

2018



Progetto Ematologia Romagna

Meccanismo oncogenetico dell'infiammazione

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Pavia



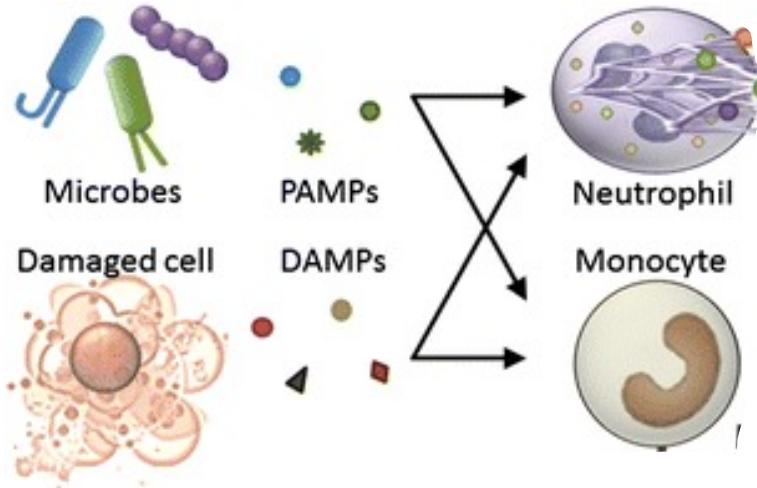
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Inflammation

Protective response of the organism to stimulation by invading pathogens or endogenous signals such as damaged cells, thus resulting in the elimination of the initial cause of injury, the clearance of necrotic cells and tissue repair



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PAMPs (pathogen-associated molecular patterns):

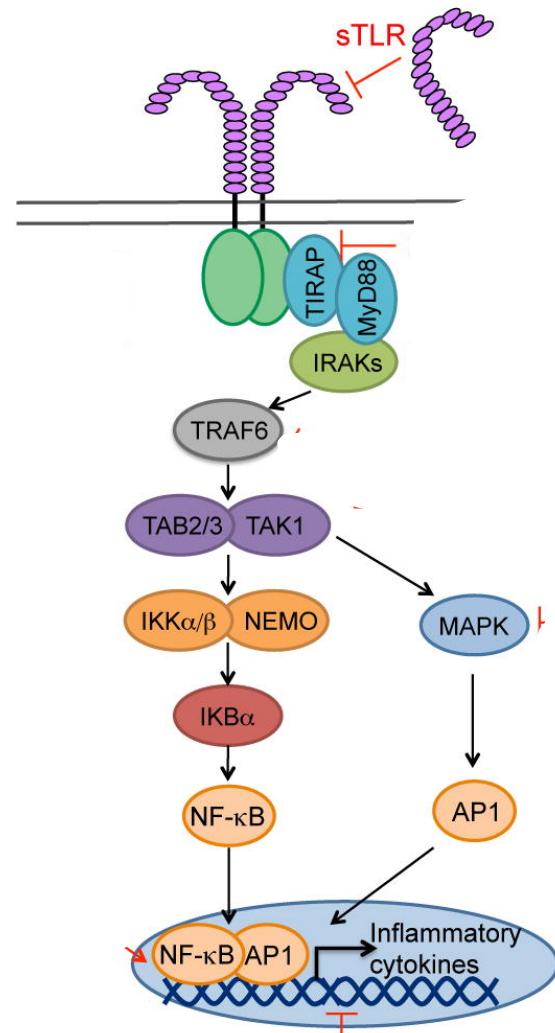
- ✓ cell wall components (LPS, β -glucan)
- ✓ flagellar components (flagellin)

DAMPS (danger-associated molecular patterns)

- ✓ Nuclear proteins (HMGB1), histones
- ✓ Purine metabolites such as ATP
- ✓ Uric acid
- ✓ Mitochondrial components (formyl peptides, mitochondrial DNA)



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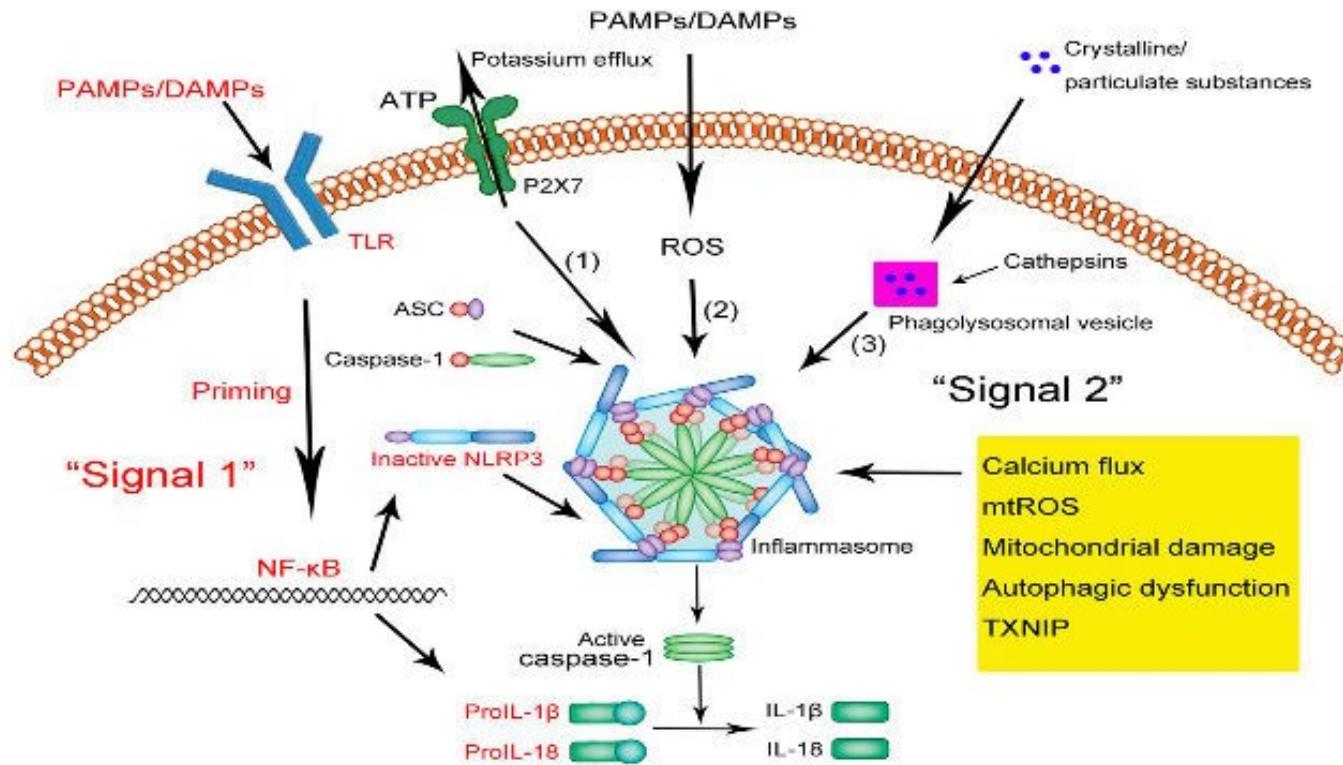
Germline-encoded pattern-recognition receptor (PRR)

