

Basis For Abnormal Desquamation And Permeability Barrier Dysfunction in RXLI

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Mutations in the gene for steroid sulfatase (SSase), are responsible for recessive x-linked ichthyosis (RXLI). As a consequence of SSase deficiency, its substrate, cholesterol sulfate (CSO₄), accumulates in the epidermis. Accumulation of this amphipathic lipid in the outer epidermis provokes both a typical scaling phenotype and permeability barrier dysfunction. Research on RXLI has illuminated several, potentially overlapping pathogenic mechanisms and provided insights about the role of SSase and CSO₄ in normal differentiation, barrier maintenance, and desquamation. We now show here that SSase is concentrated in lamellar bodies (LB), and secreted into the SC interstices, along with other LB-derived lipid hydrolases. There, it degrades CSO₄, generating some cholesterol for the barrier, while the progressive decline in CSO₄ (a serine protease (SP) inhibitor) permits corneodesmosome (CD) degradation leading to normal desquamation. Two molecular pathways contribute to disease pathogenesis in RXLI: 1) excess CSO₄ produces nonlamellar phase separation in the stratum corneum (SC) interstices, explaining the barrier abnormality. 2) The increased CSO₄ in the SC interstices inhibit activity sufficiently to delay CD degradation, leading to corneocyte retention. We also show here that increased Ca⁺⁺ in the SC interstices in RXLI could contribute to corneocyte retention, by increasing CD and interlamellar cohesion. RXLI represents one of the best understood diseases in dermatology – from the gene to the SC interstices, its etiology and pathogenesis are becoming clear, and assessment of disease mechanisms in RXLI led to new insights about the role of SSase and CSO₄ in epidermis terminal differentiation.

Key words: cholesterol sulfate/cholesterol/transepidermal water loss/serine proteases/corneodesmosomes/barrier function/ichthyosis/X-linked ichthyosis.

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The Molecular Basis for Recessive X-Linked Ichthyosis

In 1978 recessive X-linked ichthyosis (RXLI) was linked to deficiency of the microsomal enzyme, steroid sulfatase (SSase) (Koppe *et al*, 1978; Webster *et al*, 1978). The SSase gene has been the subject of considerable research because of its location on the distal tip of the short arm of the X chromosome (Tiepolo *et al*, 1977; Mohandas *et al*, 1979; Tiepolo *et al*, 1980; Shapiro, 1985; Li *et al*, 1990). Patients with RXLI display gene mutations/deletions (Ballabio *et al*, 1987; Bonifas *et al*, 1987; Conary *et al*, 1987; Gillard *et al*, 1987; Shapiro *et al*, 1989), resulting in ichthyosiform skin changes with occasional extracutaneous

organ system involvement, due to contiguous gene syndromes (Schmickel, 1986; Schnur and Trask, 1989).

SSase is a 62 kDa microsomal enzyme, responsible for hydrolyzing the 3 β -sulfate esters from both cholesterol sulfate (CSO₄) and sulfated steroid hormones, generating their nonsulfated counterparts. As a result of enzyme deficiency in RXLI, CSO₄ accumulates in skin, plasma, and red cell membranes (up to 10- to 20-fold increase (Bergner and Shapiro, 1988; Epstein *et al*, 1981). CSO₄ is carried in blood by the low-density (β -lipoprotein) fraction, producing diagnostic electrophoretic alterations (Epstein *et al*, 1981; Ibsen *et al*, 1986). The fact that CSO₄ levels in scale are still higher than levels in blood presumably explains the prominence of skin involvement in RXLI (Bergner and Shapiro, 1988; Epstein *et al*, 1981; Williams and Elias, 1981).

