue of the fact that in order to regenerate, limb tissues have to dedifferentiate. Further, dedifferentiation continues after the initial establishment of the blastema, providing a new source of cells migrating into the base of the blastema. There is no need to invoke different mechanisms for developing and regenerating limb outgrowth and pattern formation in order to accommodate the results. The recent results concerning the apparent homeotic effect of RA on amphibian tail regeneration (Mohanty-Hejmadi et al., 1992) are also interpretable by similar arguments, and are discussed in detail in Section D3 (see also Figs. 13 and 14).

C. RA AS MORPHOGEN

The view we present above is clearly at odds with the RA-as-morphogen idea that has dominated thinking in the limb field for much of the last decade. In this section we look at the origins of this idea and where it stands in the face of current knowledge about vertebrate limbs.

1. The Basic Idea

The idea that pattern formation across the anterior-posterior axis of developing limb buds is controlled by a diffusible morphogen (Tickle et al., 1975) grew out of Lewis Wolpert’s conceptualization of how positional information might, in theory, be specified (Wolpert, 1969, 1971). The one-sided nature of the interaction between anterior and posterior cells in the chick wing bud following grafting (most of the new growth is from the anterior partner, Houig, 1988; Javoy and Iteu, 1986) was consistent with the view that posterior cells signal and anterior cells respond. Reflecting the view that posterior cells can affect limb polarity, the name ZPA (“zone of polarizing activity”) was coined for the posterior, distal region of the bud with the ability to stimulate the
development of supernumerary outgrowths after transplantation to an anterior site (Balecuns et al., 1970).

As we have argued elsewhere (Bryant and Muneoka, 1986), while this view may work in principle for the anterior-posterior axis of the chick limb, it does not accommodate the data for other vertebrate limbs for several reasons, the most obvious of which is that in other developing and regenerating limbs, anterior as well as posterior cells both signal and respond at a graft interface (Muneoka and Bryant, 1984a,b; Muneoka and Murad, 1987). Nevertheless, the idea of a simple gradient of a diffusible molecule that specifies positional information has gained widespread acceptance.

When it was reported that RA-beads in chick limbs apparently mimic ZPA grafts (Tickle et al., 1982), RA began to be viewed as the putative endogenous morphogen (Eichele et al., 1985). A pivotal paper in the development of the RA-as-morphogen story was that of Thaller and Eichele (1987; see Slack, 1987) who reported on the levels of endogenous RA in chick limb buds. In that paper, they reported that while the level of retinol, the precursor to RA, is uniform and high in limb buds, RA itself is asymmetrically distributed with a 2.5-fold enrichment in the posterior one-fourth as compared to the anterior three-fourths. They subsequently reported evidence that posterior limb cells are capable of synthesizing RA from retinol (Thaller and Eichele, 1988). The discovery of several nuclear receptors for RA belonging to the steroid hormone receptor family (see Mendelsohn et al., 1992), as well as homologous receptors with unknown ligands (Mangelsdorf et al., 1990), provided a possible mechanism by which levels of RA could be transduced into the sort of differential gene expression that might account for pattern formation. However, considerations of physical chemistry led to the realization that these receptors would be saturated, even in anterior cells, given the levels of RA present and the shallow nature of the calculated gradient (see Smith et al., 1989). These concerns were overcome by the report that cellular retinoic acid binding protein (CRABP) was also distributed as a gradient across the anterior-posterior axis, but with more in the anterior than the posterior (Maden et al., 1988). It was thus concluded that a high level of CRABP in the anterior would bind up RA and thereby steepen the gradient of available RA (Maden et al., 1988).

In summary, the field of chick limb development was predisposed to conclude that RA functions as an endogenous morphogen, and the major evidence in support of the idea came from direct measurements of RA levels.

2. The Picture Is Not So Simple

The idea of RA-as-morphogen, and in fact the idea of any diffusible morphogen specifying positional information, has never been able to accommodate the data for amphibian limbs (discussed in Bryant and Muneoka, 1986). Rather, pattern formation in amphibian limb regeneration is thought of as occurring via local cell-cell interactions and intercalation (French et al., 1976; Bryant et al., 1981). RA itself has the additional problem that it affects positional information in all three limb axes; thus, it is difficult to conceptualize how a gradient of RA could specify graded positional values in three dimensions simultaneously. Since pattern regulation in the chick limb is basically one dimensional (for the reasons discussed in Section A1 above), a one-dimensional gradient of RA has not presented problems, provided that the discussion of mechanism is restricted to the chick limb.

