

Commentary

Prolactin signaling and Stat5: going their own separate ways?Cathrin Brisken¹, Ayyakkannu Ayyanan¹ and Wolfgang Doppler²¹Swiss Institute for Experimental Cancer Research, Epalinges, Switzerland²University of Innsbruck, Medical Chemistry and Biochemistry, Innsbruck, Austria**Correspondence:** Cathrin Brisken, Swiss Institute for Experimental Cancer Research, Ch. des Boveresses 155, 1066 Epalinges, Switzerland. Tel: +41 21 692 58 51; fax: +41 21 652 69 33; e-mail: Cathrin.Brisken@isrec.unil.ch

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Abstract

Miyoshi *et al.* compared the role of the prolactin receptor (PrIR) and its downstream mediator, the signal transducer and activator of transcription 5 (Stat5), in mammary epithelial cells *in vivo* by studying PrIR^{-/-} and Stat5ab^{-/-} mouse mammary epithelial transplants during pregnancy. At first glance, the two mutant epithelia appear to have similar defects in the differentiation of the alveolar epithelium. However, a closer examination by Miyoshi *et al.* revealed defects in the epithelial architecture of the smallest ducts of Stat5ab^{-/-} transplants not apparent in the PrIR^{-/-} transplants, suggesting that Stat5 is more than a simple mediator of PrIR action.

Keywords: cell adhesion, mammary epithelium, prolactin receptor, Stat5, steroid hormones**Introduction**

Pituitary prolactin is a key regulator of breast development [1–3]. Some prolactin is also produced by the breast epithelium itself, and local prolactin signaling can be deregulated during breast carcinogenesis [4,5]. Prolactin acts via the prolactin receptor (PrIR), a member of the cytokine receptor family [6], and its associated kinase Jak2 [7]. One of the key signaling molecules activated by the PrIR is the signal transducer and activator of transcription 5 (Stat5) [8,9].

A genetic approach to address and compare the contribution of PrIR and its key signaling molecule Stat5 for mammary gland differentiation during pregnancy, by deletion of their genes, is complicated by two instances. First, two closely related Stat5 genes exist, namely Stat5a and Stat5b. They share 92% of their amino acid sequence, are partially functionally redundant, and differ mainly in their carboxy-terminal region [10]. The establishment of Stat5a and Stat5b double knockout mice is therefore mandatory. Second, because PrIR-defective mice [11] or mice carry-

ing inactivated Stat5a and Stat5b (Stat5ab^{-/-}) are infertile [12], it is not possible to study the development of their mammary glands during pregnancy. To overcome the second problem, powerful transplantation techniques have been used (for a review, see [13]). In particular, mammary epithelium from infertile mice can be engrafted to mammary fat pads of wild-type prepubertal mice that are surgically cleared of the endogenous epithelium [14].

PrIR^{-/-} mammary epithelial cells (MECs) engrafted to cleared mammary fat pads of a wild-type host, and thus exposed to a normal endocrine milieu, undergo normal ductal development during puberty but fail to form alveoli and to differentiate into milk-secreting cells during pregnancy [15]. This indicates that the PrIR expressed in MECs is required for alveologenes and the associated differentiation process.

Miyoshi *et al.* used mammary epithelial transplants of Stat5ab^{-/-} epithelium and compared them directly with PrIR^{-/-} epithelial transplants [16]. At first glance, the

IL = interleukin; MAPK = mitogen-activated protein kinase; MEC = mammary epithelial cell; PI3K = PI 3-kinase; PrIR = prolactin receptor; Stat5 = signal transducer and activator of transcription 5.

expectations that the phenotypes of Stat5ab^{-/-} and PrIR^{-/-} epithelia are similar are met. But as Miyoshi *et al.* looked more closely, they uncovered thought-provoking differences that tell us we still have some significant lessons to learn about the biology of these signaling molecules.