Steroid sulfatase and cholesterol sulfate in normal epidermal physiology To examine the regulation of epidermal CSO₄ content by SSase activity in relation to normal cohesion and desquamation, epidermal cell populations were prepared from normal and RXLI epidermis, and assayed for both SSase activity and CSO₄ content (Elias *et al*, 1984). Whereas SSase activity is low in the basal and spinous layers, its levels peak in the SG (10–20 times

Abbreviations: CD, corneodesmosomes; CE, cornified envelope; Chol, cholesterol; CLE, corneocyte-bound lipid envelope; CSO₄, cholesterol sulfate; HMGCoA, hydroxymethyl glutaryl coenzyme A; LB, lamellar bodies; RXLI, recessive X-linked ichthyosis; SC, stratum corneum; SCCE, stratum corneum chymotryptic enzyme; SCTE, stratum corneum tryptic enzyme; SG, stratum granulosum; SPI, serine protease inhibitor; SSase, steroid sulfatase; TEWL, transepidermal water loss; TG-1, transglutaminase 1.

higher), and persist at equally high levels in the SC (Fig 1), where it is concentrated in membrane domains, primarily in the lower SC (Fig 2) In contrast, SSase activity is absent from both epidermal nucleated cell layers and the SC in RXLI (Kubilus *et al*, 1979; Elias *et al*, 1984). Normally, CSO₄ comprises about 5% of the total lipid of human stratum granulosum (SG), declining to about 1% of lipid mass in the outer SC, through ongoing hydrolysis of CSO₄ during SC transit (Fig 1). In RXLI, where SSase activity is absent, the SC contains up to 12% CSO₄ by lipid weight (\approx 1% of total weight), a concentration that exceeds that in hair, nail, and ungulate hoof, where CSO₄ can account for 2–5% of total weight.

Because cholesterol sulfotransferase (sulf2Bib) activity predominates in lower nucleated cell layers, while SSase peaks in the outer epidermis, Epstein *et al* (1984) proposed an “epidermal cholesterol sulfate cycle”, where cholesterol is first sulfurylated in the lower epidermis and then desulfated in outer epidermal layers, regenerating cholesterol (Fig 1). Moreover, CSO₄ levels are normally lower in the outer SC than in lower SC (Long *et al*, 1985; Ranasinghe *et al*, 1986), supporting the concept that ongoing hydrolysis of CSO₄ by SSase during SC transit may be critical for normal desquamation (Elias *et al*, 1984; Ranasinghe *et al*, 1986). The “cholesterol sulfate cycle”, however, has significance that extends well beyond desquamation (see below).

Since SSase activity reportedly is not concentrated in lamellar body (LB)-enriched fractions of whole SG (Grayson *et al*, 1985), how the enzyme reaches membrane domains has remained problematic. Though a classic microsomal enzyme, ultrastructural immunocytochemical studies have also localized the enzyme to the plasma membrane, where it localizes to coated pits (Willemssen *et al*, 1988; Dibbelt *et al*, 1989). To address the question of SSase delivery to SC, we developed a new cytochemical method, utilizing both lead- and barium-capture methods, for the ultrastructural localization of SSase activity (see legend to Fig 2 for details about one of these methods). Using these techniques, we localized SSase not only within the cytosol (i.e., microsomes), but also within LB (Fig 2B). We further demonstrated delivery of SSase to the SC interstices following LB secretion, where its activity persists into the lower SC layers, but disappears from the outer SC (Fig 2A). Thus, SSase utilizes the LB secretory system to reach sites where it participates in the regulation of barrier homeostasis and desquamation like other lipid hydrolases that are involved in the extracellular processing of secreted polar lipids (Elias and Menon, 1991).

Like SSase, CSO₄ is concentrated in the SC interstices, but in contrast to other lipid precursors, it is not present in LB (Grayson *et al*, 1985). Yet, its mode of delivery to the SC interstices can be explained readily by both its lipophilicity