In the last year or so, several findings have contributed to a reevaluation of the view that RA acts as an endogenous morphogen in chick limbs. The most direct evidence that is inconsistent with this view comes from experiments in which it was shown that the wedge of tissue next to an RA-bead contains only the most posterior edge of the pattern after 24 hr of RA exposure (Wanek et al., 1991). The extent of positional information present in that wedge of tissue corresponds to that contained in the ZPA, rather than to the anlage for digits 4, 3, and 2 as predicted from the RA-as-morphogen view (Eichele and Thaller, 1987). Other reports (although lacking cell lineage markers) have also provided evidence that RA converts cells next to the bead into ZPA cells (Summerbell and Harvey, 1983; Noji et al., 1991; Tickle, 1991). In this way, RA is not a mimic of the ZPA since the tissue next to a ZPA graft does not acquire ZPA activity (Smith, 1979).

It is not feasible to propose that once the ZPA is made, it then becomes a source of RA that reprograms cells. Since anterior cells failed to become respecified to form graded positional information in response to the exogenous RA, there is no reason to propose that they would do so in response to endogenous RA after a new ZPA has been generated. In addition, in order for the induced ZPA to become a RA source, anterior limb cells would be responding to RA autocatalytically. Such a response would preclude the development of a RA gradient that could specify graded positional information (Wanek et al., 1991).

Other evidence that is inconsistent with the idea of RA as an endogenous, diffusible morphogen comes from studies of the expression pattern of retinoic acid receptor-β (RARβ). The promoter of this gene contains an RA-responsive element within it, and therefore its expression would be expected to be elevated in places where there are elevated levels of RA. The gene has in fact been shown to be responsive to elevated levels of exogenous RA in chick limbs; however, there is no evi-
evidence of a graded distribution of RARβ across the anterior-posterior axis as predicted by the RA-as-morphogen idea. Furthermore, ZPA tissue grafted to the anterior margin does not lead to any increase in expression of RARβ in the adjacent cells (Noji et al., 1991). Two recent papers have examined the expression patterns of constructs in transgenic mice consisting of the RA-response element of the RARβ promoter fused to lacZ (Mendelsohn et al., 1991; Rossant et al., 1991). Expression of lacZ was essentially absent from the limb buds, although prominent in the adjacent trunk regions (Fig. 11). These results raise critical questions about the presence and relevance of endogenous RA for limb pattern formation.

Since the original description of a shallow gradient of RA in limb buds, an additional, endogenous retinoid (3,4-didehydroretinoic acid; ddRA) that is six times more abundant than RA and just as potent in inducing extra structures, has been identified in limb buds (Thaller and Eichele, 1990a). Since no data have been reported about the distribution of this retinoid, the current picture of the gradient of active retinoids is unclear.

The status of the reported gradient of CRABP has also become more complicated. Subsequent to the original report in the chick (Maden et al., 1988, 1989), it was reported that CRABP protein is present in mouse limb buds, but that it is not differentially distributed along the anterior-posterior limb axis (Dencker et al., 1990). In addition, studies in the mouse show that CRABP transcripts have a similar proximal-distal distribution to that of chick limb buds but they are not present in an anterior-posterior gradient (Dollé et al., 1989b). Another study on the mouse limb bud however does report an anterior-posterior gradient of CRABP transcripts (Perez-Castro et al., 1989). Most recently this issue has been further complicated by questions about cross-reactivity of the heterologous antibody used in the original study of the chick limb bud (Maden et al., 1990). Because the current view of RA as an endogenous morphogen in the chick limb is dependent on the existence of an anterior-posterior gradient of CRABP, this issue needs to be unequivocally resolved. Since there is an undisputedly nonuniform distribution of CRABP along the proximal-distal axis, either whole mounts or longitudinal sections showing proximal and distal as well as anterior and posterior on the same section would dismiss the nagging possibility that in even a slightly oblique cross section, a proximal-distal gradient could give the misleading appearance of an anterior to posterior gradient.