TLR = Toll-like receptor

AP-1 = Activator protein-1



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Inflammasome



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Inflammation key elements

Chemical mediators

- ✓ Cytokine
- ✓ Chemokines
- ✓ Growth factors

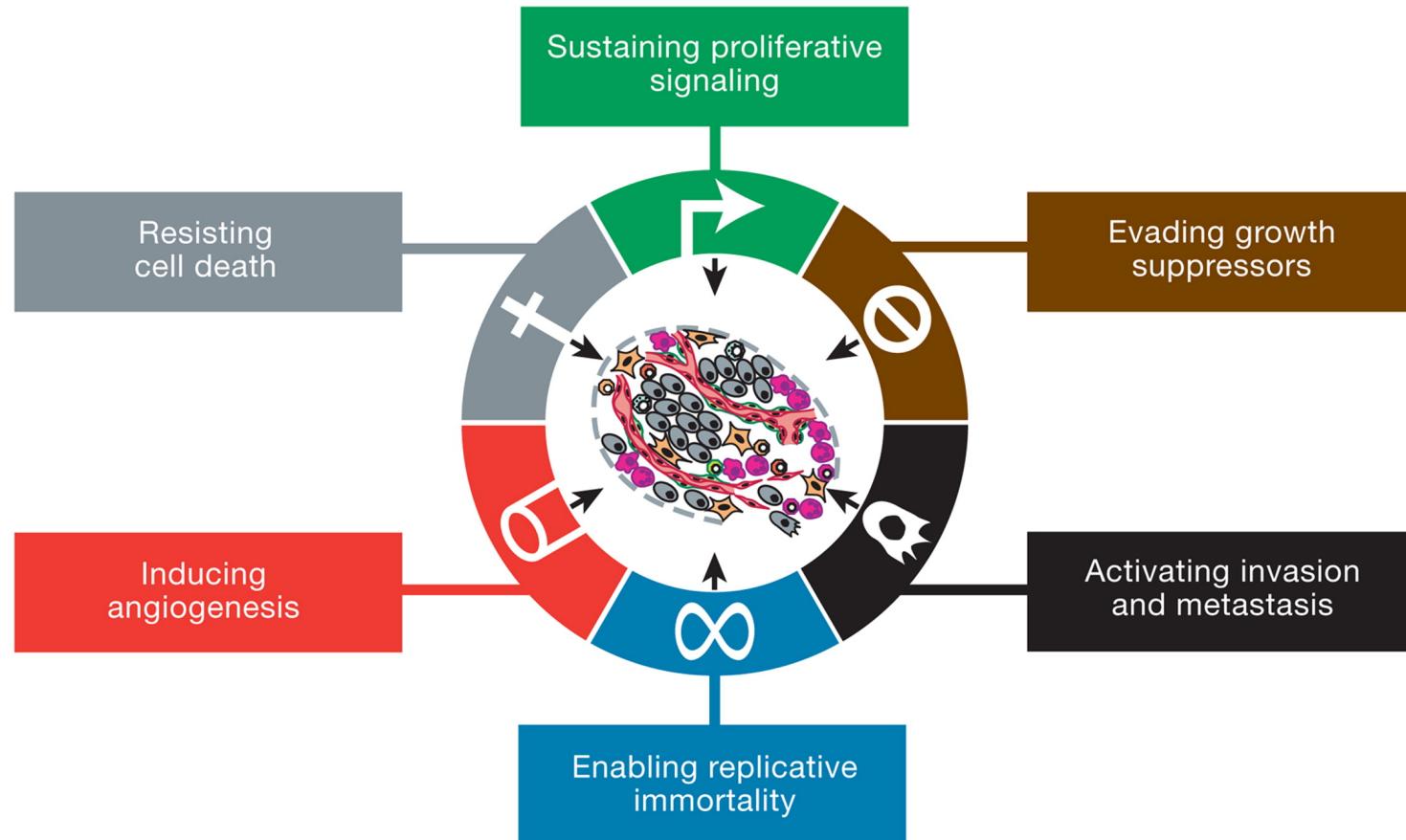
Cells

- ✓ Macrophages
- ✓ Neutrophils
- ✓ T-cells
- ✓ Endothelial cells



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Cancer hallmarks





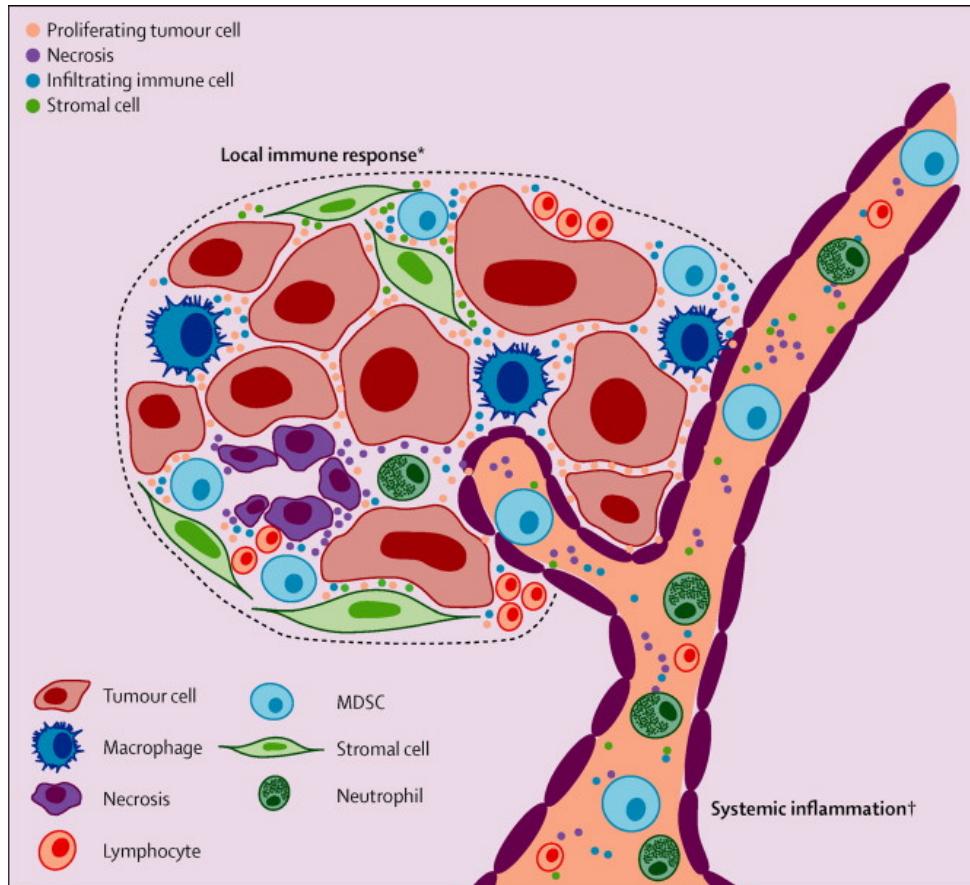
Mechanisms of cancer-related inflammation

Main characteristic	Main pathways	Specific complications
✓ Infiltration of tumor-associated macrophages with strong immunosuppressive activity	✓ M2 macrophage phenotype ✓ Checkpoint proteins PD-1, PD-L1 and CTLA-4 ✓ IL-1beta, IL-6, TNF, IL-4, IL-10, TGF-beta ✓ Pentraxin-3	✓ T cell exhaustion and anergy ✓ Tumor progression



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Highlights of onco-inflammation



1. Local and systemic dimension of inflammation



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Highlights of onco-inflammation

Prognostic use of systemic inflammation

- ✓ Neutrophil count, white cell count, neutrophil/lymphocyte ratio
- ✓ Glasgow prognostic score; albumin, CRP
- ✓ B symptoms
- ✓ Circulating myeloid derived suppressor cell (MDSC) count



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Highlights of onco-inflammation

2. Oncogenesis vs. oncoinflammation

OPEN ACCESS Freely available online



Molecular Profiling of Multiple Human Cancers Defines an Inflammatory Cancer-Associated Molecular Pattern and Uncovers KPNA2 as a Uniform Poor Prognostic Cancer Marker

Saleh M. Rachidi^{1,2}, Tingting Qin³, Shaoli Sun⁴, W. Jim Zheng^{3,5}, Zihai Li^{1,2*} 2013