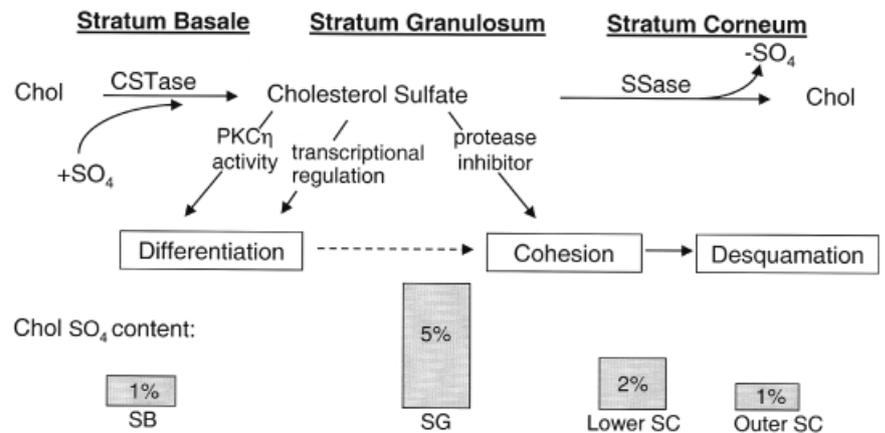
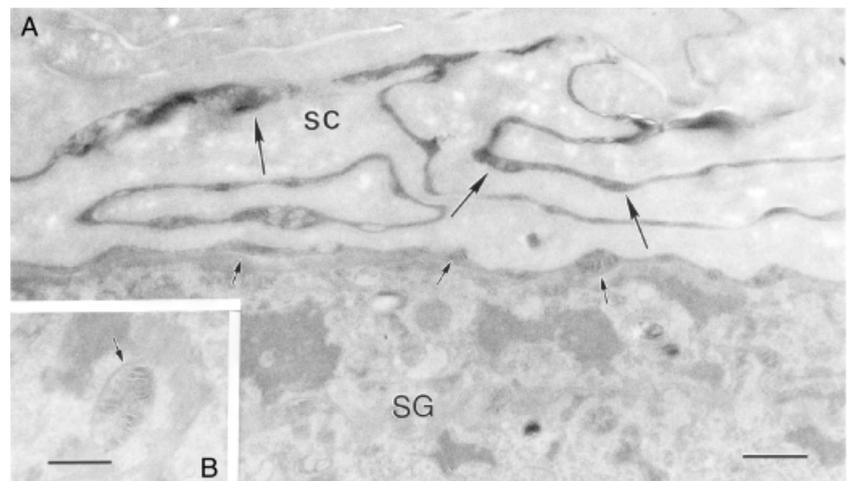


Figure 1
Cholesterol sulfate cycle in normal epidermis. Role of steroid sulfatase and cholesterol sulfate in normal epidermal physiology. Abbreviations: CSTase, cholesterol sulfotransferase; SSase, steroid sulfatase; Chol, cholesterol; PKC, protein kinase C; Chol SO₄, cholesterol sulfate; SB, stratum basale; SG, stratum granulosum; SC, stratum corneum.

Figure 2
Steroid sulfatase activity localizes to lamellar bodies and at the stratum corneum interstices. A: Enzyme activity, shown as lead precipitate (arrows), localizes to the interstices of lower SC layers. B: SSase activity localizes to lamellar body contents and limiting membranes (not shown). Abundant activity also is seen at the stratum granulosum (SG)-stratum corneum (SC) interface. Human skin samples were aldehyde-fixed, rinsed in HEPES buffer, and microwave incubated in 5 mM 4-methylumbelliferyl sulfate and 2 mg/mL lead nitrate in HEPES buffer (Rassner *et al*, 1997). Post-fixation was in reduced osmium tetroxide with microwave irradiation for 2.5 min at 37°C. A & B, Scale bars: A: 1.0 μ m; B: 0.5 μ m.



and its amphiphilicity. CSO_4 , like many oxygenated sterols (but unlike cholesterol), readily diffuses across the keratinocyte plasma membrane (Ponec and Williams, 1986). Therefore, in the absence of a lipid milieu within corneocytes, CSO_4 likely partitions preferentially to the highly hydrophobic, extracellular domains of the SC, based purely upon its solubility characteristics; i.e., it does not require delivery by a secretory organelle.

Mechanisms proposed to perturb permeability barrier homeostasis in RXLI Although patients with RXLI display only a minimal basal barrier abnormality (Lavrijsen *et al*, 1993), they demonstrate a pronounced delay in recovery kinetics following acute perturbations (Zettersten *et al*, 1998). Moreover, topically applied CSO_4 not only induces scaling (Maloney *et al*, 1984), but also a barrier abnormality (Zettersten *et al*, 1998), lending further credence to the hypothesis that excess CSO_4 plays a destabilizing role in SC barrier function. Several mechanisms have been proposed to contribute to the barrier abnormality in RXLI (Fig 3). First, CSO_4 fails to form eutectic mixtures with other SC lipids; hence, excess CSO_4 segregates into distinct nonlamellar domains in model lipid mixtures (= phase separation) (Rehfeld *et al*, 1986). Indeed, ultrastructural images of SC in RXLI show extensive, but focal nonlamellar domains, with disruption of the extracellular lamellae (Fig 3; see also Zetterston *et al*, 1998).

