CD207⁺ CD103⁺ dermal dendritic cells cross-present keratinocyte-derived antigens irrespective of the presence of Langerhans cells Sandrine Henri, Lionel Franz Poulin, Samira Tamoutounour, Laurence Ardouin, Martin Guilliams, Béatrice de Bovis, Elisabeth Devilard, Christophe Viret, Hiroaki Azukizawa, Adrien Kissenpfennig, and Bernard Malissen

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The authors regret that data was presented incorrectly in Table I of their article. The errors have been corrected in the html and pdf versions of the article. The corrected table is presented below:

Table I. Phenotype and frequency of the five skin DC subsets prior to and following their migration to CLNs

Skin subset	Phenotypes	Frequency
		%
Epidermis		
Epidermal LCs	CD207 ⁺⁺ , CD11b ^{int} , CD103 ⁻ , EpCAM ⁺⁺ , CD24 ⁺ , Sirp α ⁺	100 ± 1
Dermis		
LCs in transit	CD207 ⁺⁺ , CD11b ^{int} , CD103 ⁻ , EpCAM ⁺⁺ , CD24 ⁺ , Sirp α ⁺	6.4 ± 1
CD207+ CD103- DDCs	CD207+, CD11blow, CD103-, EpCAM-/low, CD24+, Sirpa-	2.8 ± 0.5
CD207+ CD103+ DDCs	CD207+, CD11b ^{low} , CD103+, EpCAM $^{-/low}$, CD24+, Sirp α^-	2.6 ± 0.6
CD207- CD11b- DDCs	CD207 ⁻ , CD11b ^{-/low} , CD103 ⁻ , EpCAM ⁻ , CD24 ⁻ , Sirpα ^{low}	16.3 ± 1.3
CD207 CD11b+ DDCs	CD207 ⁻ , CD11b ^{high} , CD103 ⁻ , EpCAM ⁻ , CD24 ⁻ , Sirp α ⁺	65.8 ± 5
CLNs		
mLCs	CD207+, CD11b $^{-/low} \downarrow$, CD103 $^-$, EpCAM+, CD24+, Sirp α +	9.5 ± 2.9
CD207+ CD103- mDDCs	CD207+, CD11b $^{-/low}$, CD103 $^-$, EpCAM+ \uparrow , CD24+, Sirp α + \uparrow	7.0 ± 4.2
CD207+ CD103+ mDDCs	CD207+, CD11b $^{-llow}$, CD103+, EpCAM+ \uparrow , CD24+, Sirp α^{low}	13.5 ± 4.5
CD207 CD11b mDDCs	CD207 $^-$, CD11b $^-$, CD103 bim \uparrow , EpCAM bim \uparrow , CD24 bim \uparrow , Sirp α $^+$ \uparrow	20.8 ± 2.5
CD207- CD11b+ mDDCs	CD207 $^-$, CD11b $^+$, CD103 bim \uparrow , EpCAM int \uparrow , CD24 int \uparrow , Sirp α^+	35.2 ± 2.6

The percentage \pm SEM of each subset among MHCII⁺ (epidermis and dermis) or MHCII^{high} CD11c^{inter-to-high} (CLN) cells were calculated from at least six independent experiments. Upward and downward pointing arrows indicate markers that were up-regulated or down-regulated, respectively, upon migration to CLNs. Bim, bimodal.