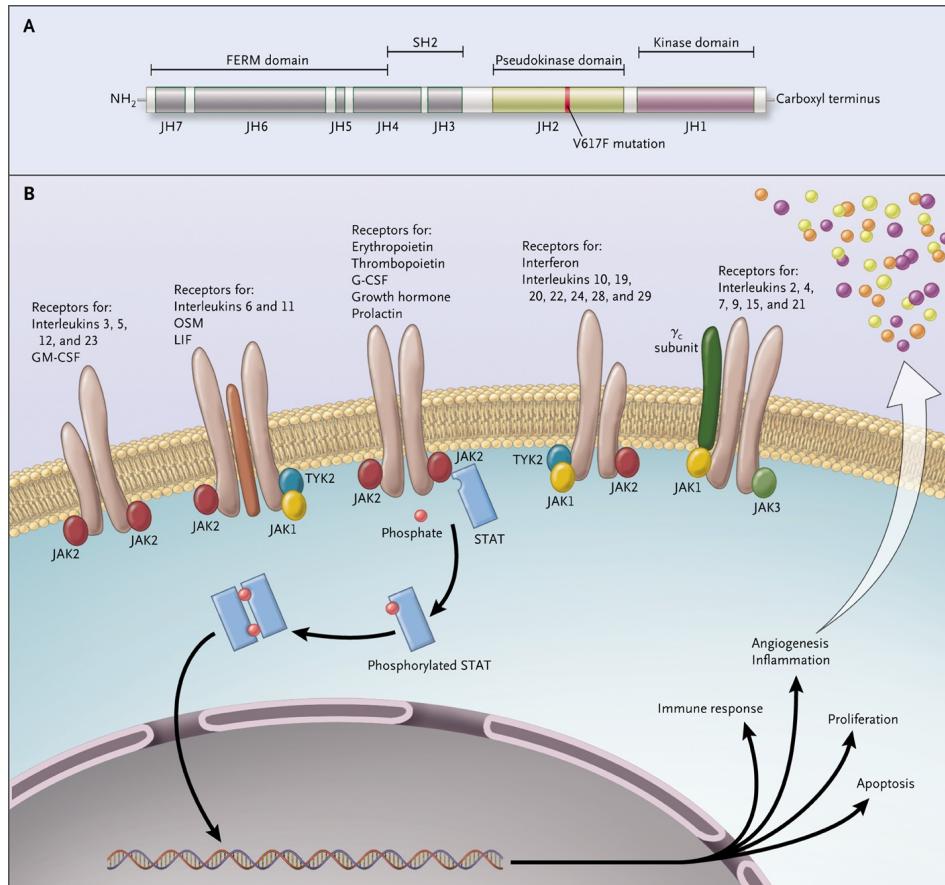
Gene expression data revealed an inflammatory signature shared by seven epithelial cancer types.

Onconflammation is a driver of oncogenesis.



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Highlights of onco-inflammation



2. Oncogenesis vs. oncoinflammation



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Highlights of onco-inflammation

2. Oncogenesis vs. oncoinflammation

OPEN

Leukemia (2018) 32, 438–449

www.nature.com/leu

ORIGINAL ARTICLE

Involvement of MAF/SPP1 axis in the development of bone marrow fibrosis in PMF patients

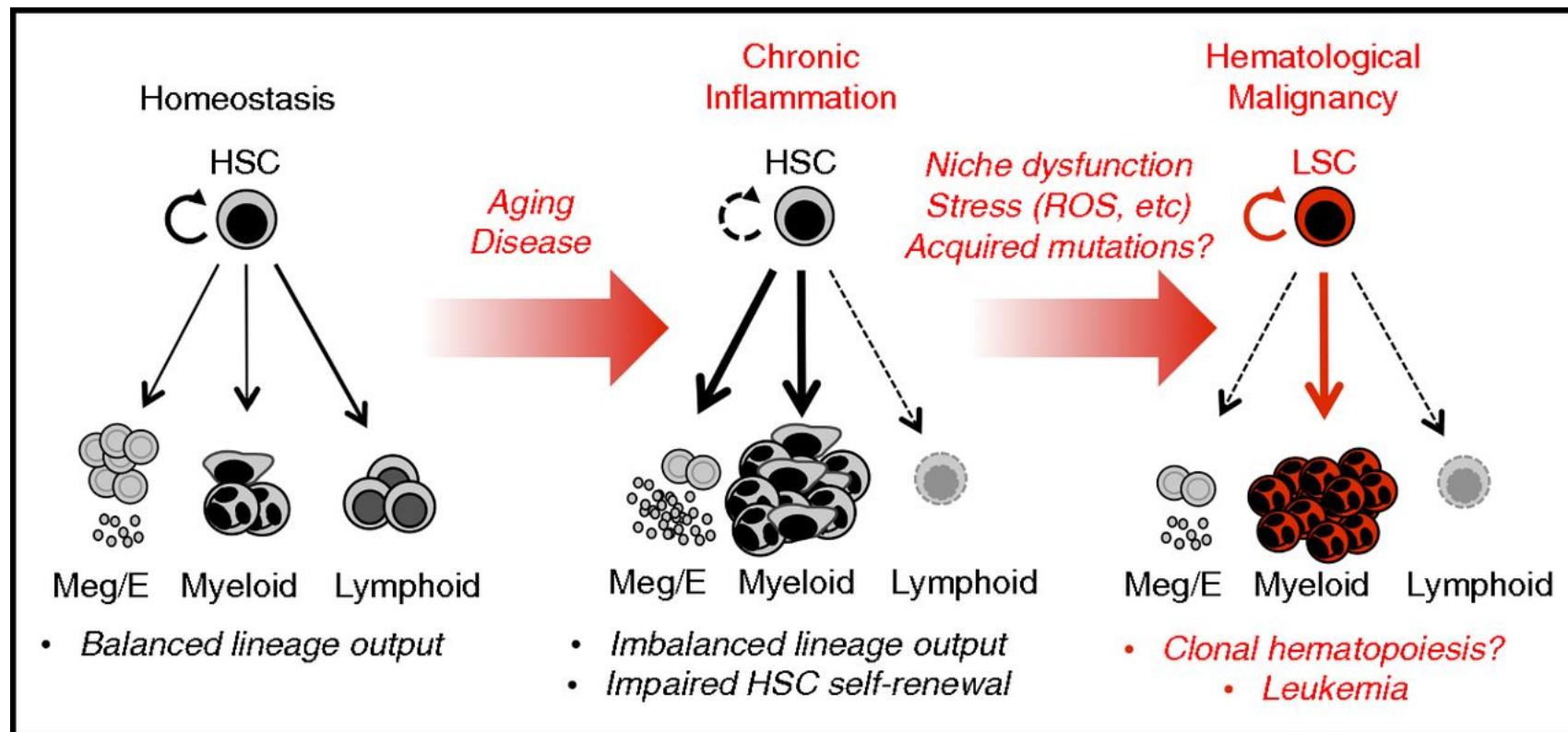
S Ruberti^{1,10}, E Bianchi^{1,10}, P Guglielmelli², S Rontauroli¹, G Barbieri¹, L Tavernari¹, T Fanelli², R Norfo^{1,3}, V Pennucci¹, G Corbizi Fattori^{2,4}, C Mannarelli^{2,4}, N Bartalucci², B Mora⁵, L Elli⁵, MA Avanzini⁶, C Rossi¹, S Salmoiraghi⁷, R Zini¹, S Salati¹, Z Prudente¹, V Rosti⁸, F Passamonti⁵, A Rambaldi⁷, S Ferrari⁹, E Tagliafico⁹, AM Vannucchi^{2,11}, R Manfredini^{1,11} on behalf of the AGIMM (AIRC Gruppo Italiano Malattie Mieloproliferative) Investigators