However, the barrier abnormality is probably due not only to excess CSO_4 , but also to decreased cholesterol (the cholesterol content of the SC in RXLI is reduced by approximately 50%) (Williams and Elias, 1981), and decreased cholesterol alone can produce abnormal extracellular lamellar membranes (Mao-Qiang *et al*, 1993). Two unrelated mechanisms have been invoked to explain the decrease in cholesterol in RXLI (Fig 4): 1) reduced generation of cholesterol from CSO_4 due to the enzyme deficiency; (2) CSO_4 -mediated inhibition of HMGCoA reductase, the rate-limiting enzyme of cholesterol synthesis (Williams *et al*, 1987). Thus, increased CSO_4 levels in RXLI could inhibit epidermal cholesterol synthesis, explaining the reduced levels of cholesterol in the SC of RXLI. Yet, topical cholesterol only partially improves barrier function in RXLI, and it only partially reverses the barrier abnormality in CSO_4 -treated mice (Maloney *et al*, 1984), suggesting that it is increased CSO_4 , rather than insufficient cholesterol, that accounts for the barrier abnormality in RXLI. Finally, CSO_4 inhibits transglutaminase 1 (TG-1)-mediated attachment of ω -hydroxyceramides to the cornified envelope (CE) *in vitro*, a step that forms the cornified-bound lipid envelope (CLE) (Nemes *et al*, 2000). Together, the CE/CLE forms a scaffold for the deposition and supramolecular organization of the SC lamellar bilayers, and in lamellar ichthyosis, decreased TG-1 activity leads to a defective CE scaffold, accounting in part for the barrier abnormality (Elias *et al*, 2002). Yet,



Figure 3
Abnormalities in barrier function in RXLI are due to lamellar-phase separation. Lamellar bilayers appear normal between numerous foci of phase separation in RXLI (A, arrows). (Sample biopsies obtained under our protocol approved by the UCSF Committee on Human Research, according to Declaration of Helsinki Guidelines). Ruthenium tetroxide postfixation. Scale bar = 0.25 μm .

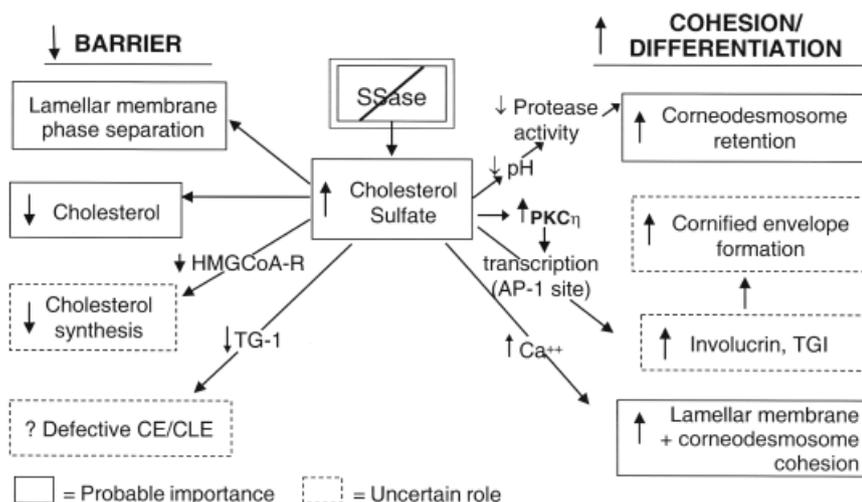


Figure 4
Pathogenic mechanisms in RXLI. Abbreviations: CE; cornified envelope; CLE, corneocyte-bound lipid envelope; TG-1, transglutaminase 1; HMGCoA-R, hydroxymethyl glutaryl/Coenzyme A reductase (for others see legend to Figure 1).

despite the proposition that increased CSO_4 could alter the barrier in RXLI via an effect on TG-1-mediated scaffold function (Nemes *et al*, 2000), the CE/CLE scaffold has not been demonstrated to be abnormal in RXLI.