Finally, the original report that posterior limb cells can convert retinol into RA was incomplete in that it did not report on whether anterior cells are the same or different in this regard (Thaller and Eichele, 1988). The RA-as-morphogen view requires that ZPA cells synthesize high levels of RA from retinol, and that RA diffuses across the field of non-ZPA cells which act as dispersed sinks (Eichele and Thaller, 1987). Accordingly, the non-ZPA cells do not synthesize high levels of RA, but be-

Fig. 11. Expression of RARE-lacZ transgene in mouse embryos. Reproduced, by permission from the publisher, from Rossant et al. (1991). (a) Mouse embryo with ~9-10 somites showing transgene expression in the trunk and its apparent absence in the head and tail regions. An interpretation of this staining pattern is that RA is present in the trunk but not in the head or the tail. (b) Later embryo with stage 3 limb bud (see Wanek et al., 1989b, for limb stages). Note the maintenance of a sharp boundary of transgene expression between the head and the trunk and the absence of transgene expression in the limb buds.
come specified as to anterior–posterior positional value by exposure to different concentrations of RA generated posteriorly. However, recent studies indicate that anterior and posterior tissue extracts synthesize RA at the same rate (see Tabin, 1991). Hence, a gradient of RA would be a reflection of intrinsic differences between anterior and posterior cells, rather than being the cause of such differences as originally proposed in the RA-as-morphogen view.

In summary, in addition to the fact that the RA-as-morphogen idea cannot accommodate the data from amphibians, where RA affects all three axes similarly, the idea that it acts as a diffusible morphogen that specifies anterior-to-posterior positional information in developing chick limb buds has not withstood the test of time.

3. Where Does That Leave the Diffusible Morphogen?

As we have argued above (see also Saunders, 1977; Saunders and Gasseling, 1983), the chick limb field was primed to find a diffusible morphogen by the concepts developed by Wolpert in the late 1960s (Wolpert, 1969). However, since that time it has become apparent that we cannot look only at developing chick limbs if we want to deduce the likely mechanisms of limb pattern formation. Other vertebrates show a much more complete spectrum of regulative abilities (see Section A1) and hence provide more information about the nature of the mechanisms involved. The cumulative evidence from all vertebrate limbs cannot be accommodated by the idea that positional information in the anterior–posterior axis is specified with reference to a diffusible morphogen (Bryant and Muneoka, 1986).

Rather than seeing the emergence of graded positional qualities in limbs or other developing systems as the result of underlying gradients of diffusible molecules, we suggest that graded properties arise in development as a consequence of interactions at the interfaces between cells with different qualities. In other words, we propose that developmental processes in limbs and possibly elsewhere are driven by discontinuities. Discontinuities are resolved by the generation of a graded series of intermediate properties that provide a smooth, seamless transition between the original discontinuities. According to this view, development consists of a successive series of events in which discontinuities are first generated then resolved. For limbs, the polar coordinate model (French et al., 1976; Bryant et al., 1981), in which the pattern of the limb is seen as being generated in the progress zone as a result of intercalation between discontinuities, continues to provide a good fit for the available data (see also Iten, 1982; Saunders and Gasseling, 1983; Javois, 1984; Wanek et al., 1981). It is possible, as we discuss below (Section D3), that the same principles, namely the emergence of graded pattern as result of discontinuities, will prove to have more widespread applicability to development as a whole.

D. RA AND THE ESTABLISHMENT OF THE LIMB FIELD

1. RA and the Primary Body Axis

It has recently become clear that RA, in addition to its effects on limb development, has profound effects on pattern formation in the primary embryonic axis (Durston et al., 1989; Mitran and Shimoni, 1989; Sive et al., 1990; Ruiz i Altaba and Jessell, 1991). Hence, Xenopus embryos treated with RA are missing the most rostral-dorsal parts of the pattern and, as reported by Cho et al. (1992), in severe examples (known as “squidy” embryos) the tail is also shortened. In addition, it has been shown that genes that are normally expressed in rostral structures are repressed by RA treatment, whereas those that are normally expressed in the trunk are expressed more strongly in the response to RA (Cho and De Robertis, 1990; Sive et al., 1990; Cho et al., 1991). Finally, implantation of RA-beads into gastrula stage chick embryos leads to the development of duplicate axes (Chen and Solursh, 1990).

