Figure 1 sup: Consort diagram of biological samples used in this study
**Figure 2 sup:** CCL5 and CD68/CD14 expression in DLBCL tumors

Correlation plots for *CCL5* and *CD68* or *CD14* mRNA expressions in tumor lesions are shown. *p* values and Pearson correlation coefficients (*r*) are indicated.
Figure 3 sup: expression of CCL and CD68 among DLBCL subtypes

CCL5 and CD68 mRNA expression among GC, ABC and unknown (unk.) DLBCL molecular subtypes. Differences were analyzed by one-way ANOVA.
Figure 4 sup: **CCL5 expression in healthy B cells**

CCL5 mRNA expression was analyzed for different stages of B cell differentiation. n ≥ 4 samples per cell population.
Figure 5 sup: chemokine expression in OCI-Ly6 and OCI-Ly18
A) Supernatants from OCI-Ly6 and OCI-Ly18 were analyzed for chemokine expression as in figure 1B. Control culture medium was also analyzed. 1:CCL1, 2:CCL2, 3:CCL3/4. Note that dot order was changed by the manufacturer between the two experiments. B) A quantification with the mean and the min/max value of the signals obtained for monocyte-specific chemokines is provided.
Figure 6 sup: CCL5 receptor expression in DLBCL tumors
Correlation plots for CCL5, CD14 or CD68 with CCR1/3/5 mRNA expressions in DLBCL tumor lesions are shown. p values and Pearson correlation coefficients (r) are indicated.