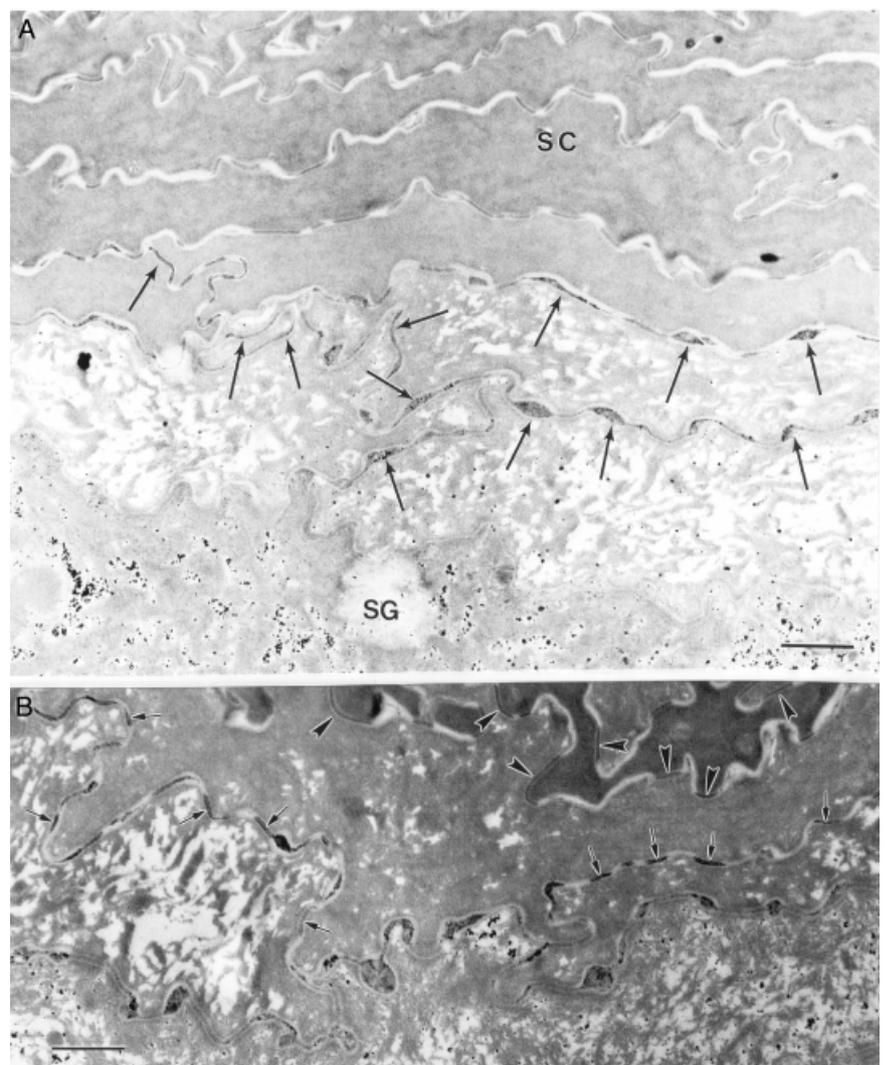
Mechanisms proposed to cause abnormal desquamation in RXLI The hyperkeratosis in RXLI is attributed to delayed desquamation rather than to excessive hyperproliferation (Williams, 1991). Both epidermal proliferation indices and transit times are normal, consistent with an absence of significant acanthosis in RXLI (Frost and Van Scott, 1966). Thus, RXLI is classified as a classic, retention-type of ichthyosis, with delayed degradation of corneodesmosomes; i.e., these structures persist high into the SC (Figs 4, 5). Two key serine proteases, the SC chymotryptic enzyme (SCCE) and the SC tryptic enzyme (SCTE), are thought to mediate desquamation, because both can degrade corneodesmosomes *in vitro* (Brattsand *et al*, 2000). While the activity of these enzymes is restricted by the acidic pH of normal SC (SCCE and SCTE exhibit neutral pH optima), the pH of the SC in RXLI is lower than that of normal SC (Ohman and Vahlquist, 1998). Hence, we hypothesized that these serine proteases might be less operative in RXLI than in normal SC. As seen in Fig 6, serine

protease activity appears to be much higher in normal than in RXLI SC (see legend for experimental details). Thus, the reduced pH of SC alone can account for the increased cohesion in RXLI.