It is conceivable that RA not only affects primary pattern formation when it is exogenously applied, but also that it plays a role in normal axis formation (see Sive et al., 1990; Durston and Otte, 1991). A likely source of RA would be Hensen’s node in chicks (and by homology, the blastopore of amphibian gastrulae) as suggested by the results of experiments involving grafts of Hensen’s node into limb buds (Hornbruch and Wolpert, 1986, Stocker and Carlson, 1990). When grafted to the anterior of the chick wing bud, Hensen’s node causes duplicated digits, consistent either with it having a posterior specification similar to that of the posterior limb field or with it being a source of RA. The node shows this activity from the definitive primitive streak stage to the time that it has completed its regression (Hornbruch and Wolpert, 1986).

The transgenic mouse studies discussed earlier, which fail to provide support for a role for RA in limb development (Mendelsohn et al., 1991; Rossant et al., 1991), at the same time provide intriguing data concerning a possible role for RA in the primary axis. In both studies, the lacZ constructs report the presence of RA in the trunk, but not in the head or the tail of early postgastrulation embryos. The rostral and caudal boundaries of expression are impressively sharp (Fig. 11). These results lend support to a role for RA in axis development and suggest that the timing of RA synthesis in gastrulation divides the body into three domains: head, trunk, and tail. Consistent with this idea is the finding discussed above that RA treatment of Xenopus embryos tends to reduce the
gastrotricia is halted prematurely (Gerhart et al., 1989) lead to rostral truncations. These results have been interpreted as showing that in order for rostral structures to form, they need to escape from a caudalizing influence in the region of the marginal zone that invaginates last (Gerhart et al., 1989). It is conceivable that this caudalizing influence is RA.

In summary, a variety of different types of evidence is accumulating that suggests that RA is involved in the specification of pattern in the primary body axis. Of interest is the finding that while evidence suggests that the trunk region contains endogenous RA, both heads and tails appear to lack it. Further, heads and tails, but not trunks, are reduced in the presence of added RA.

2. RA and the Establishment of the Limb Base

An interesting body of data that provides an important piece of the RA puzzle comes from the studies of Hornbruch and Wolpert (1991) in which they mapped the distribution of posterior activity on the flank of the prelimb bud chick embryo at different stages. In these studies, posterior activity was assessed by the ability of the graft to stimulate the formation of supernumerary pattern after grafting to the anterior of a host limb bud. The earliest stage at which flank tissue shows posterior activity overlaps with the last stage at which Hensen’s node shows activity (stage 9). At stages 10 and 11, posterior activity is found in a positionally contiguous stripe that spans the region from 2-4 somites in front of the position of the future wing, back to the node, which is 2-4 somites posterior to the future position of the wing at this stage. At later stages, the rostral extent of the posterior activity appears to move progressively more caudal. At stage 12 the rostral limit is at somite 15; at stages 13 and 14 it is at somite 17, at stage 15 it is at somite 18; and at stage 16, the beginning of visible outgrowth (Hamburger and Hamilton, 1951), the rostral limit of activity is at the level of somite 19, where it remains. Hence, during these stages the rostral limit of posterior activity shifts from opposite somite 15 to opposite somite 19. This shift matches in both magnitude and direction the backward shift of the prospective wing region mapped by Chaube (1959). The caudal extent of posterior activity has not been as completely mapped, but at stage 16, when the wing is starting to be visible, there is a 3-somite-wide region behind the wing bud that shows no posterior activity. The position of this region corresponds to the middle of the hind limb field. Hence, at this stage, posterior activity extends from the posterior of the wing bud, through the intervening flank region and into the anterior third of the leg bud region. These puzzling findings can be reconciled by the following hypothesis.