Phenotypical similarities of Stat5ab^{-/-} and PrIR^{-/-} epithelia

Whole mount microscopy of mammary glands engrafted with Stat5ab^{-/-} or PrIR^{-/-} epithelium confirms that both mutants form a normal ductal system in the adult virgin host. At the end of pregnancy, however, when the wild-type epithelium has fully expanded and the alveoli are distended by secretion, there is no alveolar development in the PrIR^{-/-} MEC and little development in the Stat5ab^{-/-} MEC. Expression of the milk proteins is substantially reduced in both of the mutant tissues. Together, these data confirm that Stat5 and the PrIR play an essential role in alveolar morphogenesis and differentiation.

Unique defects in the differentiation of the Stat5ab^{-/-} epithelium

As Miyoshi *et al.* examined the structural organization of the epithelium at the end of pregnancy in more detail, a Stat5ab^{-/-} epithelial-specific phenotype became apparent. The intraductal space, clearly discernable even in the smallest ducts of the PrIR^{-/-} epithelium, is partially obliterated in the Stat5ab^{-/-} epithelium. The epithelial cells lining in the smallest branches are of irregular shape and appear disorganized. There are multiple layers of luminal epithelial cells in the Stat5ab^{-/-} transplants and, consequently, crowded lumina.

Electron micrography reveals that microvilli on the apical surface and tight junctions are difficult to find in Stat5ab^{-/-} epithelia, and that the intercellular spaces are disrupted. It remains most puzzling that the unique Stat5ab^{-/-} defect becomes apparent specifically when alveolar morphogenesis is to take place. This suggests an essential and nonredundant role of Stat5 in the induction or repression of genes in the epithelial cells forming the alveoli versus those forming the ducts. A putative function of such genes is the control of intercellular adhesion in the nascent alveoli.

Proliferative response to estrogen and progesterone is more profoundly inhibited in the PrIR^{-/-} MEC compared with the Stat5ab^{-/-} MEC

Miyoshi *et al.* assessed the proliferative response of the mutant engrafted epithelia by injecting animals with estradiol and progesterone, and measuring 5'-bromo-2'-deoxyuridine incorporation 48 hours later. In this assay, both mutant epithelia exhibited a reduced epithelial proliferation when compared with an engrafted wild-type epithelium. Remarkably, the inhibition of the PrIR^{-/-}-defi-

cient MEC was approximately twofold higher than the inhibition of the Stat5ab^{-/-} MEC.

Since at the end of pregnancy both mutant epithelia have perfectly completed ductal morphogenesis, and this process is under the control of estrogen and progesterone [1], it is at first sight surprising that the mutant epithelia respond differently to estradiol and progesterone in the proliferation assay. Possible explanations for the failure to detect a proliferation phenotype at the end of pregnancy are that the time of hormonal stimulation is longer or because there is compensation through many additional stimuli during pregnancy, or both. Indeed, there are other examples of transient phenotypes that disappear by the end of pregnancy (e.g. wnt-4) [17].

Implications for the role of the PrIR-Jak2-Stat5 pathway in MEC differentiation

The similarities in the defects of PrIR^{-/-} and Stat5ab^{-/-} epithelia confirm an important contribution of the PrIR-Jak2-Stat5 pathway to estrogen/progesterone-induced proliferation and alveologenes. There is still much to learn regarding which genes are controlled by this pathway in the mammary epithelium.

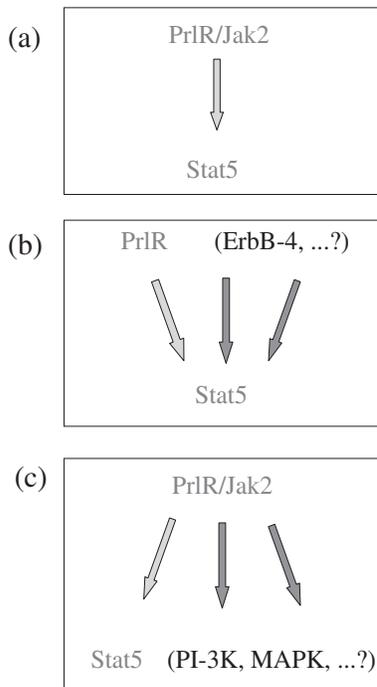
Stat5 has anti-apoptotic activity in hematopoietic tissues [18]. Whether this also applies to the mammary epithelium remains to be established.