CSO_4 could also increase SC retention through its known activity as a serine protease inhibitor (SPI): For example, CSO_4 inhibits sperm capacitation, a serine protease-catalyzed reaction (cited in Williams, 1991), and CSO_4 inhibits SC serine proteases *in vitro* (Sato *et al*, 1998). However, whether CSO_4 functions as an SPI in epidermis and its importance in relation to other known SPI of the SC (e.g., elafin, plasminogen activator inhibitor I, secretory leukocyte protease inhibitor, cystatin) is not known.

A third, unrelated mechanism which could result in increased SC cohesion posits that Ca^{++} , if present in sufficient quantities, could cross-link the highly anionic SO_4 group on adjacent lamellar bilayers (Epstein *et al*, 1984). Indeed, CSO_4 -containing liposomes aggregate avidly in the presence of calcium (Abraham *et al*, 1987). Moreover, Ca^{++} , which is required for desmosome stabilization, could also stabilize attachments between opposing corneodesmosomes. To address this potential mechanism, we employed ion precipitation cytochemistry to localize Ca^{++} in RXLI *versus*. normal SC (Menon *et al*, 1985). Whereas

Figure 5
Calcium preferentially localizes to corneodesmosomes in extracellular spaces of lower SC in RXLI. Ca^{++} precipitates in normal epidermis do not extend above the stratum granulosum (SG); i.e., into the stratum corneum (SC) (Menon *et al*, 1985). In contrast, Ca^{++} precipitates extend above the SG-SC interface in RXLI, where they are restricted to the extracellular domains (Figure 5A). Corneodesmosomes are increased at all levels of SC (arrowheads). In many sites, Ca^{++} appears to specifically segregate with corneodesmosomes (arrows). A+B, osmium tetroxide postfixation. Scale bars = 1 μm .