We suggest that RA is involved in the specification of positional information in the flank mesenchyme during gastrulation, with Hensen’s node the most likely source of endogenous RA. As prospective flank cells invaginate through the primitive streak they are exposed to high levels of RA emanating from the nearby Hensen’s node. We hypothesize that flank cells are set by this exposure to RA to a positional value that can be described as posterior-ventral-proximal with respect to the (future) limb, just as occurs experimentally when older limb bud cells are exposed to RA. We have argued that positional diversity is essential for limb outgrowth; hence, we propose that limb outgrowth is initiated by the arrival of migrating cells into the region of the future limb buds. We propose that these cells arrive from more anterior-dorsal regions and bear corresponding anterior-dorsal positional values. The interaction of the new arrivals with the posterior-ventral-proximal cells of the flank is predicted to lead to intercalation of the limb base and subsequent limb outgrowth as described previously (French et al., 1978; Bryant et al., 1981, 1987). The directional intercalation discussed earlier (Section A2) would make it possible for interactions at the interface between two oppositely specified groups of cells to generate a complete limb base that was oriented appropriately with respect to the main axes of the body (Fig. 12). The idea that fields can arise at the interfaces between differently determined regions has also been explored by Meinhardt (1991) to account for the origin of the limb field. The idea that cells with anterior-dorsal positional value arrive on the flank could account for the change in distribution of cells with posterior activity, as migration and intercalation displace them to their final locations. Hence, in the case of the forelimb, interaction between the posterior-ventral and anterior-dorsal cells will result in the caudal displacement of the posterior-ventral cells. In the case of the hindlimb, the arrival and growth of anterior-dorsal cells will lead to the apparent gap in the otherwise continuous stretch of cells with posterior properties.

Consistent with these ideas is the fact that although posterior limb properties can be shown to be present on the flank of the chick embryo from stage 9 onward (Hornbruch and Wolpert, 1991), the prospective limb regions are incapable of forming a limb after isolation from their normal surroundings before stage 15 for the wing, and later for the leg (Kieny, 1969, 1971; Pinot, 1969, 1970). Between stages 10 and 15 wings can develop from the relevant flank region only if the normally adjacent somites are included with the graft. Somites from other regions are ineffective substitutes in these experi-
Fig. 12. Limb bud initiation. (a) Interaction at a boundary between prelimb bud flank cells with a posterior-ventral-proximal positional identity (4) and recently immigrating cells with a more anterior-dorsal-proximal position value (10). Arrows indicate directional intercalation inferred from several experiments. See text for details. (b) After intercalation, a limb base with a full circumference has been generated. Distal outgrowth can proceed from this base as described previously (Bryant et al., 1981).

ments. Other experiments have shown that foil barriers separating the prospective wing region in the lateral plate from the somites between stages 10 and 15 inhibit wing development in situ (Stephens et al., 1991). Hence, despite the presence of posterior properties, something further is needed before limb outgrowth can occur. Additional experiments (Stephens et al., 1991) have pinpointed the location of the required material to the intermediate mesenchyme lying between the edge of the somites and the lateral plate mesoderm between stages 13 and 15. Further, it appears that the missing ingredient has arrived in the limb field by stage 15, when grafts of prospective limb fields to the coelom can differentiate into limbs (Kieny, 1969, 1971; Pinot, 1969, 1970). This stage immediately precedes the first detectable signs of limb outgrowth at stage 16. We propose that the ingredient that is missing between stages 9 and 15 is cells that migrate into the limb field from more anterior-dorsal regions of the embryo. The newly arriving cells interact with the posterior-ventral-proximal cells already present, stimulating intercalation and the onset of limb outgrowth.

Three migratory cell populations are candidates for involvement in the establishment of the limb base: neural crest, somitic, and nephric. Although there are not many studies of this region using modern cell tracing techniques, Chevallier (1977) demonstrated a level-specific cellular contribution from the somitic mesoderm to the shoulder girdle in chicks using the chick-quail cell marker. The level specificity of this contribution is in contrast to the lack of specificity in the contribution from the somites to the limb musculature (Chevallier et al., 1977). Since all three of the candidate cell types are located close to one another and are to some degree intermingled, this finding does not specifically rule out either neural crest or nephric cells carried along with the somites as the relevant cells.