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Highlights of onco-inflammation

3. Age/inflammation determine the fate of HSCs





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Highlights of onco-inflammation

3. Age/inflammation determine the fate of HSCs

Leukemia Research 60 (2017) 18–23



Contents lists available at ScienceDirect

Leukemia Research
journal homepage: www.elsevier.com/locate/leukres



Research paper

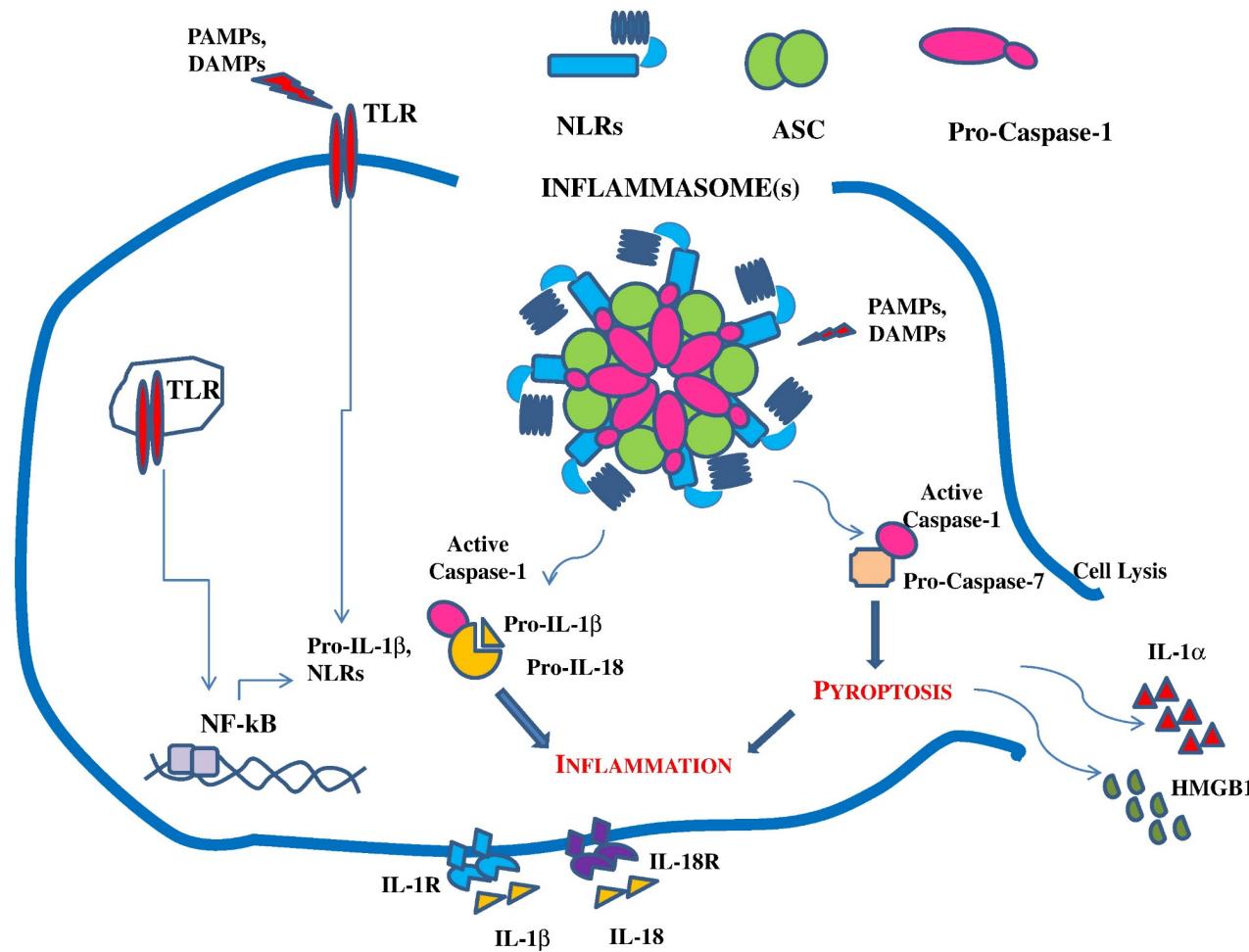
Primary myelofibrosis: Older age and high *JAK2V617F* allele burden are associated with elevated plasma high-sensitivity C-reactive protein levels and a phenotype of progressive disease

Giovanni Barosi^{a,*}, Margherita Massa^b, Rita Campanelli^a, Gabriela Fois^a, Paolo Catarsi^a, Gianluca Viarengo^c, Laura Villani^a, Valentina Poletto^a, Tiziana Bosoni^d, Umberto Magrini^a, Robert P. Gale^e, Vittorio Rosti^a