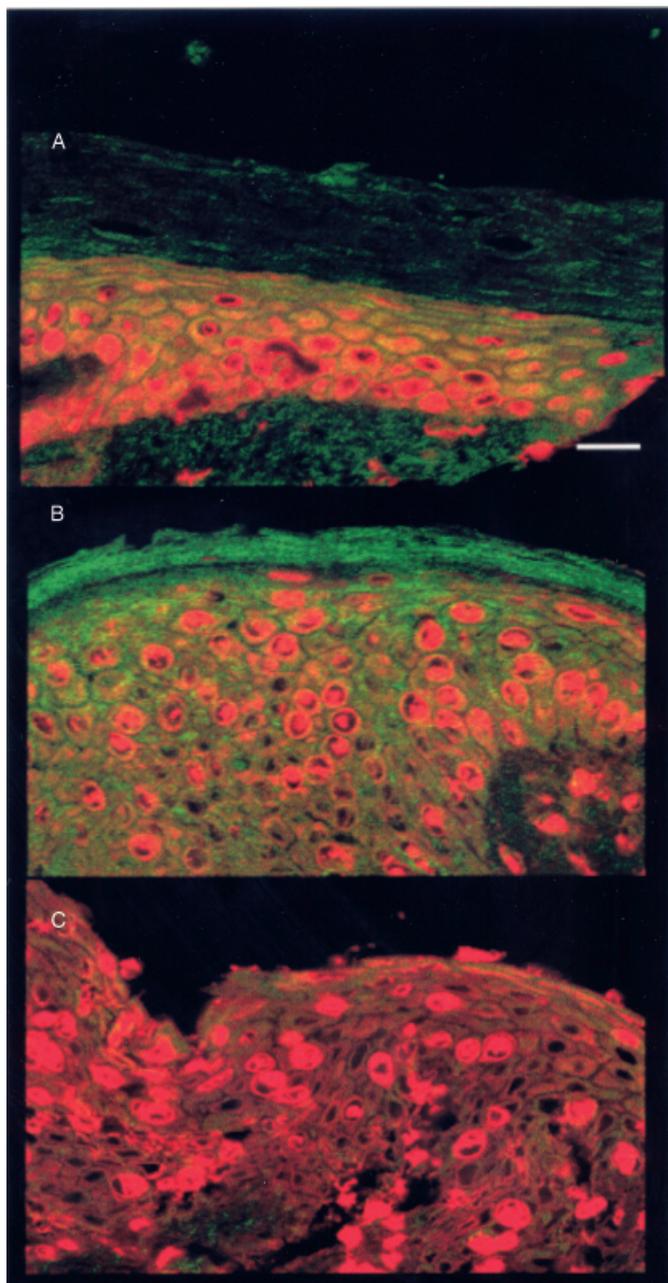


Figure 6

Serine protease activity is reduced in RXLI SC. Frozen sections (8 μm) from an X-linked ichthyosis adult (A) and from the surgical excision margins of an adult normal (B) were rinsed with a washing solution (1% Tween 20 in deionized water) and incubated at 37°C for 1 h with 250 μL of BODIPY-FI-casein in either deionized water (2 $\mu\text{L}/\text{mL}$), 1 mM cholesterol sulfate in DMSO, or DMSO alone, as described recently (Hachem *et al*, 2003); C: normal + cholesterol sulfate. All sections then were rinsed with the 1% Tween 20 washing solution, coverslipped, and visualized under a confocal microscope (Leica TCS SP, Heidelberg, Germany) at an excitation wavelength of 485 nm and an emission wavelength of 530 nm. Scale bar = 10 μm .

lower SC; and therefore, this mechanism, along with Ca^{++} bridging of adjacent lamellar bilayers (Epstein *et al*, 1984), could further contribute to increased corneocyte retention in RXLI.

Regulation of differentiation by cholesterol sulfate Sulfation of cholesterol by cholesterol sulfotransferase is a step that is intimately linked to differentiation in keratinizing epithelia. For example, CSO_4 levels are several orders of magnitude higher in keratinizing than in mucosal epithelia, and induction of mucous metaplasia in keratinizing epithelia by application of exogenous retinoids induces a dramatic decline in CSO_4 levels. CSO_4 is known to stimulate epidermal differentiation by at least two related mechanisms (Fig 4). 1) It activates the η isoform of protein kinase C (Denning *et al*, 1995), which in turn stimulates the phosphorylation of differentiation-linked proteins, assessed as increased CE formation (Fig 1). 2) CSO_4 is a transcriptional regulator of TG-1 and involucrin expression, operating through an AP-1 binding site in the promoter region (Kawabe *et al*, 1998; Hanley *et al*, 2001). It is likely that these two mechanisms are linked as shown in Fig 4: PKC activation by CSO_4 could phosphorylate AP-1, leading to enhanced transcriptional regulation of TG-1 and involucrin. A key issue that remains unresolved, however, is whether the increased levels of CSO_4 in RXLI cause increased corneocyte differentiation, and whether such increased differentiation could provoke corneocyte retention. Such a mechanism could also, however, potentially explain the unique properties of hair, nail, and ungulate hooves, where CSO_4 levels are much higher than in normal SC, and even higher than in the SC of RXLI.

In summary, we have traced the evolution of our understanding of the pathogenesis of RXLI, one of the best understood of all dermatologic disease entities. Elucidation of pathogenic mechanisms in RXLI has led to new concepts about the normal regulation of epidermal differentiation, permeability barrier homeostasis, and cohesion/desquamation.

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Ca^{++} is largely excluded from the intercellular spaces of normal SC (not shown; see Menon *et al*, 1985), the lower SC in RXLI demonstrates abundant Ca^{++} in extracellular domains (Fig 5A,B). Moreover, the Ca^{++} preferentially localizes along the external face of opposing corneodesmosomes (Fig 5B). Thus, the delayed degradation of corneodesmosomes in RXLI correlates with increased Ca^{++} in the

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