In molecular terms, it is possible that the homeobox gene XlHbox 1 identifies the proposed anterior-dorsal cell population that migrates to the forelimb base to initiate outgrowth. Oliver et al. (1988) describe the expression of the XlHbox 1 long protein in a rostral-caudally restricted zone in the pre-limb bud region of the flank of Xenopus. Scattered cells expressing XlHbox 1 protein are present in the myotomes of the pre-limb bud region, and these have been tentatively identified as neural crest cells (Oliver et al., 1988), lending credence to the suggestion that XlHbox 1 cells could migrate ventrally toward the flank. As the limb forms, cells that are positive for XlHbox 1 protein are found in the anterior half of the bud. A similar pattern of protein expression has been documented in the limbs of mice, chick, and zebrafish (Oliver et al., 1988; 1990; Molven et al., 1990). Pronin expression is graded within limb buds, with the highest expression in proximal-anterior cells and lowest in posterior and distal cells. This graded distribution could reflect a dilution of the XlHbox 1 protein as XlHbox 1-positive cells are stimulated to divide as a result of their proposed interactions with the posterior-ventral-proximal cells of the flank. Expansion of the expression domain of XlHbox 1 protein in chick limbs that will later duplicate their pattern after exposure to RA-beads or ZPA grafts (Oliver et al., 1990) could result secondarily from the interaction between the cells that express XlHbox 1 protein and the ZPA cells that are either grafted or converted into ZPA by the RA-bead (Wanek et al., 1991). Duplication of the shoulder girdle is associated with expansion of the expression domain of XlHbox 1 (Oliver et al., 1990), as expected if this gene identifies cells that contribute to the girdle during normal development.

In summary, we suggest that the primary role of RA in limb development is most likely to be in the establish-
ment of flank mesenchyme with posterior-ventral-proximal information with respect to the limb. It is possible that Hensen's node is the source of endogenous RA and that invaginating lateral plate mesoderm cells are exposed to this source, thereby acquiring a posterior-ventral-proximal positional value. A subsequent influx of cells into the limb region from a more anterior-dorsal location is proposed. There is evidence for the forelimb that this population originates in the somites adjacent to the limb region and contributes to the formation of the girdle. It is possible that this population of cells is the same as that which expresses XlHbox 1. This anterior-dorsal cell population provides the necessary levelspecific positional diversity to stimulate intercalation and the formation of the entire limb pattern by mechanisms outlined in detail elsewhere (French et al., 1976; Bryant et al., 1981, 1987).

3. RA and the Transformation of Tails into Limbs

The most recent listing in the catalogue of the developmental consequences of RA is the most amazing to date. Amputated tails of marbled balloon frog tadpoles (Uperodon systoma) exposed to RA during the first few days of regeneration regenerate legs from the tail blastema (Mohanty-Hejmadi et al., 1992) (Fig. 13).

Based on the ideas we have outlined in this article, we offer an interpretation, shown in schematic form in Fig. 14, where we have identified different positions along the rostral-caudal axis of the body with numbers I through VII. We have proposed that the normal role of RA is in the establishment of a trunk region (i.e., level IV) that is distinct from the RA-negative head and tail (see Fig. 11). In development RA exposure converts the gastrulating flank cells of the trunk to a positional value that is equivalent to level IV, also described as posterior-ventral-proximal with respect to the limb. Limb cells exposed to exogenous RA later in development also acquire this same positional value, with the consequences that we have discussed in earlier sections of this article.

In the balloon frog experiments, we propose that RA affects tail blastema cells the same way that it affects limb blastema cells, developing limb cells, and gastrulating flank cells. the cells acquire a positional value that is posterior-ventral-proximal with respect to the limb; in other words, they acquire the flank positional value (level IV), even though they are located at the tail tip. Uniformity of positional value within the blastema is predicted to bring regeneration to a halt, and indeed previous reports of exposure of tail regenerates to RA have shown inhibition of growth (Pietseh, 1987) and truneation (Niazi and Saxena, 1979). In the balloon frog experiments, the animals were removed from RA after a few days. We suggest that after removal from RA, de-differentiation of tail cells continues, bringing cells with level VI positional value into contact with the blastema cells with level IV value. In previous studies we have demonstrated that intercalation occurs along the rostral-caudal axis of newt tails (Iten and Bryant, 1976). In the case of the balloon frog, rostral-caudal intercalation between blastema and stump will lead to a reversed-polarity intercalated region that spans the region from which hind limbs normally develop, shown here as lying between body positions IV and V. Further, the tail regenerate will be able to resume outgrowth using the diverse positional information from the dedifferentiated stump cells that are added to the blastema after RA treatment is ended, in much the same way as proposed for limb regenerates. In this case, regeneration in a caudal direction from level IV will generate another region that spans the position from which hind limbs originate (between IV and V). The prediction that more than a single pair of hindlegs develops in these experiments is borne out by the results (Mohanty-Hejmadi et al., 1992). We interpret the multiple legs illustrated in Fig. 13 as basically two sets of hindlegs (as in the example in Fig. 13a) which are then able to duplicate
whether intercalation along the rostral-caudal axis of the body plays a role in the development of the primary axis. The idea that graded positional qualities could arise from initial discontinuities is not inconsistent with some features of primary axis formation. Axis development is preceded by the generation of a major discontinuity at gastrulation. Several lines of data suggest that the body is initially divided into three regions, head, trunk, and tail (discussed in Section D1 above).