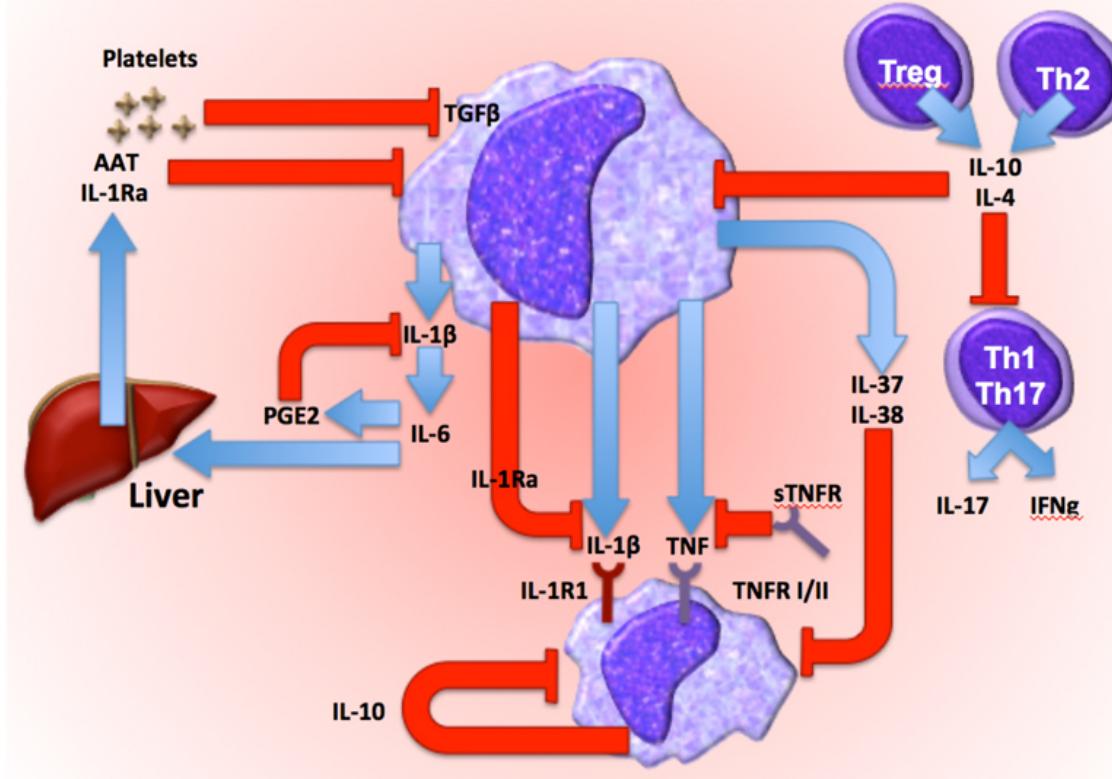
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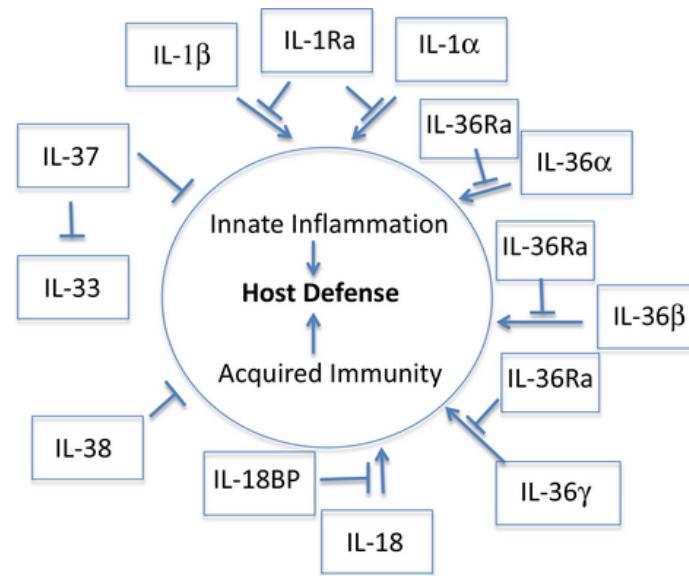
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Resolution of inflammation



