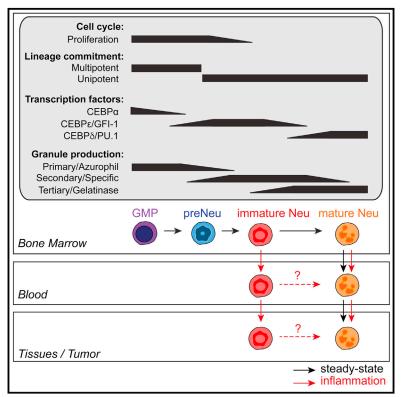
Immunity

Developmental Analysis of Bone Marrow Neutrophils Reveals Populations Specialized in Expansion, Trafficking, and Effector Functions

Graphical Abstract



Authors

Maximilien Evrard, Immanuel W.H. Kwok, Shu Zhen Chong, ..., Andrés Hidalgo, Florent Ginhoux, Lai Guan Ng

Correspondence

maximilien.evrard@unimelb.edu.au (M.E.), ng_lai_guan@immunol.a-star. edu.sg (L.G.N.)

In Brief

The neutrophil differentiation pathway is poorly defined. Evrard et. al. demonstrate a workflow of characterizing bone marrow neutrophil subsets on the basis of their proliferative capacity and molecular signatures and thereby define the developmental trajectory and functional properties of neutrophils.

Highlights

- Proliferation activity identifies committed neutrophil precursor in mice and humans
- Neutrophil subsets possess distinct transcriptomic and functional signatures
- Defect in neutrophil development leads to impaired neutrophil-mediated responses
- Increased circulating immature neutrophils are associated with cancer progression

Evrard et al., 2018, Immunity *48*, 364–379 February 20, 2018 © 2018 Elsevier Inc. https://doi.org/10.1016/j.immuni.2018.02.002



Developmental Analysis of Bone Marrow Neutrophils **Reveals Populations Specialized in Expansion,** Trafficking, and Effector Functions

Maximilien Evrard,^{1,10,11,12,*} Immanuel W.H. Kwok,^{1,2,11} Shu Zhen Chong,¹ Karen W.W. Teng,¹ Etienne Becht,¹ Jinmiao Chen,¹ Je Lin Sieow,¹ Hweixian Leong Penny,¹ Goh Chi Ching,⁷ Sapna Devi,¹ José Maria Adrover,³ Jackson L.Y. Li,^{1,3} Ka Hang Liong,¹ Leonard Tan,¹ Zhiyong Poon,⁴ Shihui Foo,¹ Jia Wang Chua,¹ I-Hsin Su,² Karl Balabanian,⁵ Françoise Bachelerie,⁵ Subhra K. Biswas,¹ Anis Larbi,¹ William Y.K. Hwang,⁴ Vikas Madan,⁶ H. Phillip Koeffler,^{6,7,8} Siew Cheng Wong,¹ Evan W. Newell,¹ Andrés Hidalgo,^{3,9} Florent Ginhoux,¹ and Lai Guan Ng^{1,2,13,*} ¹Singapore Immunology Network, Agency for Science, Technology and Research, Biopolis, 138648 Singapore

²School of Biological Sciences, Nanyang Technological University, 637551 Singapore

³Area of Cell and Developmental Biology, Fundación Centro Nacional de Investigaciones Cardiovasculares, Madrid 28029, Spain

⁴Department of Hematology, Singapore General Hospital, 169856 Singapore

⁵Inflammation Chemokines and Immunopathology, INSERM, UMR 996, Faculté de Médecine, Université Paris-Sud, Université Paris-Saclay, 92140 Clamart, France

⁶Cancer Science Institute of Singapore, National University of Singapore, 117599 Singapore

⁷Cedars-Sinai Medical Center, Division of Hematology/Oncology, UCLA School of Medicine, Los Angeles, CA, USA

⁸Department of Hematology-Oncology, National University Cancer Institute of Singapore, National University Hospital, 119074 Singapore

⁹Institute for Cardiovascular Prevention, LMU, Munich 80336, Germany

¹⁰Present address: Department of Microbiology and Immunology, Peter Doherty Institute for infection and Immunity, University of Melbourne, Melbourne, Victoria 3000, Australia

¹¹These authors contributed equally

¹²Present address: Department of Microbiology and Immunology, Peter Doherty Institute for Infection and Immunity,

The University of Melbourne, Melbourne, Victoria 3000, Australia

13Lead contact

*Correspondence: maximilien.evrard@unimelb.edu.au (M.E.), ng_lai_guan@immunol.a-star.edu.sg (L.G.N.) https://doi.org/10.1016/j.immuni.2018.02.002

SUMMARY

Neutrophils are specialized innate cells that require constant replenishment from proliferative bone marrow (BM) precursors as a result of their short half-life. Although it is established that neutrophils are derived from the granulocyte-macrophage progenitor (GMP), the differentiation pathways from GMP to functional mature neutrophils are poorly defined. Using mass cytometry (CyTOF) and cellcycle-based analysis, we identified three neutrophil subsets within the BM: a committed proliferative neutrophil precursor (preNeu) which differentiates into non-proliferating immature neutrophils and mature neutrophils. Transcriptomic profiling and functional analysis revealed that preNeu require the C/EBP ε transcription factor for their generation from the GMP, and their proliferative program is substituted by a gain of migratory and effector function as they mature. preNeus expand under microbial and tumoral stress, and immature neutrophils are recruited to the periphery of tumor-bearing mice. In summary, our study identifies specialized BM granulocytic populations that ensure supply under homeostasis and stress responses.

INTRODUCTION

Neutrophils are the most abundant immune cell type in human peripheral blood and act as the first responders during sterile and microbial insults. They elicit powerful effector functions to eliminate foreign threats and play crucial roles in tissue remodeling (Borregaard, 2010; Kolaczkowska and Kubes, 2013). Neutrophils are short lived; they have an estimated half-life of 19 hr in humans (Lahoz-Beneytez et al., 2016; Tak et al., 2013). Therefore, neutrophils must be constantly replenished; an impairment in their production and migration leads to neutropenia and lifethreatening conditions (Summers et al., 2010).

Historically, neutrophil development has been defined by histological staining and electron microscopy into stages based on neutrophil size, nucleus morphology, and cytosol coloration. After maturation, neutrophils are retained in the BM through CXCR4 chemokine receptor signaling, and CXCR2 signaling drives their release into the circulation (Devi et al., 2013). During inflammation, increased amounts of granulocyte-colony-stimulating factor (G-CSF) can potentiate mobilization of neutrophils from the BM by lowering the threshold of their release (Kim et al., 2006) and increasing the amounts of mobilizing signals (i.e., CXCL1) (Köhler et al., 2011).

It is believed that neutrophils consist of a homogeneous population. However, this view is rapidly evolving thanks to increasing reports of neutrophil heterogeneity (reviewed in (Silvestre-Roig et al., 2016)). Notably, these studies focused

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