By searching for genes controlled by this pathway, Miyoshi *et al.* analyzed the expression of genes that have been previously implicated in cellular adhesion and secretory differentiation. They determined Cx 32 as a putative target gene of the PrIR-Jak2-Stat5 pathway. Cx 32 is a connexin induced at lactation that has a potential role in the establishment of the secretory phenotype. Cx 32 expression is absent in the mutant epithelia. Whether the failure of expression is due to its direct regulation by Stat5 or to a more indirect dependence on the secretory phenotype remains to be established.

The observed subtle differences in the phenotype of PrIR^{-/-} and Stat5ab^{-/-} epithelia indicate that, in addition to the classical linear PrIR-Jak2-Stat5 pathway, Stat5-independent pathways triggered by the activated PrIR and its associated Jak2 contribute to the proliferative response of the epithelium. Other pathways that activate Stat5 but do not involve the PrIR are also mandatory for alveologenes (Fig. 1).

PrIR-independent activation of Stat5

The unique defect of the Stat5ab^{-/-} mutant epithelium indicates a role of additional pathways not dependent on the PrIR for the induction of Stat5. This is supported by the reported activation of Stat5 in the mammary epithelium by epidermal growth factor [19] and by the requirement

Figure 1

Interdependence of prolactin receptor (PrIR) and signal transducer and activator of transcription 5 (Stat5) signaling. **(a)** The 'classical' PrIR-Jak2-Stat5 pathway contributes to alveolar development. **(b)** Alternative routes to activate Stat5 are required for alveologenesis. **(c)** Potential contribution of multiple PrIR-dependent pathways to the proliferative response to estrogen and progesterone. PI-3K, PI 3-kinase; MAPK, mitogen-activated protein kinase.

for ErbB-4-triggered activation of Stat5 to trigger lobulo-alveolar development during pregnancy [20].

Stat5-independent pathways triggered by the activated PrIR

As observed with other cytokine receptors with associated Jak kinases, activation of the PrIR/Jak2 by prolactin leads to the induction of other signaling pathways, such as the mitogen-activated protein kinase (MAPK) and PI 3-kinase (PI3K) pathways. The relative signaling output of these different pathways appears to be crucial for the biological effect, indicating that the regulation of the balance between these pathways is of utmost importance for development and differentiation.

A recent example for a specific regulator of this balance is the suppressor of cytokine signaling SOCS-3, which has been shown to lead to the selective inhibition of Stat5 signaling but not of the MAPK pathway after IL-2 stimulation [21]. SOCS-3 might have a similar function in the mammary gland, where it is differentially regulated during development [22].

A further point to consider is that extensive crosstalk occurs between prolactin and other hormonal signaling pathways; that is, estrogen induces expression of the progesterone receptor [23] and PrIR [24], prolactin signaling induces the expression of the estrogen receptor ER α and ER β in granulosa cells [25,26], and prolactin has been shown to upregulate estrogen receptor expression in cultured mammary epithelial cells [27]. It is therefore important to examine the expression of the estrogen receptor and the progesterone receptor in the mutant epithelia to assess whether the number of cells, which are responsive to the hormones, is reduced in the mutant epithelia.

Finally, another possibility is that local prolactin signaling is important to the proliferative response to estrogen and progesterone. Both estrogen and progesterone act by paracrine mechanisms to induce proliferation, and it is conceivable that locally secreted prolactin is one of the mediators of these effects.

Conclusion

Stat5 was originally cloned as the 'mammary gland factor' and functionally characterized as a mediator of prolactin-induced β -casein expression. The literature suggested that the basic role of Stat5 in the mammary gland was to mediate prolactin signaling, while the PrIR in turn relied heavily on Stat5 to mediate its effects.

Miyoshi *et al.* provide evidence that there is more than this simple mutual relationship between PrIR and Stat5. For example, inactivation of Stat5 leads to much more severe defects in the intercellular adhesion of epithelial cells than PrIR deletion, whereas PrIR deletion has a more dramatic effect on proliferation than a deletion of Stat5. Further analysis of the role of these signaling pathways will provide important insight into mammary gland morphogenesis and differentiation.

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