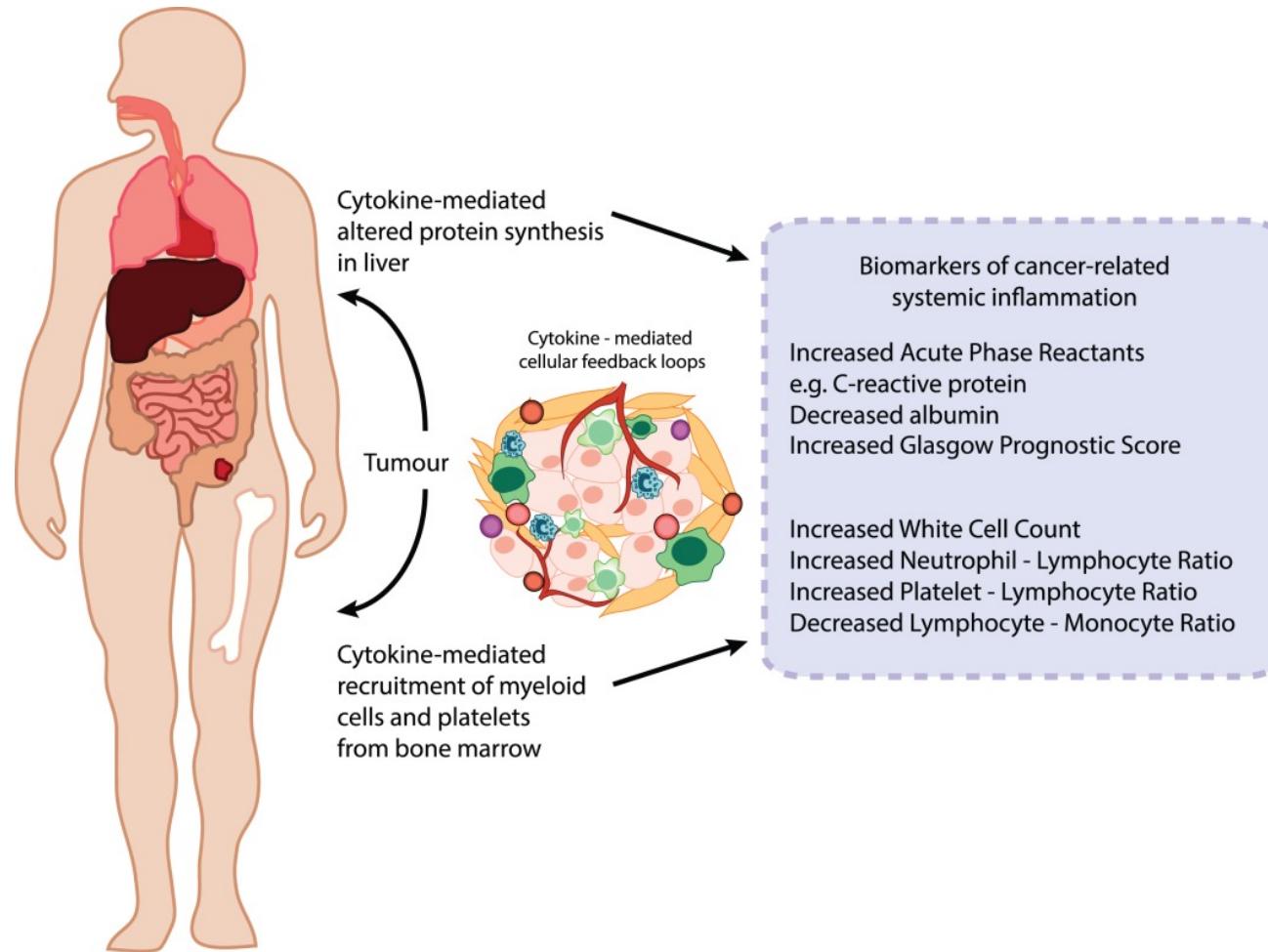


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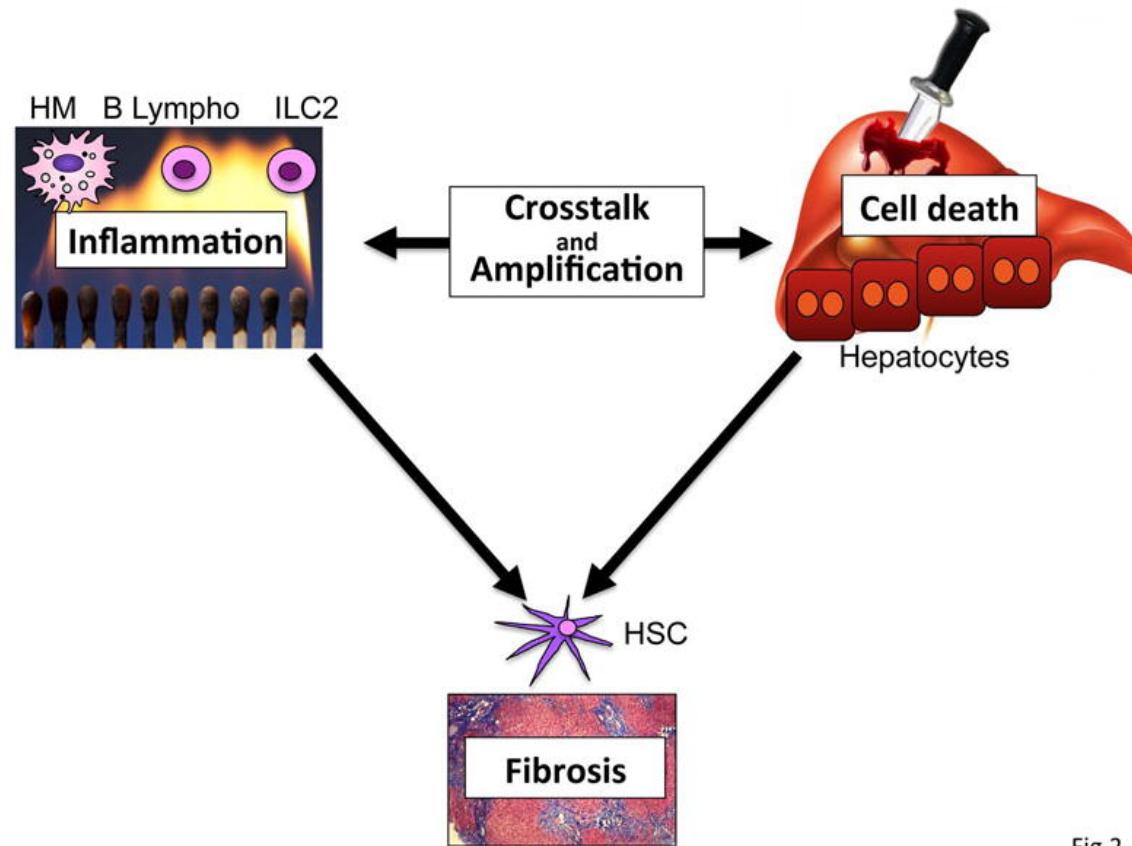