Slack (1991) has recently shown for the origin of the mesoderm in amphibians that when inducing and responding cells are separated by a filter, the induction is incomplete, suggesting that local cell-cell interactions are normally involved. Blumberg et al. (1991) showed that in addition to the homeobox gene goosecoid, thought to be involved in the unique properties of the organizer itself (Cho et al., 1992), the only other classes of homeobox-containing genes present in the organizer region of Xenopus embryos are homologs of labial and caudal, which are involved in specification of the ends of the Drosophila body axis. This led Blumberg et al. (1991) to speculate that the ends of the body axis might be established first, followed by intercalation to generate the middle regions of the axis.

In summary, the apparent homeotic change of tail to limb in regenerating frogs can be understood in terms of the proposed unitary effect of RA in changing pattern formation-competent cells to a posterior-ventral-proximal (i.e., flank) positional value, followed by intercalation along the rostral-caudal axis of the body to generate two additional pairs of hind limb sites on the tail. We raise the issue of a role for intercalation in primary axis formation for further consideration.

E. CONCLUSIONS

In this Review we have described a mechanism by which the diverse effects of RA on limbs can be understood. According to this view, RA has a single effect on pattern formation-competent cells: it converts them to a posterior-ventral-proximal positional value (with respect to the limb) that is synonymous with the positional value of the prelimb bud flank. Short range interactions between cells with discontinuous positional information, followed by growth and the intercalation of intervening qualities of positional information can account for the diverse outcomes of RA treatment. The view we have presented can account for the teratogenic effects of RA, as well as the different effects of RA on the anterior-posterior, dorsal-ventral, and proximal-distal axes of the limb, and for the apparent homeosis from tail to limb. In normal development, RA appears to be involved in the division of the primary body axis into head, trunk, and tail regions and in the establishment of
a population of flank cells with posterior–ventral–proximal specification. Both limb and tail cells retain the ability to respond to added RA throughout the period that they are actively engaged in pattern formation. Focusing on how RA changes the positional identities of cells to posterior–ventral–proximal might be expected to lead to insights into the molecular basis of positional information.

The explanatory power of the polar coordinate model and the role of cell–cell interactions in the development of insects has successfully served as guides for the exploration of the molecular basis of pattern formation (Martinez Arias, 1989; Wilkins and Gubb, 1991). In vertebrates, on the other hand, there has been a tendency to move directly from the tissue level to molecular explanations with the result that analysis at the cellular level has been more or less bypassed. We suggest that the behavior of cells, particularly local cell–cell interactions, and the principles of the polar coordinate model provide a workable cellular level of explanation for the phenomena of limb pattern formation. The challenge is to use the predicted cellular properties to assist in the discovery of the molecules involved. Hence, we need to discover the receptors and ligands that allow cells to detect similarities and differences between themselves and their neighbors, the mechanisms by which this information is transduced into the inhibition or stimulation of growth, the mechanisms by which cells in the cell cycle are able to acquire a molecular identity that is intermediate between those of the surrounding cells, the molecular basis of directional and shortest route intercalation, and the mechanism of distal transformation. All of these and more are direct questions about the molecular basis of pattern formation that arise from an understanding of the underlying cellular properties of the system and that can be used to guide an exploration into the molecular basis of pattern formation.

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REFERENCES


REVIEWS