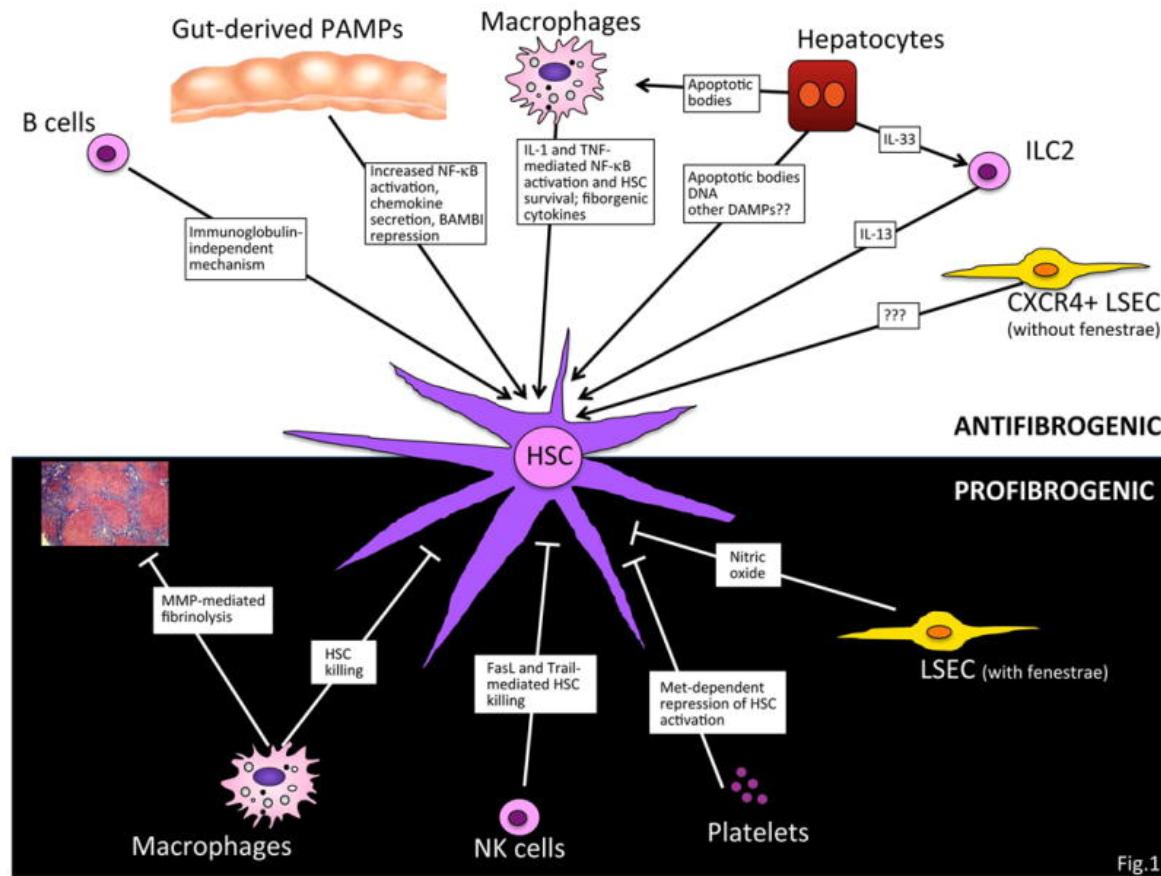


Fig.2

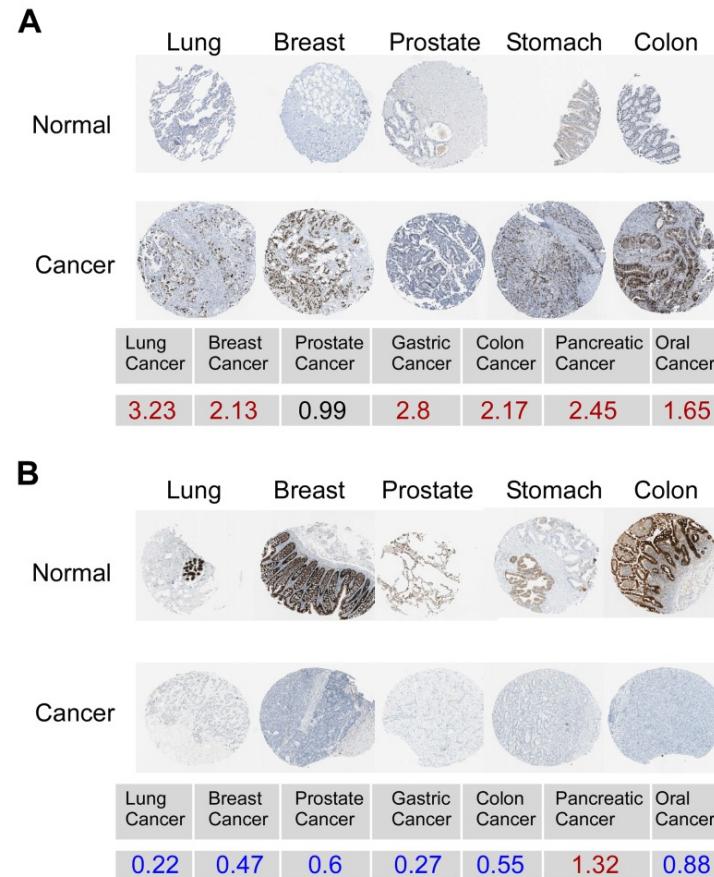


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Induction of inflammation

